Cause Group Code: C01E-17-PCB Chesapeake Bay and Tidal Tributaries

Cause Location: Chesapeake Bay mainstem and its small coastal tidal tributaries

Cause City/County: Accomack County; Chesapeake Bay - County Not Applicable; Gloucester County; Lancaster County; Mathews County; Middlesex County; Norfolk; Northampton County; Northumberland County; Poquoson City; Virginia Beach; York County

Use(s): Fish Consumption

Causes(s)/VA Category: PCBs in Fish Tissue/5A

Cause Description: The Chesapeake Bay and its small coastal tidal tributaries are included under the 12/13/2004 VDH Fish Consumption Advisories for PCBs. No more than two meals/month are recommended of anadromous (coastal) striped bass.

Also, VDH issued additional fish consumption advisory for PCBs in the Mobjack Bay and its tributaries, particularly the East, West, and Ware Rivers (on 12/13/2004) and in the Piankatank River from Rt. 17 to Deep Point Boat Landing (10/7/2009). No more than two meals/month of gizzard shad are recommended.

The advisories are based on the results of DEQ's fish tissue monitoring program, which show elevated PCBs levels in several monitoring sites within the basin, including:

7-COC000.40 in Cockrell Creek 7-DYM000.00 in Dymer Creek 7-DRN001.43 in Dragon Swamp 7-PNK019.85 and 7-PNK015.49 in the Piankatank River 7-MLF002.45 in Milford Haven 7-WIN000.88 in Winter Harbor 7-EST002.65 in the East River 7-WAR005.77 in the Ware River

Note: there were previous exceedances at 7-GWR007.97 in the Great Wicomico River, 7-IND001.80 in Indian Creek, and 7-NOR003.65 in the North River. Re-sampling at those stations was acceptable in the 2022 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_OYS01B24 / Oyster Creek / Portion of VDH condemnation 018-053A, 1/4/2005 within RPPMH. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.071

Chesapeake Bay and Tidal Tributaries

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
PCBs in Fish Tissue - Total Impaired Size by Water Type:	0.071		

Sources: Atmospheric Deposition - Toxics; Source Unknown

Cause Group Code: CRRMH-DO-BAY Corrotoman River Mesohaline Estuary (CRRMH)

Cause Location: The Corrotoman River and its tidal tributaries (CRRMH).

Cause City/County: Lancaster County

Use(s): Aquatic Life; Open-Water Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/4A

Cause Description: The mainstem Corrotoman River was included in EPA's 1998 Overlist. The Chesapeake Bay water quality standards were implemented during the 2006 cycle.

The Corrotoman River mesohaline estuary fails the Chesapeake Bay Open Water Subuse's summer 30-day mean dissolved oxygen criterion. The Open Water rest-of-year criteria was met in the 2024 cycle. There is insufficient data to assess the other dissolved oxygen criteria.

The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, CRRMH is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BES01A98 / Bells Creek / As described in VDH shellfish condemnation 58B, 4/28/1997. CRRMH	4A	Dissolved Oxygen	2006	L	0.055
VAP-E26E_BLD01A98 / Belwood Swamp / Tidal limit to its mouth at the Western Branch Corrotoman River. CRRMH	4A	Dissolved Oxygen	2006	L	0.009
VAP-E26E_CRR01A00 / Corrotoman River / The mainstem of the Corrotoman River within segment CRRMH.	4A	Dissolved Oxygen	1998	L	3.769
VAP-E26E_CTM01A00 / Eastern Branch Corrotoman River / Non-administrative portion of VDH-DSS Condemnations 021U-058B and condemnations 021U-058C & -058D, 11/15/2022. Shortened and split in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.429
VAP-E26E_CTM01B24 / Eastern Branch Corrotoman River, UT / Administratively condemned portion of VDH shellfish condemnation 021U-058B, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.007
VAP-E26E_CTM01C20 / Eastern Branch Corrotoman River / Portion of VDH shellfish condemnation 058C, 4/28/1997 that is seasonally condemned. Expanded and merged in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.186
VAP-E26E_CTM02A08 / Eastern Branch Corrotoman River, UT / Described in VDH Shellfish Condemnation 021U-058E, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.011

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM03A08 / Eastern Branch Corrotoman River / Downstream boundary of VDH condemnation 021-058C, 4/28/1997 to rivermile 0.88. CRRMH	4A	Dissolved Oxygen	2006	L	0.348
VAP-E26E_CTM03B24 / Eastern Branch Corrotoman River / Rivermile 0.88 to mouth. CRRMH	4A	Dissolved Oxygen	2006	L	0.410
VAP-E26E_CTO01A02 / Western Branch Corrotoman River / As described in VDH shellfish condemnation 021U-132A, 11/15/2022, not otherwise segmented. CRRMH	4A	Dissolved Oxygen	2006	L	0.452
VAP-E26E_CTO01B12 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 included in 021U-132S64, 11/15/20202. CRRMH	4A	Dissolved Oxygen	2006	L	0.046
VAP-E26E_CTO01C22 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 open in VDH-DSS condemnation 021U-132, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.100
VAP-E26E_CTO02A06 / Western Branch Corrotoman River / Mainstem Western Corrotoman River from rivermile 2.21 to the mouth. Split in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.795
VAP-E26E_CTO02B24 / Western Branch Corrotoman River / Mainstem from the downstream boundary of VDH Shellfish Condemnation 132A, 4/28/1997 to rivermile 2.21. Split in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.414
VAP-E26E_DAS01A02 / Davis Creek / As described in VDH-DSS condemnation 021-132S63, 11/15/2022. Segment expanded in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.043
VAP-E26E_EWE01B20 / Ewells Prong / As described in VDH shellfish condemnation 187A, 4/28/1997. Merged in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.036
VAP-E26E_EWE02A08 / Ewells Prong / Portion of VDH Shellfish Condemnation 021L-187S53, 11/15/2022 not included on 187A or 187B, 4/28/1997. Expanded in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.032

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_HLS01A00 / Hills Creek / As described in VDH shellfish condemnation 021U-058A, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.038
VAP-E26E_HLS01B20 / Hills Creek / The portion of VDH shellfish condemnation 58A, 4/28/1997 seasonally condemned/open (021U-058S59, 11/15/2022). CRRMH	4A	Dissolved Oxygen	2006	L	0.024
VAP-E26E_JON01A08 / John Creek / Described in VDH-DSS Condemnation 021U-132210, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.036
VAP-E26E_JON02A08 / John Creek / Downstream of VDH-DSS condemnation 021U-132S210, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.016
VAP-E26E_LIT01A06 / Little Branch / Tidal limit to mouth at Western Branch Corrotoman River CRRMH	4A	Dissolved Oxygen	2006	L	0.114
VAP-E26E_LOW01A08 / Lowrey Creek / Described in VDH Shellfish Condemnation 021U-132S62, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.028
VAP-E26E_MIP01A00 / Millenbeck Prong / Portion of VDH shellfish condemnation 187B, 4/28/1997 seasonally condemned in 021L-187S53, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.004
VAP-E26E_MIP01B20 / Millenbeck Prong / Described in VDH shellfish condemnation 021L-187C, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.037
VAP-E26E_MOR01B12 / Moran Creek / Described in VDH-DSS condemnation 021L-198S57, 11/15/2021. Expanded in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.045
VAP-E26E_MOR01C22 / Moran Creek, UT / Described in VDH-DSS condemnation 021L-198D, 11/15/2021. Merged in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.049
VAP-E26E_MOR02A08 / Moran Creek / Downstream of VD-DSS condemnation 021L-198, 11/15/2021. CRRMH	4A	Dissolved Oxygen	2006	L	0.060
VAP-E26E_MYE01A00 / Myer Creek / As described in VDH shellfish condemnation 198, 4/28/1997. CRRMH	4A	Dissolved Oxygen	2006	L	0.081

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MYE01B02 / Myer Creek, UT / As described in VDH-DSS condemnation 021L-198S58, 11/15/2022. Expanded in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.042
VAP-E26E_MYE01D18 / Myer Creek / Portion of VDH-DSS condemnation 021L-198A, 11/15/2022 open in 198, 4/28/1997. CRRMH	4A	Dissolved Oxygen	2006	L	0.095
VAP-E26E_MYE03A08 / Myer Creek / Downstream of condemnations to mouth at Corrotoman River. Split in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.442
VAP-E26E_MYE03B24 / Myer Creek / Described in VDH-DSS condemnation 021L-198S56, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.028
VAP-E26E_SEN01A00 / Senior Creek / As described in VDH shellfish condemnation 021U-132B and -132C, 11/15/2022. Expanded in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.053
VAP-E26E_SEN01B20 / Senior Creek / As described in VDH shellfish condemnation 021U-132S105, 11/15/2022. Shrank in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.017
VAP-E26E_TAY01A00 / Taylor Creek / As described in VDH-DSS condemnations 021L-198B and -C, 11/15/2021. CRRMH	4A	Dissolved Oxygen	2006	L	0.068
VAP-E26E_TAY02A08 / Taylor Creek / Portion of VDH-DSS condemnation 205, 4/28/1997 within 021L-198S55, 11/15/2021. CRRMH	4A	Dissolved Oxygen	2006	L	0.123
VAP-E26E_TAY03A24 / Taylor Creek / Portion of VDH-DSS Shellfish Condemnation 021L-198S55, 11/15/2021 not included in VDH condemnation 205, 4/28/1997. CRRMH	4A	Dissolved Oxygen	2006	L	0.011
VAP-E26E_TON01A00 / Town Creek / Described in VDH shellfish condemnation 021L-187S54, 11/15/2022. Shrank in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.018

(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHR01A00 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187SS52, 11/15/2022. Expanded and split in the 2024 cycle. CRRMH	4A	Dissolved Oxygen	2006	L	0.052
VAP-E26E_WHR01B24 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187A, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.045
VAP-E26E_WHR01C24 / Whitehouse Creek, UT / As described in VDH shellfish condemnation 021L-187B, 11/15/2022. CRRMH	4A	Dissolved Oxygen	2006	L	0.005
VAP-E26E_ZZZ02A14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA70 CRRMH	4A	Dissolved Oxygen	2006	L	0.091
VAP-E26E_ZZZ02C14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA72 CRRMH	4A	Dissolved Oxygen	2006	L	0.445

Corrotoman River Mesohaline Estuary (CRRMH)

Aquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 9.204	Reservoir (Acres)	River (Miles)
Corrotoman River N	Mesohaline Estuary (CRRMH)	Estuary	Reservoir	River
Open-Water Aqu	atic Life Dissolved Oxygen - Total Impaired Size by Water Type:	(Sq. Miles) 9.204	(Acres)	(Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: CRRMH-SAV-BAY Corrotoman River Mesohaline Estuary (CRRMH)

Cause Location: The Corrotoman River and its tidal tributaries (CRRMH).

Cause City/County: Lancaster County

Use(s): Aquatic Life; Shallow-Water Submerged Aquatic Vegetation

Causes(s)/VA Category: Aquatic Plants (Macrophytes)/4A

Cause Description: The mesohaline Corrotoman River (CRRMH) has been impaired of the Chesapeake Bay Shallow Water Subuse since the 2012 cycle. CRRMH did not meet the Shallow Water Subuse's submerged aquatic vegetation acreage criterion or the water clarity acreage criterion in the 2024 cycle. The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, CRRMH is considered Category 4A.

The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, CRRMH is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BES01A98 / Bells Creek / As described in VDH shellfish condemnation 58B, 4/28/1997. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.055
VAP-E26E_BLD01A98 / Belwood Swamp / Tidal limit to its mouth at the Western Branch Corrotoman River. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.009
VAP-E26E_CRR01A00 / Corrotoman River / The mainstem of the Corrotoman River within segment CRRMH.	4A	Aquatic Plants (Macrophytes)	2012	L	3.769
VAP-E26E_CTM01A00 / Eastern Branch Corrotoman River / Non-administrative portion of VDH-DSS Condemnations 021U-058B and condemnations 021U-058C & -058D, 11/15/2022. Shortened and split in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.429
VAP-E26E_CTM01B24 / Eastern Branch Corrotoman River, UT / Administratively condemned portion of VDH shellfish condemnation 021U-058B, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.007
VAP-E26E_CTM01C20 / Eastern Branch Corrotoman River / Portion of VDH shellfish condemnation 058C, 4/28/1997 that is seasonally condemned. Expanded and merged in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.186
VAP-E26E_CTM02A08 / Eastern Branch Corrotoman River, UT / Described in VDH Shellfish Condemnation 021U-058E, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.011

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM03A08 / Eastern Branch Corrotoman River / Downstream boundary of VDH condemnation 021-058C, 4/28/1997 to rivermile 0.88. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.348
VAP-E26E_CTM03B24 / Eastern Branch Corrotoman River / Rivermile 0.88 to mouth. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.410
VAP-E26E_CTO01A02 / Western Branch Corrotoman River / As described in VDH shellfish condemnation 021U-132A, 11/15/2022, not otherwise segmented. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.452
VAP-E26E_CTO01B12 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 included in 021U-132S64, 11/15/20202. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.046
VAP-E26E_CTO01C22 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 open in VDH-DSS condemnation 021U-132, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.100
VAP-E26E_CTO02A06 / Western Branch Corrotoman River / Mainstem Western Corrotoman River from rivermile 2.21 to the mouth. Split in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.795
VAP-E26E_CTO02B24 / Western Branch Corrotoman River / Mainstem from the downstream boundary of VDH Shellfish Condemnation 132A, 4/28/1997 to rivermile 2.21. Split in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.414
VAP-E26E_DAS01A02 / Davis Creek / As described in VDH-DSS condemnation 021-132S63, 11/15/2022. Segment expanded in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.043
VAP-E26E_EWE01B20 / Ewells Prong / As described in VDH shellfish condemnation 187A, 4/28/1997. Merged in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.036
VAP-E26E_EWE02A08 / Ewells Prong / Portion of VDH Shellfish Condemnation 021L-187S53, 11/15/2022 not included on 187A or 187B, 4/28/1997. Expanded in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.032

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_HLS01A00 / Hills Creek / As described in VDH shellfish condemnation 021U-058A, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.038
VAP-E26E_HLS01B20 / Hills Creek / The portion of VDH shellfish condemnation 58A, 4/28/1997 seasonally condemned/open (021U-058S59, 11/15/2022). CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.024
VAP-E26E_JON01A08 / John Creek / Described in VDH-DSS Condemnation 021U-132210, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.036
VAP-E26E_JON02A08 / John Creek / Downstream of VDH-DSS condemnation 021U-132S210, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.016
VAP-E26E_LIT01A06 / Little Branch / Tidal limit to mouth at Western Branch Corrotoman River CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.114
VAP-E26E_LOW01A08 / Lowrey Creek / Described in VDH Shellfish Condemnation 021U-132S62, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.028
VAP-E26E_MIP01A00 / Millenbeck Prong / Portion of VDH shellfish condemnation 187B, 4/28/1997 seasonally condemned in 021L-187S53, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.004
VAP-E26E_MIP01B20 / Millenbeck Prong / Described in VDH shellfish condemnation 021L-187C, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.037
VAP-E26E_MOR01B12 / Moran Creek / Described in VDH-DSS condemnation 021L-198S57, 11/15/2021. Expanded in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.045
VAP-E26E_MOR01C22 / Moran Creek, UT / Described in VDH-DSS condemnation 021L-198D, 11/15/2021. Merged in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.049
VAP-E26E_MOR02A08 / Moran Creek / Downstream of VD-DSS condemnation 021L-198, 11/15/2021. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.060
VAP-E26E_MYE01A00 / Myer Creek / As described in VDH shellfish condemnation 198, 4/28/1997. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.081

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MYE01B02 / Myer Creek, UT / As described in VDH-DSS condemnation 021L-198S58, 11/15/2022. Expanded in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.042
VAP-E26E_MYE01D18 / Myer Creek / Portion of VDH-DSS condemnation 021L-198A, 11/15/2022 open in 198, 4/28/1997. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.095
VAP-E26E_MYE03A08 / Myer Creek / Downstream of condemnations to mouth at Corrotoman River. Split in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.442
VAP-E26E_MYE03B24 / Myer Creek / Described in VDH-DSS condemnation 021L-198S56, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.028
VAP-E26E_SEN01A00 / Senior Creek / As described in VDH shellfish condemnation 021U-132B and -132C, 11/15/2022. Expanded in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.053
VAP-E26E_SEN01B20 / Senior Creek / As described in VDH shellfish condemnation 021U-132S105, 11/15/2022. Shrank in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.017
VAP-E26E_TAY01A00 / Taylor Creek / As described in VDH-DSS condemnations 021L-198B and -C, 11/15/2021. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.068
VAP-E26E_TAY02A08 / Taylor Creek / Portion of VDH-DSS condemnation 205, 4/28/1997 within 021L-198S55, 11/15/2021. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.123
VAP-E26E_TAY03A24 / Taylor Creek / Portion of VDH-DSS Shellfish Condemnation 021L-198S55, 11/15/2021 not included in VDH condemnation 205, 4/28/1997. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.011
VAP-E26E_TON01A00 / Town Creek / Described in VDH shellfish condemnation 021L-187S54, 11/15/2022. Shrank in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.018

(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHR01A00 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187SS52, 11/15/2022. Expanded and split in the 2024 cycle. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.052
VAP-E26E_WHR01B24 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187A, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.045
VAP-E26E_WHR01C24 / Whitehouse Creek, UT / As described in VDH shellfish condemnation 021L-187B, 11/15/2022. CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.005
VAP-E26E_ZZZ02A14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA70 CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.091
VAP-E26E_ZZZ02C14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA72 CRRMH	4A	Aquatic Plants (Macrophytes)	2012	L	0.445

Corrotoman River Mesohaline Estuary (CRRMH)

Aquatic Life Aquatic Plants (Macrophytes) - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 9.204	Reservoir (Acres)	River (Miles)
Corrotoman River Mesohaline Estuary (CRRMH)		D .	D:
Shallow-Water Submerged Aquatic Vegetation Aquatic Plants (Macrophytes) - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 9.204	Reservoir (Acres)	River (Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E01R-01-BAC Thumb Run

Cause Location: Begins at the confluence of West Branch Thumb Run and East Branch Thumb Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-THU004.69 at Route 688 (Leeds Manor Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

The Thumb Run Watershed bacteria TMDL (Eq ID POL0117) was approved by the EPA on 05/31/2002 (Fed ID 24413). The SWCB approved the TMDL on 06/17/2004. A bacteria TMDL Implementation Plan for the Thumb Run watershed (ID 98) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_THU01A00 / Thumb Run / Segment begins at the confluence of West Branch Thumb Run and East Branch Thumb Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2010	L	7.68

Thumb Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
]	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	,		7.68

Cause Group Code: E01R-01-BEN Thumb Run, East Branch

Cause Location: Begins at the headwaters of East Branch Thumb Run and continues downstream until the confluence of East Branch to the mainstem Thumb Run.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2018 Assessment: A total of three biological monitoring events in 2011 and 2012 at DEQ station 3-THM001.40 at Route 647 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_THM01A02 / Thumb Run, East Branch / Segment begins at the headwaters of East Branch Thumb Run and continues downstream until the confluence of East Branch to the mainstem Thumb Run.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	6.59

Thumb Run, East Branch

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			6.59

Sources: Source Unknown

Cause Group Code: E01R-02-BAC Thumb Run, West Branch

Cause Location: Begins at the headwaters of West Branch Thumb Run and continues downstream until the confluence of West Branch to the mainstem Thumb Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-THW004.68 at Route 635 (Hume Rd).

A new TMDL is not required for this impaired segment of West Branch Thumb Run because the downstream Thumb Run Watershed bacteria TMDL (Fed ID 24413, 05/31/2002) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0117). A bacteria TMDL Implementation Plan for the Thumb Run watershed (ID 98) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_THW01A02 / Thumb Run, West Branch / Segment starts at the headwaters of West Branch Thumb Run and continues downstream until the confluence of West Branch to the mainstem Thumb Run.	4A	Escherichia coli (E. coli)	2002	L	12.09

Thumb Run,	West Branch			
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			12.09

Cause Group Code: E01R-02-BEN Unnamed Tributary to Thumb Run, West Branch

Cause Location: Segment begins at the headwaters of an unnamed tributary to West Branch Thumb Run and continues downstream until the confluence with West Branch Thumb Run.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2018 Assessment: A total of two biological monitoring events in 2011 at DEQ station 3-XHU000.04 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_XHU01A14 / Unnamed Tributary to Thumb Run, West Branch / Segment begins at the headwaters of an unnamed tributary to West Branch Thumb Run and continues downstream until the confluence with West Branch Thumb Run.	5A	Benthic Macroinvertebrates Bioassessments	2014	L	0.8

Unnamed Tributary to Thumb Run, West Branch

	Estuary	Reservoir	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			0.8	

Sources: Source Unknown

Cause Group Code: E01R-03-BAC Rappahannock River

Cause Location: Begins at the headwaters of the Rappahannock River and continues downstream until the confluence with Fiery Run. Begins again at the confluence with the Jordan River, at rivermile 175.58, and continues downstream until the confluence with an unnamed tributary to the Rappahannock River, at rivermile 173.41.

Cause City/County: Fauquier County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were collected for the 2024 assessment from DEQ station 3-RPP175.51 at Route 647 (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: E. coli bacteria criterion excursions (7 of 33 samples - 21.2%).

2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples and the geomean was exceeded in at least one 90-day period with 10+ samples at DEQ station 3-RPP186.59 at Route 635; additionally, there were two or more STV exceedances in at least one 90-day period with <10 samples

The Upper Rappahannock River Watershed bacteria TMDL for the Rappahannock River (1) watershed (Eq ID POL0516) was approved by the EPA on 01/23/2008 (Fed ID 33913). The SWCB approved the TMDL on 07/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_RPP02A00 / Rappahannock River / Segment begins at the confluence with the Jordan River, at rivermile 175.58, and continues downstream until the confluence with an unnamed tributary to the Rappahannock River, at rivermile 173.41.	4A	Escherichia coli (E. coli)	2006	L	2.28
VAN-E01R_RPP03A04 / Rappahannock River / Segment begins at the headwaters of the Rappahannock River and continues downstream until the confluence with Fiery Run.	4A	Escherichia coli (E. coli)	2020	L	7.77

Rappahannock River				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			10.05	

Cause Group Code: E01R-03-BEN Hittles Mill Stream

Cause Location: Begins at the confluence with Bearwallow Creek and Bolton Branch and continues downstream to the confluence with Jordan River.

Cause City/County: Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of four biological monitoring events in 2017 and 2018 at DEQ station 3-HIT003.43 at Route 631 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_HIT01A14 / Hittles Mill Stream / Segment begins at the confluence with Bearwallow Creek and Bolton Branch and continues downstream to the confluence with Jordan River.	5A	Benthic Macroinvertebrates Bioassessments	2020	L	3.83
Hittles Mill Stream		Estuary	Rese	ervoir Riv	ver

	Estuary	$\operatorname{Reservoir}$	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			3.83	

Sources: Source Unknown

Cause Group Code: E01R-04-BAC Thumb Run, East Branch

Cause Location: Begins at the headwaters of East Branch Thumb Run and continues downstream until the confluence of East Branch to the mainstem Thumb Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-THM001.40 at Route 647 (Cresthill Rd).

A new TMDL is not required for this impaired segment of Thumb Run, East Branch because the downstream Thumb Run Watershed bacteria TMDL (Fed ID 24413, 05/31/2002) included modeling, source identification, and reductions that covered the entire Thumb Run watershed (Eq ID POL0117). The bacteria TMDL Implementation Plan for the Thumb Run watershed (ID 98) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_THM01A02 / Thumb Run, East Branch / Segment begins at the headwaters of East Branch Thumb Run and continues downstream until the confluence of East Branch to the mainstem Thumb Run.	4A	Escherichia coli (E. coli)	2004	L	6.59

Thumb Run, East Branch				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) -	Total Impaired Size by Water Type:			6.59

Cause Group Code: E01R-05-BAC Fiery Run

Cause Location: Begins at the headwaters of Fiery Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-FIR002.35 at Route 635 (Hume Rd).

A new TMDL is not required for this impaired segment of Fiery Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33913, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (1) watershed (Eq ID POL0516).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_FIR01A04 / Fiery Run / Segment begins at the headwaters of Fiery Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2010	L	9.39

Fiery Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(54. 11105)	(110105)	9.39

Cause Group Code: E01R-06-BAC Jordan River

Cause Location: Begins at the start of Class III water at rivermile 10.9 and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-JOR000.50 at Route 637 (N Poes Road).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-JOR007.56 at Route 522.

A new TMDL is not required for this impaired segment of Fiery Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33913, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (1) watershed (Eq ID POL0516).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_JOR01A04 / Jordan River / Segment begins at the confluence of Hittles Mill Stream, at rivermile 7.05, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2012	L	7.05
VAN-E01R_JOR02A20 / Jordan River / Segment begins at the start of Class III water at rivermile 10.9 and continues downstream to the confluence with Hittles Mill Stream.	4A	Escherichia coli (E. coli)	2020	L	3.85

Jordan River

D		•	Reservoir	
Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(Sq. Miles)	(Acres)	(Miles) 10.9

Cause Group Code: E01R-07-BAC Buck Run

Cause Location: Begins at the headwaters of Buck Run to the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BUC001.54 at Route 735 (Keyser Rd).

A new TMDL is not required for this impaired segment of Buck Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33913, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (1) watershed (Eq ID POL0516).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_BUC01A10 / Buck Run / Segment begins at the headwaters of Buck Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2016	L	9.76

Buck Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, - , ,		9.76

Cause Group Code: E01R-08-BAC Rappahannock River

Cause Location: Begins at the confluence with an unnamed tributary to the Rappahannock River, at rivermile 173.41, and continues downstream until the mouth of watershed E01R, at the confluence with Thumb Run.

Cause City/County: Fauquier County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP170.36 at Route 645 (Tapps Ford Rd).

A new TMDL is not required for this impaired segment of the Rappahannock River because the downstream Upper Rappahannock River bacteria TMDL (Fed ID 33951, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (2) watershed (Eq ID POL0508).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_RPP01A04 / Rappahannock River / Segment begins at the confluence with an unnamed tributary to the Rappahannock River, at rivermile 173.41, and continues downstream until the mouth of watershed E01R, at the confluence with Thumb Run.	4A	Escherichia coli (E. coli)	2020	L	4.09

Rappahannock River				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.09	

Cause Group Code: E01R-09-BAC Indian Run

Cause Location: Begins at the confluence with an unnamed tributary at rivermile 1.87 and continues downstream to the confluence with Hittles Mill Stream.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-INA000.38 at Route 637 (Jericho Rd).

A new TMDL is not required for this impaired segment of Fiery Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33913, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (1) watershed (Eq ID POL0516).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E01R_INA01A20 / Indian Run / Segment begins at the confluence with an unnamed tributary at rivermile 1.87 and continues downstream to the confluence with Hittles Mill Stream.	4A	Escherichia coli (E. coli)	2020	L	1.88

Indian Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /	, , ,	1.88

Cause Group Code: E02R-01-BAC Carter Run

Cause Location: Begins at the confluence with Horner Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-CAE000.25 at Route 688 (Leeds Manor Rd).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-CAE006.32 at Route 738 (Wilson Rd).

The Carter Run Watershed bacteria TMDL (Eq ID POL0155) was approved by the EPA on 03/10/2005 (Fed ID 24414). The SWCB approved the modified TMDL on 12/20/2005. The bacteria TMDL Implementation Plan for the Carter Run watershed (ID 99) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_CAE01A00 / Carter Run / Segment begins at the confluence with South Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	1998	L	3.62
VAN-E02R_CAE02A04 / Carter Run / Segment begins at the uppermost point of the PWS designation (five miles upstream from Waterloo) and continues downstream until the confluence with South Run.	4A	Escherichia coli (E. coli)	2006	L	1.38
VAN-E02R_CAE02B12 / Carter Run / Segment begins at the confluence with Horner Run and continues downstream until the beginning of the PWS designation (five miles upstream from Waterloo).	4A	Escherichia coli (E. coli)	2006	L	7.39

Carter Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	× - /	``	12.39

Sources: Grazing in Riparian or Shoreline Zones; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E02R-01-BEN Great Run

Cause Location: Begins at the confluence with an unnamed tributary to Great Run (streamcode XAC) at rivermile 7.20 (approximately 0.6 rivermile downstream from Route 802) and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2018 Assessment: A total of three biological monitoring events in 2011 and 2012 at DEQ station 3-GRT001.70 at Route 687 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_GRT01A00 / Great Run / Segment begins at the confluence with an unnamed tributary to Great Run, approximately 1.0 rivermile upstream of Route 687, and continues downstream until the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	2.82
VAN-E02R_GRT02A04 / Great Run / Segment begins at the confluence of an unnamed tributary to Great Run, at approximately rivermile 5.5, and continues downstream until the confluence with an unnamed tributary to Great Run, approximately 1.0 rivermile upstream of Route 687.	5A	Benthic Macroinvertebrates Bioassessments	2010	L	2.85
VAN-E02R_GRT03A02 / Great Run / Segment begins at the confluence with an unnamed tributary to Great Run at rivermile 7.20 (streamcode XAC) and continues downstream until the confluence with another unnamed tributary at approximately rivermile 5.5.	5A	Benthic Macroinvertebrates Bioassessments	2010	L	1.54

Great Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			7.21

Sources: Source Unknown

Cause Group Code: E02R-02-BAC Great Run

Cause Location: Begins at the headwaters of Great Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GRT001.70 at Route 687 (Opal Rd).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GRT007.72 at Route 802 (Springs Rd).

The Great Run Watershed bacteria TMDL (Eq ID POL0156) was approved by the EPA on 03/10/2005 (Fed ID 23325). The SWCB approved the TMDL on 12/20/2005. The bacteria TMDL Implementation Plan for the Great Run watershed (ID 160) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_GRT01A00 / Great Run / Segment begins at the confluence with an unnamed tributary to Great Run, approximately 1.0 rivermile upstream of Route 687, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2004	L	2.82
VAN-E02R_GRT02A04 / Great Run / Segment begins at the confluence of an unnamed tributary to Great Run, at approximately rivermile 5.5, and continues downstream until the confluence with an unnamed tributary to Great Run, approximately 1.0 rivermile upstream of Route 687.	4A	Escherichia coli (E. coli)	2004	L	2.85
VAN-E02R_GRT03A02 / Great Run / Segment begins at the confluence with an unnamed tributary to Great Run at rivermile 7.20 (streamcode XAC) and continues downstream until the confluence with another unnamed tributary at approximately rivermile 5.5.	4A	Escherichia coli (E. coli)	2004	L	1.54
VAN-E02R_GRT04A04 / Great Run / Segment begins at the headwaters of Great Run and continues downstream until the confluence with an unnamed tributary to Great Run (streamcode XAC), at rivermile 7.20.	4A	Escherichia coli (E. coli)	2004	L	9.46

Great Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			16.67

Sources: Grazing in Riparian or Shoreline Zones; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E02R-02-BEN Tanner Branch

Cause Location: Begins at the headwaters of Tanner Branch and continues downstream to the confluence with South Run.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2022 at DEQ station 3-TAN002.09 at ~04 miles below Route 733 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

VAN-E02R_TAN01A24 / Tanner Branch / Segment begins at the headwaters of Tanner Branch and EA Benthic Macroinvertebrates	
continues downstream to the confluence with South Run. 5A 5A Bioassessments 2024 L	2.79

Tanner Branch

Aquatic Life	Estuary (Sq. Miles)	$\begin{array}{c} \text{Reservoir} \\ \text{(Acres)} \end{array}$	River (Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			2.79

Sources: Source Unknown

Cause Group Code: E02R-03-BAC Rappahannock River

Cause Location: Begins at the dam at Waterloo (at rivermile 163.4) and continues downstream until the confluence with the Hazel River at rivermile 147.52.

Cause City/County: Culpeper County; Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP150.32.

2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP157.95 at Route 802 (Springs Rd).

2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP163.41 at Route 613 (Waterloo Rd).

A new TMDL is not required for this impaired segment of the Rappahannock River because the downstream Upper Rappahannock River bacteria TMDL (Fed ID 33951, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (2) watershed (Eq ID POL0508).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_RPP01A02 / Rappahannock River / Segment begins at the confluence with Great Run, at rivermile 154.9, and continues downstream until the confluence with the Hazel River, at rivermile 147.52.	4A	Escherichia coli (E. coli)	2006	L	7.05
VAN-E02R_RPP02A04 / Rappahannock River / Segment begins at the confluence with a tributary to the Rappahannock River at rivermile 160.4 and continues downstream until the confluence with Great Run.	4A	Escherichia coli (E. coli)	2020	L	6.24
VAN-E02R_RPP03A04 / Rappahannock River / Segment begins below the dam at Waterloo (rivermile 163.4) and continues downstream until the confluence with a tributary to the Rappahannock River at rivermile 160.4.	4A	Escherichia coli (E. coli)	2016	L	2.99

Rappahannock River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /	× ,	16.28

Cause Group Code: E02R-04-BAC Barrows Run

Cause Location: Begins at the headwaters of Barrows Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2012 Assessment: E. coli bacteria criterion excursions (3 of 4 samples - 75.0%) at DEQ station 3-BRW000.29 at Springs Drive.

A new TMDL is not required for this impaired segment of Barrows Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33951, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (2) watershed (Eq ID POL0508).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_BRW01A06 / Barrows Run / Segment begins at the headwaters of Barrows Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2006	L	4.53

Barrows Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		× /	4.53

Cause Group Code: E02R-05-BAC South Run

Cause Location: Begins at the confluence with Tanner Branch and continues downstream until the confluence with Carter Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-SUT002.62 at Route 737 (Conde Rd).

A new TMDL is not required for this impaired segment of South Run because the downstream Carter Run bacteria TMDL (Fed ID 24414, 03/10/2005) included modeling, source identification, and reductions that covered the entire Carter Run watershed (Eq ID POL0508). A bacteria TMDL Implementation Plan for the Carter Run watershed (ID 99) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_SUT01A04 / South Run / Segment begins at the beginning of the PWS designation and continues downstream until the confluence with Carter Run.	4A	Escherichia coli (E. coli)	2006	L	1.38
VAN-E02R_SUT01B12 / South Run / Segment begins at the confluence with Tanner Branch and continues downstream to the beginning of the PWS designation.	4A	Escherichia coli (E. coli)	2006	L	2.72

South Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli $({\rm E.~coli})$ - Total Impaired Size by Water Ty	ype:	× ,	4.1

Sources: Grazing in Riparian or Shoreline Zones; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E02R-07-BAC Glascock Run

Cause Location: Begins at the headwaters of Glascock Run, and continues downstream to the confluence with Bee Branch.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2016 Assessment: E. coli bacteria criterion excursions (7 of 11 samples - 63.6%) at DEQ station 3-GLC002.03 at Citation Drive.

A new TMDL is not required for this impaired segment of Glascock Run because the downstream Upper Rappahannock River bacteria TMDL (Fed ID 33951, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (2) watershed (Eq ID POL0508).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_GLC02A12 / Glascock Run / Segment begins at the beginning of the PWS designation (five miles upstream from Waterloo) and continues downstream to the confluence with Bee Branch.	4A	Escherichia coli (E. coli)	2012	L	1.44
VAN-E02R_GLC02B12 / Glascock Run / Segment begins at the headwaters of Glascock Run, and continuous downstream to the beginning of the PWS designation (five miles upstream from Waterloo).	4A	Escherichia coli (E. coli)	2012	L	2.38

Glascock Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water T	ype:	· · · · ·	3.82

Cause Group Code: E02R-08-BAC Horner Run

Cause Location: Begins at the perennial headwaters of Horner Run and continues downstream to the confluence with Carter Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (2 of 12 samples - 16.7%) at DEQ station 3-HRN000.80 at Route 691.

A new TMDL is not required for this impaired segment of Horner Run because the downstream Carter Run Watershed bacteria TMDL (Fed ID 24414, 03/10/2005) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0155). The bacteria TMDL Implementation Plan for the Carter Run watershed (ID 99) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E02R_HRN01A20 / Horner Run / Segment begins at the perennial headwaters of Horner Run and continues downstream to the confluence with Carter Run.	4A	Escherichia coli (E. coli)	2020	L	2.36

Horner Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		2.36

Sources: Grazing in Riparian or Shoreline Zones; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E03R-01-BAC Hughes River

Cause Location: Begins at the confluence with Kilbys Creek and continues downstream until the confluence with the Hazel River.

Cause City/County: Culpeper County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-HUE000.20 at Route 644 (Reva Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

The Upper Rappahannock River Watershed bacteria TMDL for the Hughes River watershed (Eq ID POL0512) was approved by the EPA on 01/23/2008 (Fed ID 33916). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Hughes River Run watershed (ID 269) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E03R_HUE01A00 / Hughes River / Segment begins at the confluence with Kilbys Creek and continues downstream until the confluence with the Hazel River.	4A	Escherichia coli (E. coli)	2004	L	3.85

Hughes River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.85

Cause Group Code: E03R-01-BEN Popham Run

Cause Location: Begins at the confluence with Ragged Run and continues downstream until the confluence with the Hughes River.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2022 Assessment: A total of three biological monitoring events in 2015 and 2016 at DEQ station 3-POH000.48 at Route 603 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E03R_POH01A02 / Popham Run / Segment begins at the confluence with Ragged Run and continues downstream until the confluence with the Hughes River.	5A	Benthic Macroinvertebrates Bioassessments	2018	L	2.22

Popham Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			2.22

Sources: Source Unknown

Cause Group Code: E03R-01-TEMP Hughes River

Cause Location: Begins at the upper crossing of Route 707 near the confluence of Rocky Run and continues downstream until the crossing of Route 231.

Cause City/County: Madison County; Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Temperature/5A

Cause Description: Excursions greater than the maximum temperature criterion for stockable trout waters (2 of 10 samples - 20%) at DEQ station 3-HUE007.31 at Route 707 (Sharp Rock Rd).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAN-E03R_HUE02A02 / Hughes River / Segment begins at the upper crossing of Route 707 near the confluence of Rocky Run and continues downstream until the crossing of Route 231.	5A	Temperature		2008	L	3.21
Hughes River Aquatic Life Temperature - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles)	Rese (Ac		River (Miles) 3.21

Sources: Source Unknown

Cause Group Code: E03R-02-BAC Popham Run

Cause Location: Begins at the confluence with Ragged Run and continues downstream until the confluence with the Hughes River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-POH000.48 at Route 603 (Hughes River Rd).

A new TMDL is not required for this impaired segment of Popham Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33916, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hughes River watershed (Eq ID POL0512). A bacteria TMDL Implementation Plan for the Hughes River watershed (ID 269) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E03R_POH01A02 / Popham Run / Segment begins at the confluence with Ragged Run and continues downstream until the confluence with the Hughes River.	4A	Escherichia coli (E. coli)	2012	L	2.22

Popham Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:		`	2.22

Cause Group Code: E03R-02-BEN Kilbys Creek

Cause Location: Begins at the confluence with an unnamed tributary at Route 231 and continues downstream to the confluence with Hughes River.

Cause City/County: Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2022 at DEQ station 3-KIL001.73 at 05 miles below Route 231 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water
VAN-E03R_KIL01A24 / Kilbys Creek / Segment begins at the confluence with an unnamed tributary at Route 231 and continues downstream to the confluence with Hughes River.	5A	Benthic Macroin Bioassessments	vertebrates	2024	L	2.36
Kilbys Creek			Estuary	Rose	rvoir	River
Aquatic Life			(Sq. Miles)		res)	(Miles)
Benthic Macroinvertebrates Bioassessments - T	otal Impaire	ed Size by Water				
		Type:				2.36

Sources: Source Unknown

Cause Group Code: E04R-01-BAC Hazel River

Cause Location: Begins at the confluence of an unnamed tributary to Hazel River at rivermile 36.80, approximately 1.6 rivermiles upstream of Route 607, and continues downstream until the confluence with an unnamed tributary to the Hazel River, at rivermile 16.03.

Cause City/County: Culpeper County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-HAZ018.29 at Route 729 (Eggbornsville Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-HAZ032.54 at Route 644.

2020 Assessment: E. coli bacteria criterion excursions (5 of 11 samples - 45.5%) at DEQ station 3-HAZ034.96 at Route 607.

The Upper Rappahannock River Watershed bacteria TMDL for the Hazel River (1) watershed (Eq ID POL0514) was approved by the EPA on 01/23/2008 (Fed ID 33915). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Hazel River watershed (ID 157) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E04R_HAZ01B00 / Hazel River / Segment begins at the confluence with Blackwater Creek and continues downstream until the confluence with an unnamed tributary to the Hazel River, at rivermile 16.03.	4A	Escherichia coli (E. coli)	2002	L	5.78
VAN-E04R_HAZ02A02 / Hazel River / Segment begins at the Route 707 bridge and continues downstream until the confluence with the Hughes River.	4A	Escherichia coli (E. coli)	2022	L	0.83
VAN-E04R_HAZ02B06 / Hazel River / Segment begins at the confluence of an unnamed tributary to Hazel River at rivermile 36.80, approximately 1.6 rivermiles upstream of Route 607, and continues downstream until the Route 707 bridge. DGIF/DWR Class ii water.	4A	Escherichia coli (E. coli)	2016	L	3.64

Hazel River

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		10.25

Cause Group Code: E04R-01-TEMP Hazel River

Cause Location: Begins at the crossing with the Shenandoah National Park boundary and continues downstream until the Route 707 bridge. DGIF/DWR Class ii water.

Cause City/County: Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Temperature/5A

Cause Description: 2020 Assessment: Excursions greater than the maximum temperature criterion for natural trout waters (3 of 11 samples - 27.3%) at DEQ station 3-HAZ034.96 at Route 607.

Excursions greater than the maximum temperature criterion for natural trout waters (7 of 23 samples - 30.4%) at DEQ station 3-HAZ039.26 at Route 618.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E04R_HAZ02B06 / Hazel River / Segment begins at the confluence of an unnamed tributary to Hazel River at rivermile 36.80, approximately 1.6 rivermiles upstream of Route 607, and continues downstream until the Route 707 bridge. DGIF/DWR Class ii water.	5A	Temperature	2016	L	3.64
VAN-E04R_HAZ03A02 / Hazel River / Segment begins at the crossing with the Shenandoah National Park boundary and continues downstream until the confluence to an unnamed tributary to the Hazel River, at rivermile 36.80. DGIF/DWR Class ii water.	5A	Temperature	2018	L	6.78
Hazel River					

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Temperature - Total Impaired Size by Water Type:			10.42

Sources: Source Unknown

Cause Group Code: E04R-02-BAC Blackwater Creek

Cause Location: Headwaters of Blackwater Creek, downstream to the confluence with the Hazel River.

Cause City/County: Culpeper County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2014 Assessment: E. coli bacteria criterion excursions (4 of 12 samples - 33.3%) at DEQ station 3-BLC001.08 at Route 615.

A new TMDL is not required for this impaired segment of Blackwater Creek because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33915, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (1) watershed (Eq ID POL0514). A bacteria TMDL Implementation Plan for the Hazel River watershed (ID 157) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E04R_BLC01A10 / Blackwater Creek / Headwaters of Blackwater Creek to the confluence with the Hazel River	4A	Escherichia coli (E. coli)	2010	L	8.98

-		•	Reservoir	
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			8.98

Cause Group Code: E04R-03-BAC Hazel River

Cause Location: Begins at the confluence with an unnamed tributary to Hazel River at rivermile 16.03 and continues downstream to the confluence with Thornton River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 12 samples - 25.0%) at DEQ station 3-HAZ013.23 at Route 640.

A new TMDL is not required for this impaired segment of Hazel River because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entre Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Hazel River watershed (ID 157) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E04R_HAZ01A20 / Hazel River / Segment begins at the confluence with an unnamed tributary to Hazel River at rivermile 16.03 and continues downstream to the confluence with Thornton River.	4A	Escherichia coli (E. coli)	2020	L	2.97

Hazel River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
Esche	aerichia coli (E. coli) - Total Impaired Size by Water Type:	. – ,	. ,	2.97

Cause Group Code: E05R-01-BAC Rush River

Cause Location: Begins at the confluence with unnamed tributary at approximately rivermile 7.12 and continues downstream until the confluence with Big Branch, approximately 0.98 rivermile upstream of Route 621.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RUS005.24 at Route 626 (Tiger Valley Rd).

There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RUS007.41 at Route 624 (Sunnyside Rd).

The Upper Rappahannock River Watershed bacteria TMDL for the Rush River watershed (Eq ID POL0513) was approved by the EPA on 01/23/2008 (Fed ID 33914). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Rush River watershed (ID 270) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_RUS02A02 / Rush River / Segment begins at the confluence with unnamed tributary at approximately rivermile 7.12 and continues downstream until the confluence with Big Branch, approximately 0.98 rivermile upstream of Route 621.	4A	Escherichia coli (E. coli)	2002	L	2.78
VAN-E05R_RUS02B18 / Rush River / Segment begins at the confluence with Big Devils Stairs, at rivermile 10.2, and continues downstream until the confluence with an unnamed tributary at approximately rivermile 7.12.	4A	Escherichia coli (E. coli)	2002	L	3.09

Recreation		Estuary (Sq. Miles)	Reservoir		
neereation		(bq. mines)	(meres)	× /	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			5.87	

Sources: Livestock (Grazing or Feeding Operations); On-site Treatment Systems (Septic Systems and Similar Decentralized Systems); Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Puch Divor

Cause Group Code: E05R-01-BEN Thornton River

Cause Location: Begins at the Sperryville Main Street crossing and continues downstream until the confluence with the North Fork Thornton River.

Cause City/County: Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring event in 2022 at DEQ station 3-THO022.27 (above the confluence with NF Thornton River) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_THO03A02 / Thornton River / Segment begins at the Sperryville Main Street crossing and continues downstream until the confluence with the North Fork Thornton River.	5A	Benthic Macroinvertebrates Bioassessments	2010	L	0.87

Thornton River

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			0.87

Sources: Source Unknown

Cause Group Code: E05R-02-BAC Thornton River

Cause Location: Begins at rivermile 25.7 on the Thornton River, where the Class VI designation ends, and continues downstream until the Sperryville Main Street crossing. Begins again at the confluence with White Walnut Run, approximately 0.8 rivermile downstream of Route 621, and continues downstream to the confluence with the Rush River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-THO024.12 at Route 667 (Woodward Rd).

Insufficient E. coli data were available from DEQ station 3-THO014.37 at Route 626 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: E. coli bacteria criterion excursions (5 of 32 samples - 15.6%).

A new TMDL is not required for this impairment of Thornton River because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_THO01A02 / Thornton River / Segment begins at the confluence with White Walnut Run, approximately 0.8 rivermile downstream of Route 621, and continues downstream to the confluence with the Rush River.	4A	Escherichia coli (E. coli)	2006	L	3.45

Thornton River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.45

Cause Group Code: E05R-02-BEN Rush River

Cause Location: Begins at the confluence with unnamed tributary at approximately rivermile 7.12 and continues downstream until the confluence with Big Branch, approximately 0.98 rivermile upstream of Route 621.

Cause City/County: Rappahannock County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of six biological monitoring events in 2017, 2019, and 2022 at DEQ station 3-RUS006.49 at Route 628 (Fodderstack Road) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_RUS02A02 / Rush River / Segment begins at the confluence with unnamed tributary at approximately rivermile 7.12 and continues downstream until the confluence with Big Branch, approximately 0.98 rivermile upstream of Route 621.	5A	Benthic Macroinvertebrates Bioassessments	2022	L	2.78

Rush River

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			2.78

Sources: Source Unknown

Cause Group Code: E05R-03-BAC Big Branch

Cause Location: Segment begins at the headwaters of Big Branch and continues downstream until the confluence with the Rush River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (6 of 11 samples - 54.5%) at DEQ station 3-BIG001.15 at Route 211.

A new TMDL is not required for this impaired segment of Big Branch because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_BIG01A08 / Big Branch / Segment begins at the headwaters of Big Branch and continues downstream until the confluence with the Rush River.	4A	Escherichia coli (E. coli)	2010	L	3.05

Big Branch

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia con (E. con) - Total impaned Size by Water Type.			3.05

Cause Group Code: E05R-04-BAC Rush River

Cause Location: Begins at the confluence with Big Branch and continues downstream until the confluence with the Covington River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RUS003.23 at Route 621 (Hunters Rd).

A new TMDL is not required for this impaired segment of the Rush River because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_RUS01B08 / Rush River / Segment begins at the confluence with Big Branch and continues downstream until the confluence with the Covington River.	4A	Escherichia coli (E. coli)	2024	L	3.36

Rush River

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		3.36

Cause Group Code: E05R-05-BAC Thornton River, North Fork

Cause Location: Begins at the confluence with the Piney River and continues downstream until the confluence with the main stem of the Thornton River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples and the geomean was exceeded in at least one 90-day period with 10+ samples at DEQ station 3-THR000.50 at Route 211 / 522 (Lee Hwy).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_THR01A02 / Thornton River, North Fork / Segment begins at the confluence with the Piney River and continues downstream until the confluence with the main stem of the Thornton River.	4A	Escherichia coli (E. coli)	2022	L	1.94

Thornton River, North Fork

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.94

Cause Group Code: E05R-06-BAC Covington River

Cause Location: Begins at the headwaters of Covington River and continues downstream until the confluence with the Rush River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-COV001.95 at Route 621 (Hunters Rd) and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-COV005.20 at Route 622.

This impairment is nested in the downstream Upper Rappahannock River Watershed bacteria TMDL for the Hazel River (2) watershed and a bacteria TMDL Implementation Plan for the Thornton River watershed is complete.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E05R_COV01A02 / Covington River / Segment begins at the headwaters of Covington River and continues downstream until the confluence with the Rush River.	4A	Escherichia coli (E. coli)	2024	L	7.38

Covington River			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			7.38

Cause Group Code: E06R-01-BAC Thornton River

Cause Location: Begins at the confluence with Mill Run, at rivermile 8.65, and continues downstream until the confluence with an unnamed tributary to the Thornton River, at rivermile 3.25.

Cause City/County: Culpeper County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-THO006.50 at Route 729 (Richmond Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples and a geomean exceedance in at least one 90-day period with 10+ samples.

A new TMDL is not required for this impaired segment of the Thornton River because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E06R_THO02A02 / Thornton River / Segment begins at the confluence with Mill Run and continues downstream until the confluence with unnamed tributary to Thornton River 3-XHH.	4A	Escherichia coli (E. coli)	2006	L	5.52

Thornton River

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type	e:		5.52

Cause Group Code: E06R-02-BAC Battle Run

Cause Location: Begins at the confluence with an unnamed tributary to Battle Run, at rivermile 2.27, and continues downstream until the confluence with the Thornton River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BTL000.94 at Route 729 (Richmond Rd).

A new TMDL is not required for this impaired segment of Battle Run because the downstream Upper Rappahannock River Watershed TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E06R_BTL01A02 / Battle Run / Segment begins at the confluence with an unnamed tributary to Battle Run, at rivermile 2.27, and continues downstream until the confluence with the Thornton River.	4A	Escherichia coli (E. coli)	2008	L	2.24

Battle Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	
	Loonorionia con (Li con) Total impared size sj trater 19por			

Cause Group Code: E06R-03-BAC Unnamed tributary to Thornton River

Cause Location: Begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with the Thornton River.

Cause City/County: Culpeper County; Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2016 Assessment: E. coli bacteria criterion excursions (4 of 12 samples - 33.3%) at DEQ station 3-XHH000.24 at Route 626.

A new TMDL is not required for this impaired segment of the unnamed tributary to Thornton River because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E06R_XHH01A12 / Unnamed tributary to Thornton River / Segment begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with the Thornton River.	4A	Escherichia coli (E. coli)	2012	L	5.02

Unnamed tributary to Thornton River

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E06R-04-BAC Mill Run

Cause Location: Begins at the perennial headwaters at Route 618/658 and continues downstream to the confluence with Thornton River.

Cause City/County: Rappahannock County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MLR001.11 at Route 618.

A new TMDL is not required for this impaired segment of Mill Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517). A bacteria TMDL Implementation Plan for the Thornton River watershed (ID 205) was approved by the EPA on 08/02/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E06R_MLR01A18 / Mill Run / Segment begins at the perennial headwaters at Route 618/658 and continues downstream to the confluence with Thornton River.	4A	Escherichia coli (E. coli)	2020	L	5.9

Mill Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)		
			0.0	

Cause Group Code: E07R-01-BAC Muddy Run

Cause Location: Begins at the headwaters of Muddy Run and continues downstream until the confluence with the Hazel River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MUU000.82 at Route 625.

2018 Assessment: E. coli bacteria criterion excursions (7 of 11 samples - 63.6%) at DEQ station 3-MUU008.52 at Route 632.

The Muddy Run bacteria TMDL (Eq IDs 1299 and POL0003) was approved by the EPA on 07/06/2004 (Fed ID 23326). The SWCB approved the TMDL on 12/02/2004.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E07R_MUU01A00 / Muddy Run / Segment begins at the confluence with an unnamed tributary to Muddy Run, approximately 0.2 rivermile upstream of Route 229, and continues downstream until the confluence with the Hazel River.	4A	Escherichia coli (E. coli)	1996	L	6.09
VAN-E07R_MUU02A02 / Muddy Run / Segment begins at the headwaters of Muddy Run and continues downstream until the confluence with an unnamed tributary to Muddy Run, approximately 0.2 rivermile upstream of Route 229.	4A	Escherichia coli (E. coli)	2002	L	8.25

Muddy Run

Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)		
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Sources: Grazing in Riparian or Shoreline Zones; Impacts from Land Application of Wastes; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E07R-02-BAC Hazel River

Cause Location: Begins at the confluence with Indian Run and continues downstream until the confluence with Muddy Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-HAZ005.98 at Route 625 (Ryland Chapel Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

The Upper Rappahannock River Watershed bacteria TMDL for the Hazel River (2) watershed (Eq ID POL0517) was approved by the EPA on 01/23/2008 (Fed ID 33917). The SWCB approved the TMDL on 07/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E07R_HAZ01A04 / Hazel River / Segment begins at the confluence with Indian Run and continues downstream until the confluence with Muddy Run.	4A	Escherichia coli (E. coli)	2006	L	3.36

Hazel River

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	× - /		3.36

Cause Group Code: E07R-03-BAC Indian Run

Cause Location: Begins at the confluence with an unnamed tributary to Indian Run, upstream from Route 626, and continues downstream until the confluence with the Hazel River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-IND001.14 at Route 624 (Oak Shade Rd).

A new TMDL is not required for this impaired segment of Indian Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entire Hazel River (2) watershed (Eq ID POL0517).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E07R_IND01A04 / Indian Run / Segment begins at the confluence with an unnamed tributary to Indian Run, upstream from Route 626, and continues downstream until the confluence with the Hazel River.	4A	Escherichia coli (E. coli)	2020	L	3.84

Indian Run				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.84

Cause Group Code: E07R-04-BAC Waterford Run

Cause Location: Begins at the headwaters of Waterford Run and continues downstream to the confluence with the Hazel River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-WAF000.82 (upstream of private bridge off Route 611).

A new TMDL is not required for this impaired segment of Indian Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33917, 01/23/2008) included modeling, source identification, and reductions that covered the entre Hazel River (2) watershed (Eq ID POL0517).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E07R_WAF01A10 / Waterford Run / Segment begins at the headwaters of Waterford Run and continues downstream to the confluence with the Hazel River.	4A	Escherichia coli (E. coli)	2020	L	6.23

Waterford Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			6.23

Cause Group Code: E08R-01-BAC Marsh Run

Cause Location: Begins at the headwaters of Marsh Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-MAH000.19 at Route 651 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: Excursions from the maximum E. coli bacteria criterion (10 of 21 samples - 47.6%).

Insufficient E. coli data were available from DEQ station 3-MAH004.18 at Route 668 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: Excursions from the maximum E. coli bacteria criterion (7 of 15 samples - 46.7%).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MAH008.88 at Route 17.

The Upper Rappahannock River Watershed bacteria TMDL for the Marsh Run watershed (Eq ID POL0515) was approved by the EPA on 01/23/2008 (Fed ID 34088). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Marsh Run watershed (ID 18) was approved by the EPA on 05/24/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_MAH01A00 / Marsh Run / Segment begins at the confluence with Harpers Run, at approximately rivermile 2.4, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	1996	L	2.32
VAN-E08R_MAH02A02 / Marsh Run / Segment begins at the confluence with Craig Run and continues downstream until the confluence with Harpers Run, at approximately rivermile 2.4.	4A	Escherichia coli (E. coli)	2012	L	6.01
VAN-E08R_MAH03A02 / Marsh Run / Segment begins at the headwaters of Marsh Run and continues downstream until the confluence with Craig Run.	4A	Escherichia coli (E. coli)	2008	L	3.87

Marsh Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	× /
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			12.2

Cause Group Code: E08R-01-BEN Marsh Run

Cause Location: Begins at the confluence with Craig Run and continues downstream until the confluence with Harpers Run, at approximately rivermile 2.4.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2022 at DEQ station 3-MAH004.18 at Route 668 (Savannah Branch Road) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_MAH02A02 / Marsh Run / Segment begins at the confluence with Craig Run and continues downstream until the confluence with Harpers Run, at approximately rivermile 2.4.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	6.01

Marsh Run

Aquatic Life	Estuary (Sq. Miles)	Reservoir (Acres)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water	(64. 11165)	(110105)	(111105)
Type:			6.01

Sources: Source Unknown

Cause Group Code: E08R-02-BAC Browns Run

Cause Location: Begins at the confluence with an unnamed tributary to Browns Run, near the Route 17 bridge, and continues downstream until the confluence with Marsh Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-BOS000.72 at Route 653 (Morganburg Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

The Upper Rappahannock River Watershed bacteria TMDL for the Browns Run watershed (Eq ID POL0510) was approved by the EPA on 01/23/2008 (Fed ID 33911). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Browns Run watershed (ID 17) was approved by the EPA on 05/24/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_BOS01A02 / Browns Run / Segment begins at the confluence with an unnamed tributary to Browns Run, near the Route 17 bridge, and continues downstream until the confluence with Marsh Run.	4A	Escherichia coli (E. coli)	2002	L	2.55

Browns Run

	Estuary	$\operatorname{Reservoir}$	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		2.55

Cause Group Code: E08R-03-BAC Craig Run

Cause Location: Begins at the headwaters of Craig Run and continues downstream until the confluence with Marsh Run.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-CRA000.46 at Luck Stone Rd for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

The Upper Rappahannock River Watershed bacteria TMDL for the Craig Run watershed (Eq ID POL0509) was approved by the EPA on 01/23/2008 (Fed ID 33912). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Craig Run watershed (ID 116) was approved by the EPA on 05/24/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_CRA01A02 / Craig Run / Segment begins at the headwaters of Craig Run and continues downstream until the confluence with Marsh Run.	4A	Escherichia coli (E. coli)	2004	L	3.72

Craig Run

		Estuary	Reservoir	River
Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(Sq. Miles)	(Acres)	$\frac{\text{(Miles)}}{3.72}$
	Ebenerienia con (E. con) Totar imparied size sy trater Type.			0.12

Cause Group Code: E08R-04-BAC Rappahannock River

Cause Location: Begins at the confluence with Ruffans Run and continues downstream until the confluence with Tinpot Run.

Cause City/County: Culpeper County; Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP147.49 at Route 29 (James Madison Hwy).

The Upper Rappahannock River Watershed bacteria TMDL for the Rappahannock River (2) watershed (Eq ID POL0508) was approved by the EPA on 01/23/2008 (Fed ID 33951). The SWCB approved the TMDL on 07/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_RPP02A02 / Rappahannock River / Segment begins at the confluence with Ruffans Run and continues downstream until the confluence with Tinpot Run.	4A	Escherichia coli (E. coli)	2004	L	2.11

Rappahannock River			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, _ , ,		2.11

Cause Group Code: E08R-05-BAC Rappahannock River

Cause Location: Begins at the confluence with an unnamed tributary to the Rappahannock River, at approximately rivermile 142.5, and continues downstream until the confluence with Marsh Run.

Cause City/County: Culpeper County; Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 assessment: The STV exceedance rate was greater than 10% in at least one 90-day period with 10+ samples, the geomean was exceeded in at least one 90-day period with 10+ samples, and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP142.36 at Route 620.

The Upper Rappahannock River Watershed bacteria TMDL for the Rappahannock River (3) watershed (Eq ID POL0511) was approved by the EPA on 01/23/2008 (Fed ID 33952). The SWCB approved the TMDL on 07/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_RPP01A02 / Rappahannock River / Segment begins at the confluence with an unnamed tributary to the Rappahannock River, at approximately rivermile 142.5, and continues downstream until the confluence with Marsh Run.	4A	Escherichia coli (E. coli)	2006	L	2.86

Rappahannock River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	. – ,	, , ,	2.86

Cause Group Code: E08R-06-BAC Tinpot Run

Cause Location: Begins at the confluence with an unnamed tributary to Tinpot Run, at rivermile 1.27, and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (4 of 10 samples - 40.0%) at DEQ station 3-TIN000.36 at Route 651.

A new TMDL is not required for this impaired segment of Tinpot Run because the downstream Upper Rappahannock River Watershed bacteria TMDL (Fed ID 33952, 01/23/2008) included modeling, source identification, and reductions that covered the entire Rappahannock River (3) watershed (Eq ID POL0511).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E08R_TIN01A08 / Tinpot Run / Segment begins at the confluence with an unnamed tributary to Tinpot Run, at rivermile 1.27, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2008	L	1.29

Tinpot Run			
	Estuary	$\operatorname{Reservoir}$	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaire	d Size by Water Type:		1.29

Cause Group Code: E09L-01-BAC Mountain Run Reservoir

Cause Location: Segment includes all of Mountain Run Reservoir.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: Excursions from the maximum criterion for E.coli were recorded in the pooled data for DEQ monitoring stations 3-MTN028.68 and 3-MTN029.08 (2 of 14 samples -14.3%).

A new TMDL is not required for this impairment because its is located with the watershed addressed by the Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001; Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09L_MTN02A02 / Mountain Run Reservoir / Segment includes all of Mountain Run Reservoir.	4A	Escherichia coli (E. coli)	2020	L	72.76

Mountain Run Reservoir

Recreation		Estuary (Sq. Miles)		
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		72.76	

Sources: Non-Point Source

Cause Group Code: E09L-01-CHLA Mountain Run Reservoir

Cause Location: Includes all of Mountain Run Reservoir.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Chlorophyll-a/5A

Cause Description: Chlorophyll a is assessed as impaired based on two complete monitoring years (2021 and 2022); the criterion was not met in each year.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09L_MTN02A02 / Mountain Run Reservoir / Segment includes all of Mountain Run Reservoir.	5A	Chlorophyll-a	2024	L	72.76

Mountain Run Reservoir				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Chlorophyll-a - Total Impaired Size by Water Type:		72.76	

Cause Group Code: E09L-01-DO Lake Pelham

Cause Location: Includes all of Lake Pelham.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: Excursions less than the minimum dissolved oxygen criterion (9 of 81 samples - 11.1%) at pooled lake stations 3-MTN024.05 and 3-MTN025.04.

The DO impairment may be attributable to seasonal conditions as most of the excursions occurred in non-stratified water during fall months.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09L_MTN01A02 / Lake Pelham / Segment includes all of Lake Pelham.	$5\mathrm{C}$	Dissolved Oxygen	2024	L	249.7
Lake Pelham		_	-		
Aquatic Life		Estuary (Sq. Mile			River Miles)

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Source Unknown

249.7

Dissolved Oxygen - Total Impaired Size by Water Type:

Cause Group Code: E09L-01-TP Lake Pelham

Cause Location: Includes all of Lake Pelham.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Lake Pelham

Causes(s)/VA Category: Phosphorus, Total/5A

Cause Description: Total phosphorus is assessed as impaired based on two complete monitoring years (2021 and 2022; algaecides applied); the criterion was not met in each year.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAN-E09L_MTN01A02 / Lake Pelham / Segment includes all of Lake Pelham.	$5\mathrm{A}$	Phosphorus, Total	2024	L	249.7

Lake I chiam		Fetuery	Reservoir	River	
		Dotuary	Itesei von	TUVEI	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	Phosphorus, Total - Total Impaired Size by Water Type:		249.7		

Cause Group Code: E09L-02-CHLA Lake Pelham

Cause Location: Includes all of Lake Pelham.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Chlorophyll-a/5A

Cause Description: Chlorophyll a is assessed as impaired based on two complete monitoring years (2021 and 2022); the criterion was not met in each year.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09L_MTN01A02 / Lake Pelham / Segment includes all of Lake Pelham.	$5\mathrm{A}$	Chlorophyll-a	2024	L	249.7

Lake Pelham				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Chlorophyll-a - Total Impaired Size by Water Type:		249.7	

Cause Group Code: E09L-02-DO Mountain Run Reservoir

Cause Location: Includes all of Mountain Run Reservoir.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: Excursions less than the minimum dissolved oxygen criterion (10 of 70 samples - 14.3%) at pooled lake stations 3-MTN028.68 and 3-MTN029.08.

The DO impairment may be attributable to seasonal conditions as most of the excursions occurred in non-stratified water during fall months.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09L_MTN02A02 / Mountain Run Reservoir / Segment includes all of Mountain Run Reservoir.	$5\mathrm{C}$	Dissolved Oxygen	2024	L	72.76

Mountain Run Reservoir

Aquatic Life		Estuary (Sq. Miles)		
	Dissolved Oxygen - Total Impaired Size by Water Type:	. – ,	72.76	

Cause Group Code: E09R-01-BAC Mountain Run

Cause Location: Begins at the confluence with Flat Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-MTN000.59 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MTN000.59 at Route 620 and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MTN000.59 at DEQ station 3-MTN005.79 at Route 672.

A bacteria TMDL for the Mountain Run watershed (Eq ID POL0116) was approved by the EPA on 04/27/2001 (Fed ID 24415). The SWCB approved the TMDL on 06/17/2004.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN01A00 / Mountain Run / Segment begins at the confluence with Flat Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	1996	L	7.59

Mountain Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			7.59

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E09R-01-BEN Mountain Run

Cause Location: Begins at the Route 15/29 bridge crossing and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: Six biological monitoring events in 2018, 2019, and 2022 at DEQ station 3-MTN000.59 at Route 620; four biological monitoring events in 2018 and 2019 at DEQ station 3-MTN005.79 at Route 672; one biological monitoring event in 2021 at DEQ station 3-MTN008.49 at 2.7 mile upstream from Route 672; four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN014.88 at Route 663; and four monitoring events in 2018 and 2019 at DEQ station 3-MTN021.11 at Route 799 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN01A00 / Mountain Run / Segment begins at the confluence with Flat Run and continues downstream until the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2008	L	7.59
VAN-E09R_MTN02A04 / Mountain Run / Segment begins at the confluence with Jonas Run and continues downstream until the confluence with Flat Run.	5A	Benthic Macroinvertebrates Bioassessments	2008	L	5.67
VAN-E09R_MTN03A00 / Mountain Run / Segment begins at the Route 15/29 bridge crossing and continues downstream until the confluence with Jonas Run.	5A	Benthic Macroinvertebrates Bioassessments	2008	L	6.65
VAN-E09R_MTN04A04 / Mountain Run / Segment begins at the outlet from Lake Pelham and continues downstream until the Route 15/29 bridge crossing.	5A	Benthic Macroinvertebrates Bioassessments	2020	L	4.63

Mountain Run Aquatic Life Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type: Estuary (Sq. Miles) (Acres) (Miles) 24.54

Sources: Source Unknown

Cause Group Code: E09R-01-PCB Mountain Run

Cause Location: Begins at the Route 15/29 bridge crossing near Culpeper City and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Culpeper County

Use(s): Aquatic Life; Fish Consumption; Wildlife

Causes(s)/VA Category: PCBs in Fish Tissue/5A; Polychlorinated biphenyls (PCBs)/5A

Cause Description: The fish consumption use is categorized as impaired due to a Virginia Department of Health PCB fish consumption advisory. The advisory, dated 12/13/04, limits American eel consumption to no more than two meals per month. The affected stretch of Mountain Run extends roughly 19 miles, from the Route 15/29 bridge crossing near Culpeper City downstream until the confluence with the Rappahannock River.

The following exceedances of the human health criterion of 640 picogram per liter (pg/l) for total PCBs in the water column were recorded: The median of two water samples collected at DEQ station 3-MTN005.79 at Route 672; the median of two water samples collected at DEQ station 3-MTN000.59 at Route 620; the median of three water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN010.98 at Route 669; the median of two water samples collected at DEQ station 3-MTN014.88 at Route 663

2020 Assessment: The following exceedances of the water quality criterion based fish tissue value (TV) of 18 parts per billion (ppb) for PCBs in fish tissue were recorded: four exceedances in two species of fish (American eel and yellow bullhead catfish) collected in 2013 at DEQ station 3-MTN000.59; two exceedances in two species of fish (American eel and yellow bullhead catfish) collected in 2013 at DEQ station 3-MTN005.79; and four exceedances in three species of fish (American eel, yellow bullhead catfish, and sunfish) collected in 2013 at DEQ station 3-MTN014.33.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN01A00 / Mountain Run / Segment begins at the confluence with Flat Run and continues downstream until the confluence with the Rappahannock River.	5A	PCBs in Fish Tissue	2006	Н	7.59
VAN-E09R_MTN02A04 / Mountain Run / Segment begins at the confluence with Jonas Run and continues downstream until the confluence with Flat Run.	5A	PCBs in Fish Tissue	2006	Н	5.67
VAN-E09R_MTN03A00 / Mountain Run / Segment begins at the Route 15/29 bridge crossing and continues downstream until the confluence with Jonas Run.	5A	PCBs in Fish Tissue	2006	Н	6.65

Mountain Run

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
PCBs in Fish Tissue - Total Impaired Size by Water Type:			19.91

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN01A00 / Mountain Run / Segment begins at the confluence with Flat Run and continues downstream until the confluence with the Rappahannock River.	5A	Polychlorinated biphenyls (PCBs)	2020	Н	7.59
VAN-E09R_MTN02A04 / Mountain Run / Segment begins at the confluence with Jonas Run and continues downstream until the confluence with Flat Run.	5A	Polychlorinated biphenyls (PCBs)	2020	Н	5.67
VAN-E09R_MTN03A00 / Mountain Run / Segment begins at the Route 15/29 bridge crossing and continues downstream until the confluence with Jonas Run.	5A	Polychlorinated biphenyls (PCBs)	2018	Н	6.65

Mountain Run

Aquatic Life Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles) 19.91
Mountain Run			
Fish Consumption Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles) 19.91
Mountain Run			
Wildlife Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles) 19.91

Cause Group Code: E09R-02-BAC Mountain Run

Cause Location: Segment begins at the outlet from Lake Pelham and continues downstream until the confluence with Flat Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MTN014.88 at Route 663 (Stevensburg Rd).

2020 Assessment: E. coli bacteria criterion excursions (2 of 10 samples - 20%) at DEQ station 3-MTN021.11 at Route 799.

2020 Assessment: E. coli bacteria criterion excursions (3 of 4 samples - 75.0%) at DEQ station 3-MTN022.01 at Old Brandy Road.

A new TMDL is not required for this impaired segment because the downstream Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN02A04 / Mountain Run / Segment begins at the confluence with Jonas Run and continues downstream until the confluence with Flat Run.	4A	Escherichia coli (E. coli)	2022	L	5.67
VAN-E09R_MTN03A00 / Mountain Run / Segment begins at the Route 15/29 bridge crossing and continues downstream until the confluence with Jonas Run.	4A	Escherichia coli (E. coli)	2010	L	6.65
VAN-E09R_MTN04A04 / Mountain Run / Segment begins at the outlet from Lake Pelham and continues downstream until the Route 15/29 bridge crossing.	4A	Escherichia coli (E. coli)	2016	L	4.63

Mountain Run

		Estuary	$\operatorname{Reservoir}$	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		16.95	

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E09R-02-BEN Jonas Run

Cause Location: Begins at the confluence with an unnamed tributary to Jonas Run (XDZ), at approximately rivermile 3.74, and continues downstream until the confluence with Mountain Run.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of four monitoring events in 2019 and 2020 at DEQ station 3-JOA000.80 at Route 663 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_JOA01A06 / Jonas Run / Segment begins at the confluence with an unnamed tributary to Jonas Run (XDZ), at approximately rivermile 3.74, and continues downstream until the confluence with Mountain Run.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	3.78

Jonas Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			3.78

Cause Group Code: E09R-02-PCB Mountain Run

Cause Location: Begins at the outlet from Lake Pelham and continues downstream until the Route 15/29 bridge crossing.

Cause City/County: Culpeper County

Use(s): Fish Consumption

Causes(s)/VA Category: PCBs in Fish Tissue/5A; Polychlorinated biphenyls (PCBs)/5A

Cause Description: 2020 Assessment: Exceedances of the water quality criterion based fish tissue value (TV) of 18 parts per billion (ppb) for polychlorinated biphenyls (PCBs) in fish tissue were recorded in three species of fish (white sucker, American eel, and yellow bullhead catfish) in four total samples collected in 2013 at DEQ station 3-MTN022.21.

2020 Assessment: Two exceedances of the human health criteria of 640 picogram per liter (pg/l) for total polychlorinated biphenyls (PCBs) in the water column were recorded in samples collected at DEQ station 3-MTN021.11 at Route 799.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN04A04 / Mountain Run / Segment begins at the outlet from Lake Pelham and continues downstream until the Route 15/29 bridge crossing.	5A	PCBs in Fish Tissue	2016	Н	4.63

Mountain Run

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
PCBs in Fish Tissue - Total Impaired Size by Water Type:			4.63

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN04A04 / Mountain Run / Segment begins at the outlet from Lake Pelham and continues downstream until the Route 15/29 bridge crossing.	5A	Polychlorinated biphenyls (PCBs)	2018	Н	4.63

Mountain Run

	Estuary	Reservoir	River	
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)	
Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	· - /		4.63	

Cause Group Code: E09R-03-BAC Mountain Run

Cause Location: Begins at the confluence with an unnamed tributary that flows from Caymore Lake and continues downstream until Lake Pelham.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (4 of 18 samples - 22.2%) at DEQ station 3-MTN027.08 at Route 641.

A new TMDL is not required for this impaired segment of Mountain Run because the downstream Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_MTN05A04 / Mountain Run / Segment begins at the confluence with an unnamed tributary that flows from Caymore Lake and continues downstream until Lake Pelham.	4A	Escherichia coli (E. coli)	2006	L	1.63

Mountain Run

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		1.63

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E09R-03-BEN Unnamed tributary to Jonas Run

Cause Location: Begins at the confluence with an unnamed tributary (downstream from Swan Dam) and continues downstream to the confluence with Jonas Run.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2018 at DEQ station 3-XMO000.41 (0.02 mile downstream from Route 685) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_XMO01A20 / Unnamed tributary to Jonas Run / Segment begins at the confluence with an unnamed tributary (downstream from Swan Dam) and continues downstream to the confluence with Jonas Run.	5A	Benthic Macroinvertebrates Bioassessments	2020	L	0.53

Unnamed tributary to Jonas Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			0.53

Cause Group Code: E09R-03-PCB Unnamed tributaries to Mountain Run

Cause Location: Unnamed tributaries 3-XBE and 3-XIH, from their perennial headwaters downstream to their confluences with Mountain Run.

Cause City/County: Culpeper County

Use(s): Aquatic Life; Fish Consumption; Wildlife

Causes(s)/VA Category: Polychlorinated biphenyls (PCBs)/5A

Cause Description: The median of two samples collected in 2018 and 2021 at DEQ station 3-XBE000.19 at Yancey Street exceeds the human health criterion for total polychlorinated biphenyls (PCBs) in the water column.

The median of two samples collected in 2018 at DEQ station 3-XIH000.06 at the end of Spring Street exceeds the human health criterion for total polychlorinated biphenyls (PCBs) in the water column

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_XBE01A18 / Unnamed tributary to Mountain Run / Segment begins at the perennial headwaters and continues downstream to the confluence with Mountain Run.	$5\mathrm{A}$	Polychlorinated biphenyls (PCBs)	2020	Н	0.60
VAN-E09R_XIH01A18 / Unnamed tributary to Mountain Run / Segment begins at the perennial headwaters and continues downstream to the confluence with Mountain Run.	$5\mathrm{A}$	Polychlorinated biphenyls (PCBs)	2020	Н	1.12

Unnamed tributaries to Mountain Run

Aquatic Life Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles) 1.72	
Unnamed tributaries to Mountain Run		р	D:	
Fish Consumption				
Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	(5q. miles)	(ACIES)	(1.72)	
Unnamed tributaries to Mountain Run				
	Estuary	Reservoir	River	
Wildlife Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type:	(Sq. Miles)	(Acres)	(Miles) 1.72	
 Fish Consumption Polychlorinated biphenyls (PCBs) - Total Impaired Size by Water Type: Unnamed tributaries to Mountain Run Wildlife 	Estuary (Sq. Miles) Estuary (Sq. Miles)	Reservoir (Acres) Reservoir (Acres)	River (Miles)	

Cause Group Code: E09R-04-BAC Jonas Run

Cause Location: Begins at the confluence with an unnamed tributary to Jonas Run (XDZ), at approximately rivermile 3.74, and continues downstream until the confluence with Mountain Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 7 samples - 42.9%) at DEQ station 3-JOA000.80 at Route 663 (Stevensburg Road).

A new TMDL is not required for this impaired segment of Jonas Run because the downstream Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_JOA01A06 / Jonas Run / Segment begins at the confluence with an unnamed tributary to Jonas Run (XDZ), at approximately rivermile 3.74, and continues downstream until the confluence with Mountain Run.	4A	Escherichia coli (E. coli)	2008	L	3.78

Jonas Run				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.78

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E09R-04-BEN Flat Run

Cause Location: Begins at the headwaters of Flat Run and continues downstream until the confluence with Mountain Run.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of four monitoring events in 2019 and 2020 at station 3-FLA001.93 at Route 675 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_FLA01A08 / Flat Run / Segment begins at the headwaters of Flat Run and continues downstream until the confluence with Mountain Run.	5A	Benthic Macroinvertebrates Bioassessments	2022	L	6.23
Run.					

Flat Run

Aquatic Life	Estuary (Sq. Miles)	$\begin{array}{c} \text{Reservoir} \\ \text{(Acres)} \end{array}$	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			6.23

Cause Group Code: E09R-05-BAC Flat Run

Cause Location: Begins at the headwaters of Flat Run and continues downstream until the confluence with Mountain Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 7 samples - 42.9%) at DEQ station 3-FLA001.93 at Route 675.

A new TMDL is not required for this impaired segment of Flat Run because the downstream Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_FLA01A08 / Flat Run / Segment begins at the headwaters of Flat Run and continues downstream until the confluence with Mountain Run.	4A	Escherichia coli (E. coli)	2014	L	6.23

Flat Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			6.23

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E09R-06-BAC Unnamed tributary to Jonas Run

Cause Location: Begins at the confluence with an unnamed tributary (downstream from Swan Dam) and continues downstream to the confluence with Jonas Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-XMO000.44 at Route 685.

A new TMDL is not required for this impaired segment of an unnamed tributary to Jonas Run because the downstream Mountain Run bacteria TMDL (Fed ID 24415, 04/27/2001) included modeling, source identification, and reductions that covered the entre watershed (Eq ID POL0116).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E09R_XMO01A20 / Unnamed tributary to Jonas Run / Segment begins at the confluence with an unnamed tributary (downstream from Swan Dam) and continues downstream to the confluence with Jonas Run.	4A	Escherichia coli (E. coli)	2020	L	0.53

Unnamed tri	butary to Jonas Run				
		Estuary	Reservoir	River	
Recreation	L	(Sq. Miles)	(Acres)	(Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		0.53	

Sources: Grazing in Riparian or Shoreline Zones; Impervious Surface/Parking Lot Runoff; Livestock (Grazing or Feeding Operations); Manure Runoff; Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E10R-01-BAC Deep Run

Cause Location: Begins at the headwaters of Deep Run and continues downstream until the confluence with Pine Branch. Begins again at the confluence with Green Branch (at rivermile 4.75) and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fauquier County; Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-DPR001.70 at Route 17 (Marsh Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-DPR008.98 at Route 634 (Elk Run Church Rd).

The Deep Run bacteria TMDL (Eq ID POL0115) was approved by the EPA on 05/26/2004 (Fed ID 24417). The SWCB approved the TMDL on 08/31/2004. A bacteria TMDL Implementation Plan for the Deep Run watershed (ID 58) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E10R_DPR01A00 / Deep Run / Segment begins at the confluence with Green Branch, at rivermile 4.75, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	1996	L	4.93
VAN-E10R_DPR03A02 / Deep Run / Segment begins at the headwaters of Deep Run and continues downstream until the confluence with Pine Branch.	4A	Escherichia coli (E. coli)	2014	L	3.75

Deep Run

		Estuary	Reservoir	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		8.68	

Sources: Grazing in Riparian or Shoreline Zones; Impacts from Land Application of Wastes; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E10R-01-BEN Sumerduck Run

Cause Location: Begins at the confluence with an unnamed tributary to Sumerduck Run, approximately 0.55 rivermile upstream of Route 632, and continues downstream until the confluence with another unnamed tributary, at Route 631.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2020 Assessment: A total of four biological monitoring events in 2013 and 2014 at DEQ station 3-SMR004.81 at Route 632 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community (2020 Assessment).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E10R_SMR02A06 / Sumerduck Run / Segment begins at the confluence with an unnamed tributary to Sumerduck Run, approximately 0.55 rivermile upstream of Route 632, and continues downstream until the confluence with another unnamed tributary, at Route 631.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	1.86

Sumerduck Run				
	Estuary	Reservoir	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			1.86	

Cause Group Code: E10R-02-BEN Unnamed tributary to Pine Branch

Cause Location: Begins at the perennial headwaters and continues downstream to the confluence with Pine Branch.

Cause City/County: Fauquier County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2021 at DEQ station 3-XMT000.25, approximately 0.25 mile above the confluence with Pine Branch, resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E10R_XMT01A24 / Unnamed tributary to Pine Branch / Segment begins at the perennial headwaters and continues downstream to the confluence with Pine Branch.	$5\mathrm{A}$	Benthic Macroinvertebrates Bioassessments	2024	L	1.97

Unnamed tributary to Pine Branch

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			1.97

Cause Group Code: E10R-03-BAC Alcotti Run

Cause Location: Begins at the headwaters of Alcotti Run and continues downstream until the confluence with Deep Run.

Cause City/County: Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-ALC000.45 at Hazel Lane.

A new TMDL is not required for this impaired segment of Alcotti Run because the downstream Deep Run bacteria TMDL (Fed ID 24417, 05/26/2004) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0115). A bacteria TMDL Implementation Plan for the Deep Run watershed (ID 58) was approved by the EPA on 05/22/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E10R_ALC01A00 / Alcotti Run / Segment begins at the headwaters of Alcotti Run and continues downstream until the confluence with Deep Run.	4A	Escherichia coli (E. coli)	2012	L	5.17

Alcotti Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	,		5.17

Sources: Grazing in Riparian or Shoreline Zones; Impacts from Land Application of Wastes; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E10R-04-BAC Sumerduck Run

Cause Location: Begins at the confluence with an unnamed tributary to Sumerduck Run, approximately 0.55 rivermile upstream of Route 632, and continues downstream until the confluence with another unnamed tributary, at Route 631.

Cause City/County: Fauquier County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/5A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-SMR004.81 at Route 632 (Union Church Rd).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E10R_SMR02A06 / Sumerduck Run / Segment begins at the confluence with an unnamed tributary to Sumerduck Run, approximately 0.55 rivermile upstream of Route 632, and continues downstream until the confluence with another unnamed tributary, at Route 631.	5A	Escherichia coli (E. coli)	2016	L	1.86

 ${\rm Sumerduck}\; {\rm Run}$

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.86

Cause Group Code: E11R-01-BAC Garth Run

Cause Location: Begins at the headwaters of Garth Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GAR000.95 at Route 718 (Wallace Gap Ln).

Insufficient E. coli data were available from DEQ station 3-GAR005.59 at Route 615 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

A new TMDL is not required for this impaired segment of Garth Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Garth Run watershed (ID 78) was approved by the EPA on 12/31/2015

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E11R_GAR01A02 / Garth Run / Segment begins at the Route 665 crossing, at approximately rivermile 1.9, and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2014	L	1.61
VAN-E11R_GAR02A06 / Garth Run / Segment begins at the headwaters of Garth Run and continues downstream until the Route 665 crossing, at approximately rivermile 1.9. DGIF/DWR Class iii water.	4A	Escherichia coli (E. coli)	2018	L	5.82

Garth Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E11R-01-BEN Conway River

Cause Location: Segment begins at the confluence with an unnamed tributary to the Conway River, approximately 0.6 rivermile upstream from Route 230, and continues downstream until the confluence with the Rapidan River.

Cause City/County: Greene County; Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2014 Assessment: A total of three biological monitoring events in 2007 and 2008 at DEQ station 3-CON002.26 at Route 230 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E11R_CON01A04 / Conway River / Segment begins at the beginning of the PWS designation, and continues downstream until the confluence with the Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2010	L	0.33
VAN-E11R_CON01B12 / Conway River / Segment begins at the confluence with an unnamed tributary to the Conway River, approximately 0.6 rivermile upstream from Route 230, and continues downstream until the start of the PWS designated area.	5A	Benthic Macroinvertebrates Bioassessments	2010	L	2.67

Conway River

	Estuary	Reservoir	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			3	

Cause Group Code: E12R-01-BAC Rapidan River

Cause Location: Begins at the confluence with the Conway River and continues downstream until the confluence with Rippin Run.

Cause City/County: Greene County; Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-RAP066.54 at Route 29 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

A new TMDL is not required for this impaired segment of Rapidan River because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E12R_RAP01A00 / Rapidan River / Segment begins at the end of the public water supply designation area, approximately 0.43 rivermiles upstream from the Route 29 crossing, and continues downstream until the confluence with Rippin Run.	4A	Escherichia coli (E. coli)	2006	L	2.34
VAN-E12R_RAP01B06 / Rapidan River / Segment begins at the confluence with the Conway River and continues downstream until the end of the public water supply designation area, approximately 0.43 rivermiles upstream from the Route 29 crossing.	4A	Escherichia coli (E. coli)	2006	L	4.93

Rapidan River

	Estuary	$\operatorname{Reservoir}$	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			7.27	

Cause Group Code: E12R-01-BEN Rippin Run

Cause Location: Begins at the confluence with White Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Greene County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2016 Assessment: A total of two biological monitoring events in 2010 at DEQ station 3-RIP000.22 at Route 609 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E12R_RIP01A04 / Rippin Run / Segment begins at the confluence with White Run and continues downstream until the confluence with the Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	0.6
Rippin Run			D		

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water	, _ ,	, í	, í
Type:			0.6

Cause Group Code: E12R-02-BAC Rippin Run

Cause Location: Begins at the confluence with White Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Greene County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RIP000.22 at Route 609 (Fredericksburg Rd).

A new TMDL is not required for this impaired segment of Rippin Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Rippin Run watershed (ID 72) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E12R_RIP01A04 / Rippin Run / Segment begins at the confluence with White Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2012	L	0.6

Rippin Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			0.6

Cause Group Code: E12R-02-BEN Buckner Run

Cause Location: Begins at the perennial headwaters and continues downstream to the confluence with Rapidan River.

Cause City/County: Greene County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2021 at DEQ station 3-BUK001.15 at 0.23 miles downstream from Route 609 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E12R_BUK01A24 / Buckner Run / Segment begins at the perennial headwaters and continues downstream to the confluence with Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2024	L	3.21

Buckner Run

Aquatic Life	Estuary (Sq. Miles)	$\begin{array}{c} \text{Reservoir} \\ \text{(Acres)} \end{array}$	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			3.21

Cause Group Code: E12R-03-BAC South River

Cause Location: Begins at the confluence with Henshaw Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Greene County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-SOT001.00 at Route 619 (Dundee Rd).

A new TMDL is not required for this impaired segment of South River because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Rapidan River #2 watershed (ID 73) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E12R_SOT01A04 / South River / Segment begins at the confluence with Henshaw Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2020	L	1.67

South River

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	,	, , ,	1.67

Cause Group Code: E13R-01-BAC Blue Run

Cause Location: Begins at the headwaters of Blue Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Albemarle County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-BLU000.80 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BLU002.60 at Route 20.

2020 Assessment: E. coli bacteria criterion excursions (7 of 11 samples - 63.6%) at DEQ station 3-BLU008.33 at Route 33.

The Rapidan River Basin bacteria TMDL for the Blue Run watershed (Eq ID POL0494) was approved by the EPA on 12/05/2007 (Fed ID 33865). The SWCB approved the TMDL on 07/31/2008. The Upper Rapidan River bacteria TMDL Implementation Plan for the Blue Run watershed (ID 77) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_BLU01A00 / Blue Run / Segment begins at the beginning of the PWS designation and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2002	L	0.15
VAN-E13R_BLU01B12 / Blue Run / Segment begins at the confluence with Barbour Run, approximately 0.13 rivermile upstream of the Southern Rail Road bridge, and continues downstream until the start of the PWS designation.	4A	Escherichia coli (E. coli)	2002	L	4.20
VAN-E13R_BLU02A04 / Blue Run / Segment begins at the headwaters of Blue Run and continues downstream until the confluence with Barbour Run.	4A	Escherichia coli (E. coli)	2006	L	8.38

Blue Run			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	/	. ,	12.73

Cause Group Code: E13R-01-BEN Beautiful Run

Cause Location: Begins at an unnamed tributary at rivermile 3.44, and continues downstream to another unnamed tributary, upstream of Route 620.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2018 Assessment: A total of two biological monitoring events in 2011 at DEQ station 3-BFL002.90 at Route 616 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water
VAN-E13R_BFL02A12 / Beautiful Run / Segment begins at an unnamed tributary at rivermile 3.44, and continues downstream to another unnamed tributary, upstream of Route 620.	5A	Benthic Macroin Bioassessments	vertebrates	2012	L	2.51
Beautiful Run			Estuary (Sq. Miles)		ervoir res)	River (Miles)
Benthic Macroinvertebrates Bioassessments - T	Total Impaire	ed Size by Water	(~9. 11100)	(110		(

Type:

2.51

Cause Group Code: E13R-03-BAC Unnamed tributary to Beautiful Run

Cause Location: Begins at the perennial headwaters and continues downstream to the confluence with Beautiful Run.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 5 samples - 60.0%) at DEQ station 3-XMM001.33 at Route 231.

A new TMDL is not required for this impaired segment of an unnamed tributary to Beautiful Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Beautiful Run watershed (ID 69) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_XMM01A20 / Unnamed tributary to Beautiful Run / Segment begins at the perennial headwaters and continues downstream to the confluence with Beautiful Run.	4A	Escherichia coli (E. coli)	2020	L	2.02

Unnamed tributary to Beautiful Run Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type: Escherichia coli (E. coli) - Total Impaired Size

Cause Group Code: E13R-04-BAC Unnamed tributary to the Rapidan River

Cause Location: Begins at the headwaters of the unnamed tributary and continues downstream until the confluence with the Rapidan River.

Cause City/County: Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (2 of 11 samples - 18.2%) at DEQ station 3-XEZ000.12 at Route 634.

The Rapidan River Basin bacteria TMDL for this Unnamed Tributary to the Rapidan River watershed (Eq ID POL0497) was approved by the EPA on 12/05/2007 (Fed ID 33866). The SWCB approved the TMDL on 07/31/2008. The Upper Rapidan River bacteria TMDL Implementation Plan for the Rapidan River #1 watershed (ID 76) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_XEZ01A04 / Unnamed tributary to Rapidan River / Segment begins at the headwaters of the unnamed tributary and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2004	L	2.67

Unnamed tributary to the Rapidan River				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		2.67	

Cause Group Code: E13R-05-BAC Beautiful Run

Cause Location: Begins at the headwaters and continues downstream until the confluence with the Rapidan River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BFL000.90 at Route 620 (Tatums School Rd).

2020 Assessment: E. coli bacteria criterion excursions (4 of 12 samples - 33.3%) at DEQ station 3-BFL002.90 at Route 616.

There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BFL006.28 at Route 621 (Beautiful Run Rd).

A new TMDL is not required for this impaired segment of Beautiful Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Beautiful Run watershed (ID 69) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_BFL01A04 / Beautiful Run / Segment begins at the confluence of an unnamed tributary, upstream from Route 620, and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2006	L	1.18
VAN-E13R_BFL02A12 / Beautiful Run / Segment begins at an unnamed tributary at rivermile 3.44, and continues downstream to another unnamed tributary, upstream of Route 620.	4A	Escherichia coli (E. coli)	2012	L	2.51
VAN-E13R_BFL03A16 / Beautiful Run / Segment begins at the headwaters of Beautiful Run and continues downstream to an unnamed tributary at rivermile 3.44.	4A	Escherichia coli (E. coli)	2016	L	8.46

Beautiful Rui	1				
		Estuary	Reservoir	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			12.15	

Cause Group Code: E13R-06-BAC Rapidan River

Cause Location: Begins at the confluence with Marsh Run and continues downstream until the confluence with Blue Run.

Cause City/County: Madison County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-RAP055.84 at Route 231 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: Excursions from the maximum E. coli bacteria criterion (7 of 12 samples - 58.3%).

A new TMDL is not required for this impaired segment of Rapidan River because the downstream Rapidan River Basin TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Rapidan River #2 watershed (ID 73) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_RAP02A06 / Rapidan River / Segment begins at the beginning of the PWS designation. and continues downstream until the confluence with Blue Run.	4A	Escherichia coli (E. coli)	2006	L	0.15
VAN-E13R_RAP02B12 / Rapidan River / Segment begins at the confluence with Marsh Run and continues downstream until the start of the PWS designation.	4A	Escherichia coli (E. coli)	2006	L	4.19

Rapidan River

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Tota	npaired Size by Water Type:		4.34

Cause Group Code: E13R-07-BAC Unnamed tributary to Rapidan River

Cause Location: Begins at the headwaters of the unnamed tributary and continues downstream until the confluence with the Rapidan River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (8 of 12 samples - 66.7%) at DEQ station 3-XBO000.26 at Route 621.

A new TMDL is not required for this impaired segment of the unnamed tributary to the Rapidan River because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the UT to Rapidan River #2 watershed (ID 75) was approved by the EPA on 12/31/2015

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_XBO01A04 / Unnamed tributary to Rapidan River / Segment begins at the headwaters of the unnamed tributary and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2006	L	3.11

Unnamed tributary to Rapidan River

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E13R-08-BAC Marsh Run

Cause Location: Begins at the headwaters of Marsh Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Greene County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MAS000.62 at Route 609.

The Rapidan River Basin bacteria TMDL for the Marsh Run watershed (Eq ID POL0495) was approved by the EPA on 12/05/2007 (Fed ID 33864). The SWCB approved the TMDL on 07/31/2008. The Upper Rapidan River bacteria TMDL Implementation Plan for the Marsh Run watershed (ID 74) was approved by the EPA on 12/31/2015.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_MAS01A04 / Marsh Run / Segment begins at the headwaters of Marsh Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2014	L	5.65

Marsh Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Typ	e:	× ,	5.65

Cause Group Code: E13R-09-BAC Poplar Run

Cause Location: Begins at the headwaters of Poplar Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-POL000.10 at Route 633 (Amicus Rd).

A new TMDL is not required for this impaired segment of Poplar Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33867, 12/05/2007) included modeling, source identification, and reductions that covered the entire Upper Rapidan River watershed (Eq ID POL0496). The Upper Rapidan River bacteria TMDL Implementation Plan for the Poplar Run watershed (ID 71) was approved by the EPA on 12/31/2015

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E13R_POL01A04 / Poplar Run / Segment begins at the headwaters of Poplar Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2014	L	4.15

Poplar Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, _	, , ,	4.15

Cause Group Code: E14R-01-BEN White Oak Run

Cause Location: Begins approximately 0.4 rivermile upstream from the Route 657 crossing, and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2022 at DEQ station 3-WHO001.51 at just upstream from Route 231 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E14R_WHO01A06 / White Oak Run / Segment begins approximately 0.4 rivermile upstream from the Route 657 crossing, and continues downstream until the confluence with the Robinson River.	5A	Benthic Macroinvertebrates Bioassessments	2018	L	3.19

White Oak Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			3.19

Cause Group Code: E14R-02-BAC Finks Run

Cause Location: Begins at the headwaters of Finks Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-FIK001.08 at Route 650 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2020 assessment: Excursions from the maximum E. coli bacteria criterion (3 of 11 samples - 27.3%).

A new TMDL is not required for this impaired segment of Finks Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 40412, 12/12/2005) included modeling, source identification, and reductions that covered the entire Upper Robinson River watershed (Eq ID POL0245). The Little Dark Run and Robinson River bacteria TMDL Implementation Plan for the Upper Robinson River watershed (ID 14) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E14R_FIK01A06 / Finks Run / Segment begins at the headwaters of Finks Run and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2006	L	3.17

Finks Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	/		3.17

Sources: Grazing in Riparian or Shoreline Zones; Impacts from Land Application of Wastes; Livestock (Grazing or Feeding Operations); Runoff from Forest/Grassland/Parkland; Sewage Discharges in Unsewered Areas; Wastes from Pets; Waterfowl; Wildlife Other than Waterfowl

Cause Group Code: E14R-02-TEMP Rose River

Cause Location: Begins at rivermile 2.6, approximately 0.36 rivermile downstream from the confluence with Strother Run, and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Temperature/5A

Cause Description: Excursions greater than the maximum temperature criterion for stockable trout waters (2 of 12 samples - 16.7%) at DEQ station 3-ROE000.75 at a private road.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAN-E14R_ROE01A02 / Rose River / Segment starts at rivermile 2.6, approximately 0.36 rivermile downstream from the confluence with Strother Run, and continues downstream until the confluence with the Robinson River.	5A	Temperature		2006	L	2.58
Rose River Aquatic Life Temperature - Total In	mpaired Size	by Water Type:	Estuary (Sq. Miles)		rvoir res)	River (Miles) 2.58

Cause Group Code: E14R-03-BAC White Oak Run

Cause Location: Begins approximately 0.4 rivermile upstream from the Route 657 crossing, and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-WHO001.48 at Route 231 (Blue Ridge Turnpike).

A new TMDL is not required for this impaired segment of White Oak Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E14R_WHO01A06 / White Oak Run / Segment begins approximately 0.4 rivermile upstream from the Route 657 crossing, and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2006	L	3.19

White Oak Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.19

Cause Group Code: E14R-04-BAC Leathers Run

Cause Location: Begins at the confluence with an unnamed tributary to Leathers Run, approximately 0.65 rivermile downstream from the Route 641 crossing, and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-LEA000.17 at Route 609 (W Hoover Rd).

A new TMDL is not required for this impaired segment of Leathers Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E14R_LEA01A06 / Leathers Run / Segment begins at the confluence with an unnamed tributary to Leathers Run, approximately 0.65 rivermile downstream from the Route 641 crossing, and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2006	L	2.18

Leathers Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)		
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		× ,	2.18	

Cause Group Code: E15R-01-BAC Little Dark Run

Cause Location: Begins at the headwaters of Little Dark Run and continues downstream until the confluence with Dark Run.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-LDR000.70 at Route 680 (Gate Rd).

The Robinson River and Little Dark Run bacteria TMDL for the Little Dark Run watershed (Eq ID POL0244) was approved by the EPA on 12/12/2005 (Fed ID 24418). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Little Dark Run watershed (ID 15) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_LDR01A00 / Little Dark Run / Segment begins at the confluence with an unnamed tributary to Little Dark Run, at rivermile 2.17, and continues downstream until the confluence with Dark Run.	4A	Escherichia coli (E. coli)	1998	L	4.54

Little Dark Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	
	Escherichia con (E. con) - rotar imparted bize by water rype.		4.04

Cause Group Code: E15R-02-BAC Robinson River

Cause Location: Begins at the confluence with Deep Run and continues downstream to the confluence with Beaverdam Run. Begins again at the confluence with Crooked Run, and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County; Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-ROB001.90 at Route 614 (Locust Dale Rd) for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-ROB009.93 at Route 632 (Beahm Town Road).

The Robinson River and Little Dark Run bacteria TMDL for the Lower Robinson River watershed (Eq ID POL0243) was approved by the EPA on 12/12/2005 (Fed ID 24419). The SWCB approved the TMDL on 07/31/2008. A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the U.S. EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_ROB01A00 / Robinson River / Segment begins at the confluence with Crooked Run, and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2004	L	5.32
VAN-E15R_ROB02B18 / Robinson River / Segment begins at the confluence with Deep Run and continues downstream to the confluence with Beaverdam Run.	4A	Escherichia coli (E. coli)	2022	L	8.83

Robinson River

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			14.15

Cause Group Code: E15R-02-BEN Deep Run

Cause Location: Begins at the confluence with Muddy Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2022 Assessment: A total of two biological monitoring events in 2016 at DEQ station 3-DRN001.81 at Route 638 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_DRN01A04 / Deep Run / Segment begins at the confluence with Muddy Run and continues downstream until the confluence with the Robinson River.	5A	Benthic Macroinvertebrates Bioassessments	2018	L	2.48
Deep Run					

Aquatic Life	Estuary (Sq. Miles)	$\begin{array}{c} \text{Reservoir} \\ \text{(Acres)} \end{array}$	$\begin{array}{c} \text{River} \\ \text{(Miles)} \end{array}$
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			2.40
Type:			2.48

Cause Group Code: E15R-03-BAC Deep Run

Cause Location: Begins at the confluence with Muddy Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-DRN001.81 at Route 638 (Hebron Church Rd).

A new TMDL is not required for this impaired segment of Deep Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_DRN01A04 / Deep Run / Segment begins at the confluence with Muddy Run and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2008	L	2.48

Deep Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			2.48

Cause Group Code: E15R-03-BEN Great Run

Cause Location: Begins at the headwaters of Great Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2022 Assessment: A total of two biological monitoring events in 2016 at DEQ station 3-GRA002.01 at Route 15 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_GRA01A04 / Great Run / Segment begins at the headwaters of Great Run and continues downstream until the confluence with the Robinson River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	9.31
Great Run					

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water	, _ ,	, í	`
Type:			9.31

Cause Group Code: E15R-04-BAC Crooked Run

Cause Location: Begins at the confluence with Little Crooked Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Culpeper County; Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-COO005.66 at Route 618 (Waylands Mill Rd).

A new TMDL is not required for this impaired segment of Crooked Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_COO01A04 / Crooked Run / Segment begins at the confluence with Little Crooked Run and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2008	L	7.89

Crooked Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
E	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, _ ,		7.89

Cause Group Code: E15R-04-BEN Little Dark Run

Cause Location: Begins at the headwaters of Little Dark Run and continues downstream until the confluence with Dark Run.

Cause City/County: Madison County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of five biological monitoring events in 2017, 2019, and 2021 at DEQ station 3-LDR000.70 at Route 680 (Gate Road) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_LDR01A00 / Little Dark Run / Segment begins at the confluence with an unnamed tributary to Little Dark Run, at rivermile 2.17, and continues downstream until the confluence with Dark Run.	5A	Benthic Macroinvertebrates Bioassessments	2020	L	4.54

Little Dark Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			4.54

Cause Group Code: E15R-05-BAC Great Run

Cause Location: Begins at the headwaters of Great Run and continues downstream until the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GRA002.01 at Route 15 (James Madison Hwy).

A new TMDL is not required for this impaired segment of Great Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_GRA01A04 / Great Run / Segment begins at the headwaters of Great Run and continues downstream until the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2008	L	9.31

Great RunEstuaryReservoirRiverRecreationEscherichia coli (E. coli) - Total Impaired Size by Water Type:(Sq. Miles)(Acres)(Miles)9.31

Cause Group Code: E15R-06-BAC Dark Run

Cause Location: Begins at the headwaters of Dark Run and continues to the confluence with the Robinson River.

Cause City/County: Madison County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-DAK001.18 at Route 634 (Oak Park Rd).

A new TMDL is not required for this impaired segment of Dark Run because the downstream Robinson River and Little Dark Run bacteria TMDL (Fed ID 24419, 12/12/2005) included modeling, source identification, and reductions that covered the entire Lower Robinson River watershed (Eq ID POL0243). A bacteria TMDL Implementation Plan for the Lower Robinson River watershed (ID 119) was approved by the EPA on 05/31/2011.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E15R_DAK01A10 / Dark Run / Segment begins at the headwaters of Dark Run and continues to the confluence with the Robinson River.	4A	Escherichia coli (E. coli)	2010	L	8.59

Dark Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			8.59

Cause Group Code: E16R-01-BAC Cedar Run

Cause Location: Begins at the confluence with Cabin Branch and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-CED000.59 at Route 522 (Zachary Taylor Hwy).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-CED003.52 at Route 652 (Mitchell Rd).

The Rapidan River Basin bacteria TMDL for the Cedar Run watershed (Eq ID POL0493) was approved by the EPA on 12/05/2007 (Fed ID 33868). The SWCB approved the TMDL on 07/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E16R_CED01A00 / Cedar Run / Segment begins at the confluence with Cabin Branch and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2018	L	2.26
VAN-E16R_CED02A04 / Cedar Run / Segment begins at the confluence with Buck Run and continues downstream until the confluence with Cabin Branch.	4A	Escherichia coli (E. coli)	2006	L	3.54

Cedar Run

		Estuary	Reservoir	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
Escheric	iia coli (E. coli) - Total Impaired Size by Water Type:			5.8	

Cause Group Code: E16R-01-BEN Cedar Run

Cause Location: Begins at the confluence with Cabin Branch and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2022 at DEQ station 3-CED000.59 at Route 522 (Zachary Taylor Highway) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E16R_CED01A00 / Cedar Run / Segment begins at the confluence with Cabin Branch and continues downstream until the confluence with the Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2018	L	2.26

Cedar Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			2.26

Cause Group Code: E16R-02-BAC Rapidan River

Cause Location: Begins at the confluence with an unnamed tributary to the Rapidan River, at rivermile 34.5, approximately 0.6 rivermile downstream from Route 689, and continues downstream until the confluence with Cedar Run.

Cause City/County: Culpeper County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ/USGS station 3-RAP030.21 at Route 522 (Zachary Taylor Hwy).

A new TMDL is not required for this impaired segment of the Rapidan River because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E16R_RAP01A04 / Rapidan River / Segment begins at the confluence with an unnamed tributary to the Rapidan River, at rivermile 34.5, approximately 0.6 rivermile downstream from Route 689, and continues downstream until the confluence with Cedar Run.	4A	Escherichia coli (E. coli)	2006	L	4.66

Rapidan River

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		4.66

Cause Group Code: E16R-03-BAC Rapidan River

Cause Location: Begins at the confluence with the Robinson River and continues downstream until the confluence with an unnamed tributary to the Rapidan River, at rivermile 36.6.

Cause City/County: Culpeper County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 12 samples - 25.0%) at DEQ station 3-RAP037.90 at Route 615.

A new TMDL is not required for this impaired segment of the Rapidan River because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E16R_RAP03A08 / Rapidan River / Segment begins at the confluence with the Robinson River and continues downstream until the confluence with an unnamed tributary to the Rapidan River, at rivermile 36.6.	4A	Escherichia coli (E. coli)	2008	L	3.4

Rapidan Rive	r			
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.4

Cause Group Code: E16R-04-BAC Cabin Branch

Cause Location: Begins at the perennial headwaters of Cabin Branch and continues downstream to the confluence with Cedar Run.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-CAB000.22 at Route 655 (Summerville Rd) and there were two or more STV exceedances in at least one 90-day period with <10 samples DEQ station 3-CAB002.23 at Route 615 (Rapidan Rd).

A new TMDL is not required for this impaired segment of Cabin Branch because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33868, 12/05/2007) included modeling, source identification, and reductions that covered the entre Cedar Run watershed (Eq ID POL0493).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E16R_CAB01A22 / Cabin Branch / Segment begins at the perennial headwaters of Cabin Branch and continues downstream to the confluence with Cedar Run.	4A	Escherichia coli (E. coli)	2022	L	3.19

Cabin Branch

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.19

Cause Group Code: E17R-01-BAC Mine Run

Cause Location: Begins at the confluence with Cormack Run, approximately 0.6 rivermile upstream of Route 20, and continues downstream until the confluence with the Rapidan River.

Cause City/County: Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (2 of 11 samples - 18.2%) at DEQ station 3-MIR004.05 at Route 611.

The Mountain Run and Mine Run bacteria TMDL for the Mine Run watershed (Eq ID POL0242) was approved by the EPA on 11/15/2005 (Fed ID 24420). The SWCB approved the TMDL on 09/27/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_MIR01A00 / Mine Run / Segment begins at the confluence with Cormack Run, approximately 0.6 rivermile upstream of Route 20, and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2002	L	10.5

Mine Run

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, – , ,	10.5

Cause Group Code: E17R-01-BEN Brook Run

Cause Location: Begins at the confluence with an unnamed tributary to Brook Run, at Route 647, and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of two biological monitoring events in 2018 at DEQ station 3-BRK002.52 at 0.1 mile downstream of Route 647 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_BRK01A04 / Brook Run / Segment begins at the confluence with an unnamed tributary to Brook Run. at Route 647, and continues downstream until the confluence with the Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	2.51

Brook Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			2.51

Cause Group Code: E17R-02-BAC Mountain Run

Cause Location: Begins at the headwaters of Mountain Run and continues downstream until the confluence with Mine Run.

Cause City/County: Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-MTR003.51 at Route 611 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples.

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MTR008.31 at Route 621 (Pine Stake Rd).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MTR010.60 at Route 666 (Hawfield Rd).

The Mountain Run and Mine Run bacteria TMDL for the Mountain Run watershed (Eq ID POL0241) was approved by the EPA on 11/15/2005 (Fed ID 24421). The SWCB approved the TMDL on 09/27/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_MTR01A00 / Mountain Run / Segment begins at the confluence with Mill Run, approximately 0.25 rivermile downstream of Route 617, and continues downstream until the confluence with Mine Run.	4A	Escherichia coli (E. coli)	2002	L	10.11
VAN-E17R_MTR02A02 / Mountain Run / Segment begins at the headwaters of Mountain Run and continues downstream until the confluence with Mill Run.	4A	Escherichia coli (E. coli)	2006	L	7.46

Mountain Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			17.57

Cause Group Code: E17R-02-BEN Mountain Run

Cause Location: Begins at the confluence with Mill Run, approximately 0.25 rivermile downstream of Route 617, and continues downstream until the confluence with Mine Run.

Cause City/County: Orange County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of four biological monitoring events in 2016 and 2017 at DEQ station 3-MTR003.51 at Route 611 resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_MTR01A00 / Mountain Run / Segment begins at the confluence with Mill Run, approximately 0.25 rivermile downstream of Route 617, and continues downstream until the confluence with Mine Run.	5A	Benthic Macroinvertebrates Bioassessments	2018	L	10.11

Mountain Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			10.11

Cause Group Code: E17R-03-BAC Black Walnut Run

Cause Location: Begins at the Route 621 crossing and continues downstream until the confluence with Mine Run.

Cause City/County: Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-BWR004.13 at Route 602 (Old Office Rd).

A new TMDL is not required for this impaired segment of Black Walnut Run because the downstream Mountain Run and Mine Run bacteria TMDL (Fed ID 24420, 11/15/2005) included modeling, source identification, and reductions that covered the entire Mine Run watershed (Eq ID POL0242).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_BWR01A06 / Black Walnut Run / Segment begins at the Route 621 crossing and continues downstream until the confluence with Mine Run.	4A	Escherichia coli (E. coli)	2006	L	6.48

Black Walnut Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			6.48

Cause Group Code: E17R-04-BAC Sumerduck Run

Cause Location: Begins at the confluence with Dry Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (7 of 10 samples - 70.0%) at DEQ station 3-SUM002.40 at Route 647 (Twin Mountain Road).

A new TMDL is not required for this impaired segment of Sumerduck Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_SUM01A04 / Sumerduck Run / Segment begins at the confluence with Dry Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2014	L	6.21

Sumerduck Run

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		6.21

Cause Group Code: E17R-05-BAC Potato Run

Cause Location: Begins at the headwaters of Potato Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (4 of 8 samples - 50.0%) at DEQ station 3-POT001.06 at Route 647 (Twin Mountain Road).

A new TMDL is not required for this impaired segment of Potato Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_POT01A14 / Potato Run / Segment begins at the headwaters of Potato Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2014	L	6.84

Potato Run

Pognostion		Estuary		
Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(Sq. Miles)	(Acres)	(Miles) 6.84

Cause Group Code: E17R-06-BAC Brook Run

Cause Location: Begins at the confluence with an unnamed tributary to Brook Run. at Route 647, and continues downstream until the confluence with the Rapidan River

Cause City/County: Culpeper County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples DEQ station 3-BRK002.64 at Route 647 (Batna Rd).

A new TMDL is not required for this impaired segment of Brook Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E17R_BRK01A04 / Brook Run / Segment begins at the confluence with an unnamed tributary to Brook Run. at Route 647, and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2018	L	2.51

Brook Run			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Typ	e:		2.51

Cause Group Code: E18R-01-BEN Russell Run

Cause Location: Begins at the headwaters of Russell Run and continues downstream to the confluence with the Rapidan River.

Cause City/County: Orange County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of eight biological monitoring events in 2017, 2019, 2020, 2021, and 2022 at DEQ station 3-RUL000.39 at Route 603 (Indiantown Road) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E18R_RUL01A14 / Russell Run / Segment begins at the headwaters of Russell Run and continues downstream to the confluence with the Rapidan River.	5A	Benthic Macroinvertebrates Bioassessments	2024	L	4.73

Russell Run

Aquatic Life	Estuary (Sq. Miles)	Reservoir (Acres)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			4.73

Cause Group Code: E18R-01-HG Rapidan River

Cause Location: Begins at the confluence with Flat Run and continues downstream to the confluence with the Rappahannock River.

Cause City/County: Culpeper County; Orange County; Spotsylvania County

Use(s): Fish Consumption

Causes(s)/VA Category: Mercury in Fish Tissue/5A

Cause Description: Exceedances of the water quality criterion based fish tissue value (TV) of 300 parts per billion (ppb) for mercury (Hg) in fish tissue were recorded in three species of fish (American eel, rock bass, smallmouth bass) collected in 2006 and in one species of fish (largemouth bass) collected in 2018 at DEQ station 3-RAP006.53.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E18R_RAP01A02 / Rapidan River / Segment begins at the confluence with Hunting Run, at rivermile 1.35, and continues downstream until the confluence with the Rappahannock River.	5A	Mercury in Fish Tissue	2010	L	1.24
VAN-E18R_RAP02A02 / Rapidan River / Segment begins at the confluence with Middle Run, rivermile 5.10, and continues downstream until the confluence with Hunting Run.	5A	Mercury in Fish Tissue	2010	L	3.64
VAN-E18R_RAP03A02 / Rapidan River / Segment begins at the confluence with Wilderness Run, rivermile 7.78, and continues downstream until the confluence with Middle Run.	5A	Mercury in Fish Tissue	2010	L	2.59
VAN-E18R_RAP04A04 / Rapidan River / Segment begins at the confluence with Flat Run and continues downstream until the confluence with Wilderness Run.	5A	Mercury in Fish Tissue	2010	L	2.34

Rapidan I	River
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	Estuary	Reservoir	River	
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)	
Mercury in Fish Tissue - Total Impaired Size by Water Type:			9.81	

Cause Group Code: E18R-02-BAC Wilderness Run

Cause Location: Begins at the confluence of North Wilderness Run and South Wilderness Run and continues downstream until the confluence with the Rapidan River.

Cause City/County: Orange County; Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (7 of 11 samples - 63.6%) at DEQ station 3-WIL004.00 at Route 3.

A new TMDL is not required for this impaired segment of Wilderness Run because the downstream Rapidan River Basin bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E18R_WIL01A08 / Wilderness Run / Segment begins at the confluence of North Wilderness Run and South Wilderness Run and continues downstream until the confluence with the Rapidan River.	4A	Escherichia coli (E. coli)	2008	L	5.56

Wilderness Run				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water	r Type:		5.56	

Cause Group Code: E18R-03-BAC Rapidan River

Cause Location: Begins at the boundary of the public water supply area, approximately 1.21 rivermiles upstream from the Route 3 crossing, and continues downstream until the confluence with Lick Branch.

Cause City/County: Culpeper County; Orange County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (8 of 24 samples - 33.3%) at DEQ station 3-RAP014.45 at Route 3.

A new TMDL is not required for this impaired segment of the Rapidan River because the downstream Rapidan River bacteria TMDL (Fed ID 33869, 12/05/2007) included modeling, source identification, and reductions that covered the entire Lower Rapidan River watershed (Eq ID POL0492).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E18R_RAP05A08 / Rapidan River / Segment begins at the boundary of the public water supply area, approximately 1.17 rivermiles upstream from the Route 3 crossing, and continues downstream to the confluence with Lick Branch.	4A	Escherichia coli (E. coli)	2008	L	3.41

Estuary Reservoir River	
Recreation (Sq. Miles) (Acres) (Mile	;)
Escherichia coli (E. coli) - Total Impaired Size by Water Type: 3.4	1

Cause Group Code: E19L-01-HG Motts Run Reservoir

Cause Location: Includes the entirety of Motts Run Reservoir.

Cause City/County: Spotsylvania County

Use(s): Fish Consumption

Causes(s)/VA Category: Mercury in Fish Tissue/5A

Cause Description: The fish consumption use is categorized as impaired due to a Virginia Department of Health mercury (Hg) fish consumption advisory. The advisory, dated 8/31/07, limits consumption of largemouth bass to no more than two meals per month. The affected area includes the entirety of Motts Run Reservoir.

There were also five total exceedances of the water quality criterion-based tissue value (TV) of 300 parts per billion (ppb) for mercury (Hg) in fish tissue from two species of fish (largemouth bass and bluegill sunfish) sampled in 2017 at DEQ station 3-MOT000.39 and previous exceedances in six total samples of one species of fish (largemouth bass) sampled in 2006 at DEQ station 3-MOT000.39.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E19L_MOT01A02 / Motts Run Reservoir / Segment includes the lower half of Motts Run Reservoir; beginning at rivermile 0.8 and continuing downstream until the lake's discharge.	5A	Mercury in Fish Tissue	2008	L	62.89
VAN-E19L_MOT02A02 / Motts Run Reservoir / Segment includes the upper half of Motts Run Reservoir; beginning at the upper end of the reservoir and continuing downstream until rivermile 0.8.	5A	Mercury in Fish Tissue	2008	L	74.29

Motts Run Reservoir

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
Mercury in Fish Tissue - Total Impaired Size by Water Type:		137.18	

Cause Group Code: E19R-01-BAC Horsepen Run

Cause Location: Begins at headwaters of Horsepen Run and continues downstream to the confluence with the Rappahannock River.

Cause City/County: Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/5A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (4 of 12 samples - 33.3%) at DEQ station 3-HOR000.50 at Route 655 (Holly Corner Road).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E19R_HOR01A04 / Horsepen Run / Segment begins at headwaters of Horsepen Run and continues downstream to the confluence with the Rappahannock River.	5A	Escherichia coli (E. coli)	2014	L	5.71

Horsepen Run
Recreation
Escherichia coli (E. coli) - Total Impaired Size by Water Type:
Escherichia coli (E. coli) - Total Impaired Size by Water Type:

Cause Group Code: E19R-02-BAC Mine Run

Cause Location: Begins at the headwaters of Mine Run and continues downstream to the upper end of the Motts Run Reservoir.

Cause City/County: Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/5A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (6 of 12 samples - 50.0%) at DEQ station 3-MIN002.14 at Route 620 (Spotswood Furnace Road).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E19R_MIN02A14 / Mine Run / Segment begins at the headwaters of Mine Run and continues downstream to the upper end of the Motts Run Reservoir.	5A	Escherichia coli (E. coli)	2014	L	4.01
Mine Run		Estuary	Rese	rvoir Bi	ver

		Estuary	Reservoir	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, – ,	, , ,	4.01	

Cause Group Code: E20E-01-BAC Rappahannock River

Cause Location: Begins at the fall line at Route 1 and continues downstream until the confluence with Massaponax Creek.

Cause City/County: Caroline County; Fredericksburg; King George County; Spotsylvania County; Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples and no geomean exceedances at DEQ station 3-RPP110.57 at Route 1 (Jefferson Davis Hwy).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP106.01 (upstream from the Fredericksburg Country Club).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP098.81 (near Hayfield Bar).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP091.55 near Hop Yard Bar.

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-RPP080.19 at 100 yards downstream from 301 bridge at Port Royal.

The Tidal Rappahannock River Watershed bacteria TMDL (Eq ID POL0569) was approved by the EPA on 05/05/2008 (Fed ID 34369). The SWCB approved the TMDL on 04/28/2009.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20E_RPP01A02 / Rappahannock River / Segment begins at the confluence with Little Falls Run and continues downstream until the outlet of waterbody VAN-E20E. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2006	L	0.285
VAN-E20E_RPP02A02 / Rappahannock River / Segment begins at the confluence with Deep Run and continues downstream until the confluence with Little Falls Run. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2002	L	0.135
VAN-E20E_RPP03A02 / Rappahannock River / Segment begins at the fall line at Route 1 and continues downstream until the confluence with Deep Run. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2002	L	0.195
VAN-E21E_RPP03A02 / Rappahannock River / Segment begins at the confluence with Mount Creek and continues downstream until the confluence with Mill Creek. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2006	L	1.366
VAN-E21E_RPP04A02 / Rappahannock River / Segment begins at the confluence with Ware Creek and continues downstream until the confluence with Mount Creek. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2006	L	1.206
VAN-E21E_RPP05A02 / Rappahannock River / Segment begins at the confluence with Snow Creek (at the start of E21) and continues downstream until the confluence with Ware Creek. Portion of CBP segment RPPTF.	4A	Escherichia coli (E. coli)	2006	L	0.579

Rappahannock River

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	3.766	

Cause Group Code: E20E-03-PCB Rappahannock River

Cause Location: Extends from the I-95 bridge above Fredericksburg downstream to the mouth of the river near Stingray Point, including its tributaries Hazel Run up to the I-95 bridge crossing and Claiborne Run up to the Route 1 bridge crossing.

- Cause City/County: Caroline County; Essex County; Fredericksburg; King George County; Lancaster County; Middlesex County; Richmond County; Spotsylvania County; Stafford County; Westmoreland County
- Use(s): Aquatic Life; Fish Consumption; Wildlife

Causes(s)/VA Category: PCBs in Fish Tissue/5A; Polychlorinated biphenyls (PCBs)/5A

Cause Description: The fish consumption use is categorized as impaired due to a Virginia Department of Health PCB fish consumption advisory. The advisory, dated 12/13/04, limits American eel, blue catfish, carp, channel catfish, croaker, gizzard shad, and anadromous (coastal) striped bass consumption to no more than two meals per month.

The following exceedances of the water quality criterion based tissue value (TV) of 18 parts per billion (ppb) for polychlorinated biphenyls (PCBs) in fish tissue were recorded: two exceedances in two species (blue catfish and gizzard shad) collected in 2018 at DEQ station 3-RPP080.19 on Rappahannock River; four exceedances in three species (blue catfish, largemouth bass, and gizzard shad) collected in 2018 at DEQ station 3-RPP091.55 on Rappahannock River; and three exceedances in three species (blue catfish, carp, and gizzard shad) collected in 2018 at DEQ station 3-RPP107.33 on Rappahannock River. Additionally, the previous exceedances from the 2022IR were carried forward: two exceedances in two samples of American eel collected during the same sample event in 2016 at DEQ station 3-HAL000.57 on Hazel Run; two exceedances in two samples of American eel collected during the same sample event in 2016 at DEQ station 3-HAL000.57 on Claiborne Run.

In addition, during 2018 sampling, there were exceedances in three species at 3-RPP070.00 and two species at 3-RPP042.12.

There was a previous exceedance at 3-RPP056.20; however, re-sampling in 2018 was acceptable. In addition, 2018 sampling at 3-RPP029.40 and 3-RPP008.42, as well as 2020 sampling at 3-RPP001.36 were all acceptable.

The median of two samples collected in 2021 at DEQ station 3-RPP098.81 exceeds the human health criterion for total polychlorinated biphenyls (PCBs) in the water column.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E19R_RPP01A18 / Rappahannock River / Segment begins at the I-95 bridge and continues downstream to the E19/E20 watershed boundary (at the downstream reach of the PWS designation).	5A	PCBs in Fish Tissue	2004	L	0.660
VAN-E20E_RPP01A02 / Rappahannock River / Segment begins at the confluence with Little Falls Run and continues downstream until the outlet of waterbody VAN-E20E. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	0.285
VAN-E20E_RPP02A02 / Rappahannock River / Segment begins at the confluence with Deep Run and continues downstream until the confluence with Little Falls Run. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	0.135
VAN-E20E_RPP03A02 / Rappahannock River / Segment begins at the fall line at Route 1 and continues downstream until the confluence with Deep Run. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	0.195

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_CLB01A00 / Claiborne Run / Segment begins at the Route 1 crossing of Claiborne Run and continues downstream until the confluence with the Rappahannock River.	5A	PCBs in Fish Tissue	2006	L	4.520
VAN-E20R_HAL01A00 / Hazel Run / Segment begins at the Route 95 crossing and continues downstream until the confluence with the Rappahannock River.	5A	PCBs in Fish Tissue	2006	L	4.730
VAN-E20R_RPP01A10 / Rappahannock River / Segment begins at the E19/E20 watershed boundary, and extends downstream to the end of the free flowing waters of the Rappahannock River, at the Route 1/Cambridge Street/Falmouth Bridge.	$5\mathrm{A}$	PCBs in Fish Tissue	2004	L	2.660
VAN-E21E_RPP01A02 / Rappahannock River / Segment begins at the confluence with Mill Creek, at rivermile 78.94, and continues downstream until immediately upstream of Devils Elbow, at rivermile 70.52. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2006	L	4.547
VAN-E21E_RPP03A02 / Rappahannock River / Segment begins at the confluence with Mount Creek and continues downstream until the confluence with Mill Creek. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	1.366
VAN-E21E_RPP04A02 / Rappahannock River / Segment begins at the confluence with Ware Creek and continues downstream until the confluence with Mount Creek. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	1.206
VAN-E21E_RPP05A02 / Rappahannock River / Segment begins at the confluence with Snow Creek (at the start of E21) and continues downstream until the confluence with Ware Creek. Portion of CBP segment RPPTF.	5A	PCBs in Fish Tissue	2004	L	0.579
VAP-E22E_RPP01A02 / Rappahannock River / The Rappahannock River from Devils Elbow at Toby Point and Green Bay (river mile 70.52) downstream to the tidal freshwater/oligohaline boundary at river mile 57.85. RPPTF	5A	PCBs in Fish Tissue	2006	L	5.133
VAP-E22E_RPP02A02 / Rappahannock River / The Rappahannock River from the tidal freshwater/oligohaline boundary downstream to river mile 56.21. RPPOH	5A	PCBs in Fish Tissue	2006	L	1.344
VAP-E22E_RPP02B16 / Rappahannock River / The Rappahannock River from rivermile 56.21 downstream to river mile 51.04. RPPOH	$5\mathrm{A}$	PCBs in Fish Tissue	2006	L	2.003
VAP-E22E_RPP03A02 / Rappahannock River / The Rappahannock River from river mile 51.04 to river mile 49.04. RPPOH	5A	PCBs in Fish Tissue	2006	L	2.012

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP04A02 / Rappahannock River / The Rappahannock River from river mile 49.04 downstream to the oligohaline/mesohaline boundary at approximately river mile 48.51. RPPOH	5A	PCBs in Fish Tissue	2006	L	0.942
VAP-E22E_RPP05A02 / Rappahannock River / The oligohaline/mesohaline boundary at river mile 48.51 to the downstream boundary of VDH shellfish condemnation area 025A-068B, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2006	L	6.958
VAP-E23E_RPP02A98 / Rappahannock River / Mainstem Rappahannock as described in VDH shellfish condemnation 025A-068A, 3/15/2021, excluding administratively condemned portion. RPPMH	$5\mathrm{A}$	PCBs in Fish Tissue	2006	L	8.123
VAP-E23E_RPP02B10 / Rappahannock River / Portion of mainstem Rappahannock River that is administratively condemned within condemnation 025A-068A, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.158
VAP-E23E_RPP02C12 / Rappahannock River / Portion of VDH shellfish condemnation 025A-068A, 11/14/2005 not included in 025A-068A, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.387
VAP-E24E_RPP01B14 / Garrett's Marina / As delineated in VDH shellfish condemnation 026-181A, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2008	L	0.003
VAP-E24E_RPP01B98 / Rappahannock River: Garrett's Marina / As delineated in VDH shellfish condemnation 026-181M1, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2008	L	0.025
VAP-E24E_RPP01C06 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/16/2007 (non-admin) that is open 3/15/2021. Shrank in the 2024 cycle. RPPMH	$5\mathrm{A}$	PCBs in Fish Tissue	2006	L	0.327
VAP-E24E_RPP01D10 / Rappahannock River / The portion of the Rappahannock River within the administratively closed area of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.137
VAP-E24E_RPP01E18 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/15/2021 (non-admin) Expanded in the 2024 cycle. RPPMH	$5\mathrm{A}$	PCBs in Fish Tissue	2006	L	0.378

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(continued)

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP03A00 / Rappahannock River / The Rappahannock River from the limit of VDH shellfish condemnation 068A, 11/14/2005 downstream to end of MSN (Sharps/0.7 mi DS of Mark Haven Beach) unless otherwise segmented RPPMH	5A	PCBs in Fish Tissue	2006	L	10.858
VAP-E24E_RPP03D24 / Rappahannock River / Described in VDH shellfish condemnation 026-270A, 5/25/2022. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.061
VAP-E25E_RPP01C10 / Rappahannock River: Mark Haven Beach Basin / The portion of VDH shellfish condemnation 026-181B, 1/20/2006 not administratively closed. RPPMH	5A	PCBs in Fish Tissue	2008	L	0.010
VAP-E25E_RPP01C98 / Mark Haven Beach Basin / As delineated in VDH shellfish condemnation 026-181A, 4/3/2012. RPPMH	5A	PCBs in Fish Tissue	2008	L	0.004
VAP-E25E_RPP02A02 / Rappahannock River / The mainstem of the Rappahannock River from the end of MSN (Sharps/0.7 mi DS of Mark Haven Beach to the mouth, excluding area in SFC 051A. RPPMH	5A	PCBs in Fish Tissue	2006	L	81.443
VAP-E25E_RPP03A06 / Rappahannock River / Described in VDH Shellfish Condemnation 024-070B, 11/15/2022. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.008
VAP-E25E_RPP03B16 / Rappahannock River Run Bluffs / As described in VDH shellfish condemnation 026-181B, 3/15/2021. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.003
VAP-E26E_RPP04A00 / Rappahannock River / Described in VDH Shellfish Condemnation 029-051B, 9/1/2015. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.131
VAP-E26E_RPP05A00 / Rappahannock River / Delineated in VDH-DSS condemnation 029-051C, 9/1/2015. RPPMH	5A	PCBs in Fish Tissue	2006	L	0.049
VAP-E26E_RPP07A02 / Rappahannock River / As delineated in VDH-DSS condemnation 018-053A, 7/23/2018. RPPMH	5A	PCBs in Fish Tissue	2002	L	0.139

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VAP-E26E_RPP08A22 / Rappahannock River / Portion of VDH-DSS Condemnation 032-109A, 9/15/2022 that is within the mainstem Rappahannock River. Shrank in the 2024 cycle. RPPMH	5A	PCBs in Fish Ti	ssue	2006	L	0.002
Rappahannock River Fish Consumption PCBs in Fish Tissue - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 128.948		ervoir cres)	River (Miles) 12.57
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Prior	Water
VAN-E21E_RPP05A02 / Rappahannock River / Segment begins at the confluence with Snow Creek (at the start of E21) and continues downstream until the confluence with Ware Creek. Portion of CBP segment RPPTF.	5A	Polychlorinated (PCBs)	biphenyls	2024	L	0.579
Rappahannock River Aquatic Life Polychlorinated biphenyls (PCBs) - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 0.579		ervoir cres)	River (Miles)
Rappahannock River Fish Consumption Polychlorinated biphenyls (PCBs) - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 0.579		ervoir cres)	River (Miles)
Rappahannock River Wildlife Polychlorinated biphenyls (PCBs) - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 0.579		ervoir cres)	River (Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E20R-01-BAC Claiborne Run

Cause Location: Begins at the Route 1 crossing of Claiborne Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 12 samples - 25.0%) at DEQ station 3-CLB000.50 at Naomi Road.

A new TMDL is not required for this impaired segment of Claiborne Run because the downstream bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire tidal freshwater Rappahannock River watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_CLB01A00 / Claiborne Run / Segment begins at the Route 1 crossing of Claiborne Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2004	L	4.52

Claiborne Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
Escl	cherichia coli (E. coli) - Total Impaired Size by Water Type:			4.52

Cause Group Code: E20R-01-BEN Falls Run

Cause Location: Begins at the headwaters of Falls Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Stafford County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2016 Assessment: A total of two biological monitoring events in 2009 at DEQ station 3-FAL000.13 at Washington Street resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_FAL01A04 / Falls Run / Segment begins at the headwaters of Falls Run and continues downstream until the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	7.36
Falls Run		Fatuom	Poss	ww.oir Di	iver
Aquatic Life		Estuary (Sq. Miles			Tiles)

	Estuary	Treser von	TUVEL	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			7.36	

Sources: Source Unknown

Cause Group Code: E20R-02-BAC Hazel Run

Cause Location: Begins at the Route 95 crossing and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fredericksburg; Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (2 of 12 samples - 16.7%) at DEQ station 3-HAL001.44 at Route 1 Business (Lafayette Boulevard).

A new TMDL is not required for this impaired segment of Hazel Run because the downstream Tidal freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_HAL01A00 / Hazel Run / Segment begins at the Route 95 crossing and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2004	L	4.73

Hazel Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.73

Cause Group Code: E20R-02-BEN Hazel Run

Cause Location: Begins at the Route 95 crossing and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Fredericksburg; Spotsylvania County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2016 Assessment: A total of two biological monitoring events in 2009 at DEQ station 3-HAL002.72 (upstream of Route 1) resulted in a VSCI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_HAL01A00 / Hazel Run / Segment begins at the Route 95 crossing and continues downstream until the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	4.73

Hazel Run

Aquatic Life	Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			4.73

Sources: Source Unknown

Cause Group Code: E20R-03-BAC Massaponax Creek

Cause Location: Segment begins at the confluence with an unnamed tributary to Massaponax Creek, approximately 0.25 rivermile upstream from the Route 639 bridge, and continues downstream until the confluence with another unnamed tributary, approximately 0.25 rivermile upstream of Ruffins Pond.

Cause City/County: Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MAP002.61 at Route 609 (Jim Morris Rd).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MAP007.97 at Route 1 (Jefferson Davis Hwy).

2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-MAP009.42 at Route 639 (Leavells Rd).

A new TMDL is not required for this impaired segment of Massaponax Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_MAP02A02 / Massaponax Creek / Segment begins at the confluence with an unnamed tributary to Massaponax Creek, at rivermile 2.68, and continues downstream until the confluence with another unnamed tributary, approximately 0.25 rivermile upstream of Ruffins Pond.	4A	Escherichia coli (E. coli)	2006	L	1.21
VAN-E20R_MAP02B12 / Massaponax Creek / Segment begins at the confluence with an unnamed tributary to Massaponax Creek, just upstream of Route 1, and continues downstream until the confluence with another unnamed tributary, at rivermile 2.68.	4A	Escherichia coli (E. coli)	2004	L	5.19
VAN-E20R_MAP03A02 / Massaponax Creek / Segment begins at the confluence with an unnamed tributary to Massaponax Creek, approximately 0.25 rivermile upstream from the Route 639 bridge, and continues downstream until the confluence with another unnamed tributary, just upstream from Route 1.	4A	Escherichia coli (E. coli)	2010	L	1.67

Massaponax Creek

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)		
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			8.07	

Cause Group Code: E20R-03-BEN Little Falls Run

Cause Location: Begins at the headwaters of Little Falls Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Stafford County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2020 Assessment: A total of two biological monitoring events in 2013 at DEQ station 3-LIA003.14 (0.02 miles downstream from Route 606) resulted in a VCPMI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_LIA01A04 / Little Falls Run / Segment begins at the headwaters of Little Falls Run and continues downstream until the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2016	L	4.93

Little Falls Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			4.93

Sources: Source Unknown

Cause Group Code: E20R-04-BAC Massaponax Creek

Cause Location: Begins at the confluence with an unnamed tributary, approximately 1.1 rivermiles downstream from Route 673, and continues downstream until the confluence with another unnamed tributary, approximately 0.25 rivermile upstream from Route 639.

Cause City/County: Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2014 Assessment: E. coli bacteria criterion excursions (7 of 7 samples - 100.0%) at DEQ station 3-MAP010.37 at Route 208 (Courthouse Road).

A new TMDL is not required for this impaired segment of Massaponax Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_MAP04A02 / Massaponax Creek / Segment begins at the confluence with an unnamed tributary, approximately 1.1 rivermiles downstream from Route 673, and continues downstream until the confluence with another unnamed tributary, approximately 0.25 rivermile upstream from Route 639.	4A	Escherichia coli (E. coli)	2008	L	2.17

Massaponax	Ureek				
Recreation		Estuary (Sq. Miles)	Reservoir (Acres)		
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			2.17	

Cause Group Code: E20R-04-PH Deep Run

Cause Location: Begins at the headwaters of Deep Run, and continues downstream to the confluence with an unnamed tributary at rivermile 2.19, downstream of Route 638.

Cause City/County: Spotsylvania County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: 2022 assessment: excursions less than the lower limit of the pH criterion range (5 of 28 samples - 17.9%) at National Park Service station 3DEP-06-NPS, upstream of Lee Drive.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDI Dev. Priori	Water Size
VAN-E20R_DEP03A12 / Deep Run / Segment begins at the headwaters of Deep Run and continues downstream to the confluence with an unnamed tributary at Route 638.	$5\mathrm{C}$	рН		2012	L	1.56
Deep Run Aquatic Life pH - Total Im	npaired Size	by Water Type:	Estuary (Sq. Miles)		ervoir res)	River (Miles) 1.56

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E20R-05-BAC Unnamed Tributary to Hazel Run

Cause Location: Segment begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with Hazel Run.

Cause City/County: Fredericksburg; Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (4 of 11 samples - 36.4%) at citizen monitoring station 3XHN-7-ALL.

A new TMDL is not required for this impaired segment of Hazel Run because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_XHN01A10 / Unnamed Tributary to Hazel Run / Segment begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with Hazel Run.	4A	Escherichia coli (E. coli)	2014	L	1.53

Unnamed Tributary to Hazel Run

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		1.53

Cause Group Code: E20R-05-PH Unnamed tributary to Massaponax Creek

Cause Location: Begins where XEN joins XFE and continues downstream until the confluence with Massaponax Creek at rivermile 8.06

Cause City/County: Spotsylvania County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: 2020 Assessment: Excursions less than the lower limit of the pH criterion range (2 of 11 samples - 18.2%) at DEQ station 3-XFE001.05 at Spotsylvania County Parkway.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_XFE01A02 / Unnamed tributary to Massaponax Creek / Segment begins where XEN joins XFE and continues downstream until the confluence with Massaponax Creek at rivermile 8.06	5C	рН	2016	L	1.27

Unnamed tributary to Massaponax Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:	, _		1.27

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E20R-06-BAC Unnamed tributary to Hazel Run

Cause Location: Segment begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with Hazel Run.

Cause City/County: Fredericksburg

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (2 of 11 samples - 18.2%) at citizen monitoring station 3XIA-9-ALL.

A new TMDL is not required for this impaired segment of Hazel Run because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_XIA01A12 / Unnamed tributary to Hazel Run / Segment begins at the headwaters of the unnamed tributary, and continues downstream to the confluence with Hazel Run.	4A	Escherichia coli (E. coli)	2014	L	2.23

Unnamed tributary to Hazel Run

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	(Miles)
	Escherichia coli (E. coli) - Iotal Impaired Size by Water Type:			2.23

Cause Group Code: E20R-06-PH Unnamed tributary to Deep Run

Cause Location: Begins at the headwaters and continues downstream until the confluence with Deep Run.

Cause City/County: Spotsylvania County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: Excursions less than the lower limit of the pH criterion range (16 of 24 samples - 66.7%) at citizen station 3XIL-DEEPRUN_FA-ALL

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_XIL01A24 / Unnamed tributary to Deep Run / Segment begins at the headwaters and continues downstream until the confluence with Deep Run.	5C	рН	2024	L	2.25

Unnamed tributary to Deep Run

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
*	pH - Total Impaired Size by Water Type:			2.25

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E20R-07-BAC Little Falls Run

Cause Location: Begins at the headwaters of Little Falls Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (4 of 12 samples - 33.3%) at DEQ station 3-LIA002.27 at Route 682 (Colebrooke Road).

A new TMDL is not required for this impaired segment of Little Falls Run because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_LIA01A04 / Little Falls Run / Segment begins at the headwaters of Little Falls Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2016	L	4.93

Little Falls Run

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.93

Cause Group Code: E20R-08-BAC Deep Run

Cause Location: Begins at the confluence with an unnamed tributary at Route 638 and continues downstream to the confluence with another unnamed tributary downstream of Route 756 at rivermile 0.74.

Cause City/County: Spotsylvania County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Insufficient E. coli data were available from DEQ station 3-DEP000.92 for the 2024 assessment (there was one STV exceedance in one or multiple 90-day periods but there were insufficient data to analyze geomean). The following information is carried over from the 2022 assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-DEP000.92 at Route 17 (Business) and there were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-DEP001.92 at Route 17 station 3-DEP001.59 at Latimers Knoll Ct.

A new TMDL is not required for this impaired segment of Deep Run because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_DEP02A18 / Deep Run / Segment begins at the confluence with an unnamed tributary at Route 638 and continues downstream to the confluence with another unnamed tributary downstream of Route 756 at rivermile 0.74.	4A	Escherichia coli (E. coli)	2018	L	1.67

Deep Run			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Typ	be:		1.67

Cause Group Code: E20R-09-BAC Rappahannock River

Cause Location: Begins at the E19/E20 watershed boundary, and extends downstream to the end of the free flowing waters of the Rappahannock River, at the Route 1 Alternate Bridge.

Cause City/County: Fredericksburg; Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: There were two or more STV exceedances in at least one 90-day period with <10 samples and no geomean exceedances at DEQ station 3-RPP110.57 at Route 1 (Jefferson Davis Hwy).

A new TMDL is not required for this impaired segment of Rappahannock River because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entre watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20R_RPP01A10 / Rappahannock River / Segment begins at the E19/E20 watershed boundary, and extends downstream to the end of the free flowing waters of the Rappahannock River, at the Route 1/Cambridge Street/Falmouth Bridge.	4A	Escherichia coli (E. coli)	2018	L	2.66

Rappahannoc	k River			
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			2.66

Cause Group Code: E21R-01-BAC Muddy Creek

Cause Location: Begins at the confluence with an unnamed tributary to Muddy Creek, approximately 0.7 rivermile downstream from Route 218, and continues downstream until the confluence with the Rappahannock River.

Cause City/County: King George County; Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (3 of 12 samples - 25.0%) at DEQ station 3-MUY001.43 at Route 3.

A new TMDL is not required for this impaired segment of Muddy Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_MUY01A00 / Muddy Creek / Segment begins at the confluence with White Oak Run and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2008	L	1.37
VAN-E21R_MUY01B20 / Muddy Creek / Segment begins at the confluence with an unnamed tributary to Muddy Creek, approximately 0.7 rivermile downstream from Route 218, and continues downstream until the confluence with White Oak Run.	4A	Escherichia coli (E. coli)	2008	L	2.21

Muddy Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.58

Cause Group Code: E21R-01-PH Portobago Creek

Cause Location: Begins at the confluence of two intermittent tributaries around rivermile 6.66 and extends downstream to the end of the free-flowing waters.

Cause City/County: Caroline County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: Excursions less than the lower limit of the pH criterion range (2 of 12 samples - 16.7%) at DEQ station 3-PBC003.09 at Route 17.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDL Dev. Priorit	Water Size
VAN-E21R_PBC01A10 / Portobago Creek / Segment begins at the confluence of two intermittent tributaries around rivermile 6.66 and extends downstream to the end of the free-flowing waters.	$5\mathrm{C}$	рН		2020	L	7
Portobago Creek Aquatic Life pH - Total Im	paired Size	by Water Type:	Estuary (Sq. Miles)	Rese (Ac		River (Miles) 7

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E21R-02-BAC Ware Creek

Cause Location: Segment begins at the confluence with an unnamed tributary to Ware Creek, just downstream from Burma Road, and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Caroline County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-WAE000.72 at Route 17 (Tidewater Trail).

A new TMDL is not required for this impaired segment of Ware Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_WAE01A08 / Ware Creek / Segment begins at the confluence with an unnamed tributary to Ware Creek, just downstream from Burma Road, and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2010	L	4.5

Ware Creek				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.5

Cause Group Code: E21R-02-BEN Ware Creek

Cause Location: Begins at the headwaters of Ware Creek and continues downstream until the confluence with an unnamed tributary to Ware Creek, just downstream from Burma Road.

Cause City/County: Caroline County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2008 Assessment: One biological monitoring event in 2002 at DEQ station 3-WAE005.95 (at Fort A.P. Hill) resulted in a VCPMI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_WAE02A04 / Ware Creek / Segment begins at the headwaters of Ware Creek and continues downstream until the confluence with an unnamed tributary to Ware Creek, just downstream from Burma Road.	5A	Benthic Macroinvertebrates Bioassessments	2008	L	3.07

Ware Creek

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			3.07

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Source Unknown

Cause Group Code: E21R-02-PH Ware Creek

Cause Location: Begins at the headwaters of Ware Creek and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Caroline County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: 2008 Assessment: Excursions less than the lower limit of the pH criterion range (2 of 2 samples - 100%) at DEQ station 3-WAE005.95 at Fort A.P. Hill.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_WAE02A04 / Ware Creek / Segment begins at the headwaters of Ware Creek and continues downstream until the confluence with an unnamed tributary to Ware Creek, just downstream from Burma Road.	5C	рН		2004	L	3.07
Ware Creek Aquatic Life			Estuary (Sq. Miles)	Rese) (Ac		River Miles)

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Source Unknown

3.07

pH - Total Impaired Size by Water Type:

Cause Group Code: E21R-03-BAC Gingoteague Creek

Cause Location: Begins at the confluence with an unnamed tributary to Gingoteague Creek, at rivermile 2.99, and continues downstream until tidal waters, near the confluence with the Rappahannock River.

Cause City/County: King George County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/5A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GIN002.64 at Route 625 (Salem Church Rd).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_GIN01A08 / Gingoteague Creek / Segment begins at the confluence with an unnamed tributary to Gingoteague Creek, at rivermile 2.99, and continues downstream until tidal waters, near the confluence with the Rappahannock River.	5A	Escherichia coli (E. coli)	2008	L	1.49

Gingoteague Creek

		Estuary	$\operatorname{Reservoir}$	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.49

Sources: Source Unknown

Cause Group Code: E21R-03-BEN Gingoteague Creek

Cause Location: Begins at the confluence with an unnamed tributary to Gingoteague Creek, at rivermile 2.99, and continues downstream until tidal waters, near the confluence with the Rappahannock River.

Cause City/County: King George County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2016 Assessment: Two biological monitoring events in 2010 at DEQ station 3-GIN002.64 at Route 625 resulted in a VCPMI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_GIN01A08 / Gingoteague Creek / Segment begins at the confluence with an unnamed tributary to Gingoteague Creek, at rivermile 2.99, and continues downstream until tidal waters, near the confluence with the Rappahannock River.	5A	Benthic Macroinvertebrates Bioassessments	2012	L	1.49

Gingoteague Creek

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			1.49

Sources: Source Unknown

Cause Group Code: E21R-04-BAC Goldenvale Creek

Cause Location: Begins at the confluence with Doctor Branch and continues downstream until tidal waters, near the confluence with the Rappahannock River.

Cause City/County: Caroline County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-GLL001.98 at Route 17 (Tidewater Trail).

A new TMDL is not required for this impaired segment of Rappahannock River because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entre watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_GLL01A08 / Goldenvale Creek / Segment begins at the confluence with Doctor Branch and continues downstream until tidal waters, near the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2020	L	5.31

Goldenvale Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			5.31

Cause Group Code: E21R-04-BEN Mill Creek

Cause Location: Begins at the confluence with an unnamed tributary, at rivermile 9.5, and continues downstream until the confluence with Peumansend Creek, at rivermile 6.06.

Cause City/County: Caroline County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: 2010 Assessment: Two biological monitoring events in 2004 at DEQ station 3-MIC008.55 (on Fort A.P. Hill property) resulted in a VCPMI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_MIC02A06 / Mill Creek / Segment begins at the confluence with an unnamed tributary, at rivermile 9.5, and continues downstream until the confluence with Peumansend Creek, at rivermile 6.06.	5A	Benthic Macroinvertebrates Bioassessments	2008	L	3.59

Mill Creek

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			3.59

Sources: Source Unknown

Cause Group Code: E21R-04-PH Muddy Creek

Cause Location: Begins at the confluence with White Oak Run and continues downstream until the confluence with the Rappahannock River.

Cause City/County: King George County; Stafford County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: Excursions less than the lower limit of the pH criterion range (4 of 28 samples - 14.3%) at citizen monitoring station 3MUY-1-ALL.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAN-E21R_MUY01A00 / Muddy Creek / Segment begins at the confluence with White Oak Run and continues downstream until the confluence with the Rappahannock River.	$5\mathrm{C}$	рН		2024	L	1.37
Muddy Creek			Estuary	Rese	ervoir	River
Aquatic Life pH - Total In	npaired Size	by Water Type:	(Sq. Miles)		eres)	(Miles) 1.37

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E21R-05-BAC Mount Creek

Cause Location: Begins at the confluence with West Branch and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Caroline County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2012 Assessment: E. coli bacteria criterion excursions (3 of 18 samples - 16.7%) at DEQ station 3-MTC001.94 at Route 17.

A new TMDL is not required for this impaired segment of Mount Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_MTC01A08 / Mount Creek / Segment begins at the confluence with West Branch and continues downstream until the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2008	L	4.46

Mount Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.46

Cause Group Code: E21R-05-BEN White Oak Run

Cause Location: Begins just downstream from the Route 604 crossing and continues downstream until the confluence with Muddy Creek.

Cause City/County: Stafford County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: A total of four biological monitoring events in 2021 and 2022 at DEQ station 3-WHT003.73 at Route 601 (upstream crossing) resulted in a VCPMI assessment that indicates an impaired macroinvertebrate community.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_WHT01A06 / White Oak Run / Segment begins just downstream from the Route 604 crossing and continues downstream until the confluence with Muddy Creek.	5A	Benthic Macroinvertebrates Bioassessments	2014	L	6.52

White Oak Run

	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water			
Type:			6.52

Sources: Source Unknown

Cause Group Code: E21R-05-PH Mount Creek

Cause Location: Begins at the confluence with West Branch and continues downstream until the confluence with the Rappahannock River.

Cause City/County: Caroline County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: 2014 Assessment: Excursions less than the lower limit of the pH criterion range (9 of 11 samples - 81.8%) at DEQ station 3-MTC001.94 at Route 17.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDI Dev. Priorit	Water Size
VAN-E21R_MTC01A08 / Mount Creek / Segment begins at the confluence with West Branch and continues downstream until the confluence with the Rappahannock River.	$5\mathrm{C}$	рН		2008	L	4.46
Mount Creek Aquatic Life pH - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles)	Rese (Ac		River (Miles) 4.46

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E21R-06-BAC Lambs Creek

Cause Location: Begins at the confluence with Popcastle Creek and continues downstream until tidal waters, near the confluence with the Rappahannock River.

Cause City/County: King George County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-LAM000.57 at Route 3 (Kings Hwy).

A new TMDL is not required for this impaired segment of Lambs Creek because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_LAM01A08 / Lambs Creek / Segment begins at the confluence with Popcastle Creek and continues downstream until tidal waters, near the confluence with the Rappahannock River.	4A	Escherichia coli (E. coli)	2008	L	0.54

Lambs Creek

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(~1)	()	0.54

Cause Group Code: E21R-07-BAC Mill Creek

Cause Location: Begins at the confluence with Peumansend Creek, at rivermile 6.06, and continues downstream until the tidal waters of Mill Creek.

Cause City/County: Caroline County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2020 Assessment: E. coli bacteria criterion excursions (2 of 12 samples - 16.7%) at DEQ station 3-MIC0001.66 at Route 17.

The Rappahannock River and Tributaries bacteria TMDL for the Mill Creek watershed (Eq ID 2539) was approved by the EPA on 07/10/2019 (Fed ID 11483). The SWCB approved the TMDL on 06/27/2019.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_MIC01A08 / Mill Creek / Segment begins at the confluence with Peumansend Creek, at rivermile 6.06, and continues downstream until the tidal waters of Mill Creek.	4A	Escherichia coli (E. coli)	2008	L	4.59

Mill Creek

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E21R-10-BAC Jetts Creek

Cause Location: Segment begins at the confluence of Boom Swamp with Jetts Creek, and continues downstream to the end of the free flowing waters.

Cause City/County: King George County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-JET003.49 at Route 625 (Salem Church Rd).

The Rappahannock River and Tributaries bacteria TMDL for the Jetts Creek watershed (Eq ID 2561) was approved by the EPA on 07/10/2019 (Fed ID 11483). The SWCB approved the TMDL on 06/27/2019.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_JET01A10 / Jetts Creek / Segment begins at the confluence of Boom Swamp with Jetts Creek, and continues downstream to the end of the free flowing waters.	4A	Escherichia coli (E. coli)	2010	L	1.85

Jetts Creek

Recreation		Estuary (Sq. Miles)		
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		1.85	

Cause Group Code: E21R-11-BAC Portobago Creek

Cause Location: Segment begins at the confluence of two intermittent tributaries around rivermile 6.66 and extends downstream to the end of the free-flowing waters.

Cause City/County: Caroline County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2022 Assessment: There were two or more STV exceedances in at least one 90-day period with <10 samples at DEQ station 3-PBC003.09 at Route 17 (Tidewater Trail).

The Rappahannock River and Tributaries bacteria TMDL for the Portobago Creek watershed (Eq ID 2563) was approved by the EPA on 07/10/2019 (Fed ID 11483). The SWCB approved the TMDL on 06/27/2019.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_PBC01A10 / Portobago Creek / Segment begins at the confluence of two intermittent tributaries around rivermile 6.66 and extends downstream to the end of the free-flowing waters.	4A	Escherichia coli (E. coli)	2010	L	7

Portobago Creek				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /		7	

Cause Group Code: E21R-12-BAC White Oak Run

Cause Location: Begins just downstream from the Route 604 crossing and continues downstream until the confluence with Muddy Creek.

Cause City/County: Stafford County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: 2018 Assessment: E. coli bacteria criterion excursions (2 of 12 samples - 16.7%) at DEQ station 3-WHT000.35 at Route 601 (downstream crossing).

A new TMDL is not required for this impaired segment of White Oak Run because the downstream Tidal Freshwater Rappahannock River bacteria TMDL (Fed ID 34369, 05/05/2008) included modeling, source identification, and reductions that covered the entire watershed (Eq ID POL0569).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21R_WHT01A06 / White Oak Run / Segment begins just downstream from the Route 604 crossing and continues downstream until the confluence with Muddy Creek.	4A	Escherichia coli (E. coli)	2014	L	6.52

White Oak Run

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			6.52

Cause Group Code: E22E-02-EBEN Rappahannock River

Cause Location: The mesohaline mainstem of the Rappahannock River

Cause City/County: Essex County; Lancaster County; Middlesex County; Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: Estuarine Bioassessments/5A

Cause Description: In 2004, the mesohaline portion of the mainstem Rappahannock indicated benchic impairment based on the Chesapeake Bay Benchic Index of Biological Integrity. The impairment was attributed to low oxygen and the benchic impairment was treated as a confirmation of the impairment. The mainstem remained impaired in the 2006 cycle; however, due to guidance changes the segment was 303(d) listed for estuarine bioassessments.

The segment remains impaired.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP05A02 / Rappahannock River / The oligohaline/mesohaline boundary at river mile 48.51 to the downstream boundary of VDH shellfish condemnation area 025A-068B, 3/15/2021. RPPMH	5A	Estuarine Bioassessments	2006	L	6.958
VAP-E23E_RPP02A98 / Rappahannock River / Mainstem Rappahannock as described in VDH shellfish condemnation 025A-068A, 3/15/2021, excluding administratively condemned portion. RPPMH	5A	Estuarine Bioassessments	2006	L	8.123
VAP-E23E_RPP02B10 / Rappahannock River / Portion of mainstem Rappahannock River that is administratively condemned within condemnation 025A-068A, 3/15/2021. RPPMH	5A	Estuarine Bioassessments	2006	L	0.158
VAP-E23E_RPP02C12 / Rappahannock River / Portion of VDH shellfish condemnation 025A-068A, 11/14/2005 not included in 025A-068A, 3/15/2021. RPPMH	5A	Estuarine Bioassessments	2006	L	0.387
VAP-E24E_RPP01B14 / Garrett's Marina / As delineated in VDH shellfish condemnation 026-181A, 3/15/2021. RPPMH	$5\mathrm{A}$	Estuarine Bioassessments	2008	L	0.003
VAP-E24E_RPP01B98 / Rappahannock River: Garrett's Marina / As delineated in VDH shellfish condemnation 026-181M1, 3/15/2021. RPPMH	$5\mathrm{A}$	Estuarine Bioassessments	2008	L	0.025
VAP-E24E_RPP01C06 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/16/2007 (non-admin) that is open 3/15/2021. Shrank in the 2024 cycle. RPPMH	5A	Estuarine Bioassessments	2006	L	0.327

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(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP01D10 / Rappahannock River / The portion of the Rappahannock River within the administratively closed area of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	5A	Estuarine Bioassessments	2006	L	0.137
VAP-E24E_RPP01E18 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/15/2021 (non-admin) Expanded in the 2024 cycle. RPPMH	5A	Estuarine Bioassessments	2006	L	0.378
VAP-E24E_RPP03A00 / Rappahannock River / The Rappahannock River from the limit of VDH shellfish condemnation 068A, 11/14/2005 downstream to end of MSN (Sharps/0.7 mi DS of Mark Haven Beach) unless otherwise segmented RPPMH	5A	Estuarine Bioassessments	2006	L	10.858
VAP-E24E_RPP03D24 / Rappahannock River / Described in VDH shellfish condemnation 026-270A, 5/25/2022. RPPMH	5A	Estuarine Bioassessments	2006	L	0.061
VAP-E25E_ROS02D24 / Robinson Creek / Portion of VDH-DSS Condemnation 028-177B, 1/15/2022 within Robinson Creek. RPPMH	5A	Estuarine Bioassessments	2024	L	0.039
VAP-E25E_RPP01C10 / Rappahannock River: Mark Haven Beach Basin / The portion of VDH shellfish condemnation 026-181B, 1/20/2006 not administratively closed. RPPMH	5A	Estuarine Bioassessments	2008	L	0.010
VAP-E25E_RPP01C98 / Mark Haven Beach Basin / As delineated in VDH shellfish condemnation 026-181A, 4/3/2012. RPPMH	5A	Estuarine Bioassessments	2008	L	0.004
VAP-E25E_RPP02A02 / Rappahannock River / The mainstem of the Rappahannock River from the end of MSN (Sharps/0.7 mi DS of Mark Haven Beach to the mouth, excluding area in SFC 051A. RPPMH	5A	Estuarine Bioassessments	2006	L	81.443
VAP-E25E_RPP03A06 / Rappahannock River / Described in VDH Shellfish Condemnation 024-070B, 11/15/2022. RPPMH	5A	Estuarine Bioassessments	2006	L	0.008
VAP-E25E_RPP03B16 / Rappahannock River Run Bluffs / As described in VDH shellfish condemnation 026-181B, 3/15/2021. RPPMH	$5\mathrm{A}$	Estuarine Bioassessments	2006	L	0.003
VAP-E25E_ZZZ01D14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA69. RPPMH	5A	Estuarine Bioassessments	2024	L	0.239

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(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CRR02A08 / Corrotoman River / The portion of the Corrotoman River that is within CB segment RPPMH.	5A	Estuarine Bioassessments	2006	L	1.039
VAP-E26E_MLL02A06 / Mill Creek / Downstream of VDH shellfish condemnation 103, 12/10/1991 RPPMH	5A	Estuarine Bioassessments	2024	L	0.358
VAP-E26E_RPP04A00 / Rappahannock River / Described in VDH Shellfish Condemnation 029-051B, 9/1/2015. RPPMH	5A	Estuarine Bioassessments	2006	L	0.131
VAP-E26E_RPP05A00 / Rappahannock River / Delineated in VDH-DSS condemnation 029-051C, 9/1/2015. RPPMH	5A	Estuarine Bioassessments	2006	L	0.049
VAP-E26E_RPP07A02 / Rappahannock River / As delineated in VDH-DSS condemnation 018-053A, 7/23/2018. RPPMH	5A	Estuarine Bioassessments	2006	L	0.139
VAP-E26E_RPP08A22 / Rappahannock River / Portion of VDH-DSS Condemnation 032-109A, 9/15/2022 that is within the mainstem Rappahannock River. Shrank in the 2024 cycle. RPPMH	5A	Estuarine Bioassessments	2006	L	0.002
VAP-E26E_ZZZ01E14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA74 RPPMH	5A	Estuarine Bioassessments	2024	L	0.613
Rappahannock River		Estuary	Rese	ervoir Riv	ver

	Estuary	Reservoir	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Estuarine Bioassessments - Total Impaired Size by Water Type:	111.488			

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E22E-03-BAC Peedee Creek

Cause Location: Tidal Peedee Creek

Cause City/County: Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2014 cycle, tidal Peedee Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 6/13 at 3-PEE003.97.

Tidal Peedee Creek was addressed in the Rappahannock River and Tributaries Bacterial TMDL, which was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. The impairment is Category 4A.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Reservoir

(Acres)

River

(Miles)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_PEE01A14 / Peedee Creek / Tidal portion of Peedee Creek. RPPOH	4A	Enterococcus	2014	L	0.15

Peedee Creek		
		Estuary
Recreation		(Sq. Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.15

Cause Group Code: E22E-04-BAC Occupacia Creek

Cause Location: Tidal Occupacia Creek

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2014 cycle, tidal Occupacia Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 9/11 at 3-OCC005.62.

During the 2020 cycle, monitoring at 3-OCC001.85 had an exceedance rate of 3/9.

New bacteria criteria were implemented in the 2022 cycle. There was insufficient information to assess 3-OCC001.85. However, the creek remained impaired due to four geometric mean exceedances at 3-OCC005.62.

It is nested in the bacterial TMDL for Occupacia and Farmers Hall Creeks, which was developed and was approved by the EPA on 7/30/2007; therefore, the segment is considered Category 4A. The TMDL addresses the nontidal watersheds feeding into the tidal portion and the upstream bacterial reductions should improve water quality downstream.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_OCC01A08 / Occupacia Creek / The tidal portion of Occupacia Creek RPPOH	4A	Enterococcus	2014	L	0.668

Occupacia Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.668		

Cause Group Code: E22E-05-BAC Rappahannock River

Cause Location: The Rappahannock River from the tidal freshwater/oligohaline boundary downstream to river mile 56.21.

Cause City/County: Essex County; Richmond County; Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2014 cycle, the Rappahannock River from the tidal freshwater oligohaline boundary downstream to rivermile 51.04 was impaired of the Recreation Use due to an enterococci exceedance rate of 2/12 at 3-RPP056.20.

The impairment was nested within the Upper Rappahannock River Shellfish TMDL, which was approved by the EPA on 8/10/2010 and was considered Category 4A.

However, during the 2016 cycle, the upper portion of the impairment, which was not located within the actual TMDL study area boundary, was split off and considered Category 5A.

The impairment was subsequently addressed in the Rappahannock River and Tributaries Bacterial TMDL, which was approved by the SWCB on 6/27/2019 and by the EPA on 7/20/2019, and is considered Category 4A.

New bacteria criteria were implemented in the 2022 cycle. There is insufficient information to assess the criteria; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP02A02 / Rappahannock River / The Rappahannock River from the tidal freshwater/oligohaline boundary downstream to river mile 56.21. RPPOH	4A	Enterococcus	2014	L	1.344

Rappahannock River

Recre

		Estuary	Reservoir	River	
eation		(Sq. Miles)	(Acres)	(Miles)	
	Enterococcus - Total Impaired Size by Water Type:	1.344			

Cause Group Code: E22E-06-BAC Rappahannock River

Cause Location: The oligohaline/mesohaline boundary at river mile 48.51 downstream to the limit of VDH shellfish condemnation area 025A-068B, 415/2020.

Cause City/County: Essex County; Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2014 cycle, the segment was impaired of the Recreation Use due to an enterococci exceedance rate of 4/12 at 3-RPP046.26.

It is located within the study area for the Upper Rappahannock River Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010. The enterococci impairment is considered nested within the TMDL; therefore, the segment is considered Category 4A.

New bacteria criteria were implemented in the 2022 cycle. There is insufficient information to assess the criteria at 3-RPP046.26; therefore, the impairment is carried over. In addition, monitoring at 3-RPP044.06 was impaired due to four geometric mean exceedances.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP05A02 / Rappahannock River / The oligohaline/mesohaline boundary at river mile 48.51 to the downstream boundary of VDH shellfish condemnation area 025A-068B, 3/15/2021. RPPMH	4A	Enterococcus	2014	L	6.958

Rappahannock River

Recreation		Estuary (Sq. Miles)		
	Enterococcus - Total Impaired Size by Water Type:	6.958		

Cause Group Code: E22E-07-BAC Rappahannock River

Cause Location: The Rappahannock River from rivermile 56.21 downstream to river mile 51.04.

Cause City/County: Essex County; Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

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Cause Description: During the 2014 cycle, the Rappahannock River from the tidal freshwater oligohaline boundary downstream to rivermile 51.04 was impaired of the Recreation Use due to an enterococci exceedance rate of 2/12 at 3-RPP056.20.

The impairment was nested within the Upper Rappahannock River Shellfish TMDL, which was approved by the EPA on 8/10/2010 and was considered Category 4A.

However, during the 2016 cycle, the upper portion of the impairment, which was not located within the actual TMDL study area boundary, was split off and was considered Category 5A (the TMDL was later completed in 2019). This nested segment remains Category 4A.

New bacteria criteria were implemented in the 2022 cycle. There was insufficient information to assess the criteria; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP02B16 / Rappahannock River / The Rappahannock River from rivermile 56.21 downstream to river mile 51.04. RPPOH	4A	Enterococcus	2014	L	2.003

Estuary Reservoir River	
Recreation (Sq. Miles) (Acres) (Miles)	
Enterococcus - Total Impaired Size by Water Type: 2.003	

Cause Group Code: E22E-09-BAC Waterview Creek

Cause Location: The tidal portion of Waterview Creek.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2018 cycle, tidal Waterview Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 10/12 at 3-WAR001.81.

It is located within the study area for the Upper Rappahannock River Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/31/2010. The enterococci impairment is considered nested within the TMDL; therefore, the segment is considered Category 4A.

New bacteria criteria were implemented in the 2022 cycle. The segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_WAR01A18 / Waterview Creek / Tidal portion of Waterview Creek RPPMH	4A	Enterococcus	2018	L	0.038

Waterview Creek				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.038		

Cause Group Code: E22E-10-BAC Bridge Creek

Cause Location: The tidal portion of Bridge Creek to its mouth at Occupacia Creek.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2020 cycle, tidal Bridge Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 4/9 at 3-BDG000.10.

The creek is located within the study area for the Upper Rappahannock River Watershed Bacterial TMDL (growing areas 25 and 26.) The TMDL was approved by the SWCB on 8/10/2010 and by the EPA on 12/13/2010. The impairment will be addressed during implementation and is considered nested (Category 4A.)

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data confirmed that the segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_BDG01A20 / Bridge Creek / Tidal Bridge Creek to its mouth at Occupacia Creek RPPOH	4A	Enterococcus	2020	L	0.123

Bridge Creek

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Enterococcus - Total Impaired Size by Water Type:	0.123	× ,	

Cause Group Code: E22E-11-PH Occupacia Creek

Cause Location: Tidal Occupacia Creek

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2022 cycle, tidal Occupacia Creek was impaired of the Aquatic Life Use due to a pH exceedance rate of 8/20 at 3-OCC005.62, which is located off of Route 661. The nontidal Occupacia Creek watershed is designated as Class VII waters, so it is suspected that the low pH is a natural condition.

Downstream monitoring at 3-OCC001.85 is acceptable.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAP-E22E_OCC01A08 / Occupacia Creek / The tidal portion of Occupacia Creek RPPOH	$5\mathrm{C}$	pH		2022	L	0.668
Occupacia Creek Aquatic Life pH - Total I	mpaired Size	by Water Type:	Estuary (Sq. Miles) 0.668	Rese (Ac	rvoir res)	River (Miles)

Cause Group Code: E22E-12-PH Waterview Creek

Cause Location: Tidal Waterview Creek

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2022 cycle, tidal Waterview Creek was impaired of the Aquatic Life Use due to a pH exceedance rate of 3/25 at 3-WAR001.81, which is located at Waterview Road. Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDL Dev. Priorit	Water
VAP-E22E_WAR01A18 / Waterview Creek / Tidal portion of Waterview Creek RPPMH	$5\mathrm{C}$	рН		2022	L	0.038
Waterview Creek				D		
Aquatic Life			Estuary (Sq. Miles)			River (Miles)

pH - Total Impaired Size by Water Type: 0.038

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Sediment Resuspension (Clean Sediment); Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E22R-01-BAC Occupacia Creek

Cause Location: Occupacia Creek from the Hunters Millpond Dam to the extent of tidal influence.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Occupacia Creek was initially assessed as impaired of the Recreation Use during the 2002 cycle based on fecal coliform violations at the Route 17 bridge (3-OCC010.47). In 2006 the segment was also impaired for E. coli. During the 2008 cycle, the impairment converted to E. coli with a violation rate of 3/21.

The bacterial TMDL for Occupacia and Farmers Hall Creeks was developed and was approved by the EPA on 7/30/2007; therefore, the segment is considered Category 4A.

Occupacia Creek remained impaired in the 2020 cycle due to an E.coli exceedance rate of 3/12.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_OCC01A98 / Occupacia Creek / Occupacia Creek from Hunters Millpond downstream to the tidal limit.	4A	Escherichia coli (E. coli)	2006	L	2.34

Occupacia Creek

Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	Estuary (Sq. Miles)		

Cause Group Code: E22R-02-BAC Farmers Hall Creek

Cause Location: Farmers Hall Creek from its headwaters to its tidal limit

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: In the 2004 cycle, Farmers Hall Creek was assessed as not supporting of the Recreation Use support goal based on a fecal coliform violation rate of 3/13 at the Route 631 bridge (3-FAR002.88). The TMDL was due in 2016.

The bacterial TMDL for Occupacia and Farmers Hall Creeks was approved by the EPA on 7/30/2007; therefore the segment is considered Category 4A.

The WQS has changed from fecal coliform to E. coli. No monitoring had been conducted since 2006, so the fecal coliform impairment had been carried over through the 2020 cycle.

E. coli sampling was conducted in the 2022 cycle under a revised E. coli criteria. The segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples. The impairment is being converted to E. coli.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_FAR01A04 / Farmers Hall Creek / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2022	L	4.01

Farmers Hall Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
E	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	,		4.01

Cause Group Code: E22R-04-BAC Elmwood Creek and Tributary XHY

Cause Location: The nontidal portion of Elmwood Creek and its tributary XHY in its entirety.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Elmwood Creek and its tributary were assessed as not supporting of the Recreation Use in the 2014 cycle based on multiple E. coli exceedances.

The exceedance rates were as follows in the 2016 cycle: 5/23 at 3-ELM002.23 5/13 at 3-ELM002.92 1/13 (FS) at 3-ELM004.27 4/13 at 3-XHY000.06 1/12 (FS) at 3-XHY002.50

During the 2020 cycle, the exceedance rates were 5/24 at 3-ELM002.23, 1/12 (FS) at 3-ELM002.92, and 2/13 at 3-XHY000.06.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The Rappahannock River and Tributaries Bacteria TMDL was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. The impairment was addressed in the TMDL and will be considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_ELM01A06 / Elmwood Creek and tributary XHY / Headwaters to tidal limit, including tributary XHY.	4A	Escherichia coli (E. coli)	2014	L	9.07

Elmwood Creek and Tributary XHY Recreation (Sq. Miles) (Acres) (Miles) Escherichia coli (E. coli) - Total Impaired Size by Water Type: 9.07

Cause Group Code: E22R-04-PH Elmwood Creek and Tributary XHY

Cause Location: The nontidal portion of Elmwood Creek and its tributary XHY in its entirety.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: Elmwood Creek was assessed as not supporting of the Aquatic Life Use in the 2006 cycle based on a pH exceedance rate of 4/10 at 3-ELM002.23, which is located at the Route 17 bridge.

Additional data was collected during the 2014 and 2016 cycles. The impairment was expanded to incorporate tributary XHY. The exceedance rates were as follows: 5/24 at 3-ELM002.23 5/26 at 3-ELM002.92 4/26 at 3-ELM004.27 6/26 at 3-XHY000.06 2/25 (FS) at 3-XHY002.50

During the 2020 cycle, the exceedance rates were 2/24 (FS) at 3-ELM002.23, 2/12 at 3-ELM002.92, and 3/13 at 3-XHY000.06.

Additional monitoring was conducted at 3-ELM002.92 (4/22) during the 2022 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_ELM01A06 / Elmwood Creek and tributary XHY / Headwaters to tidal limit, including tributary XHY.	$5\mathrm{C}$	рН	2006	L	9.07

Elmwood Creek and Tributary XHY

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
	pH - Total Impaired Size by Water Type:	× - /	``	9.07

Cause Group Code: E22R-05-BAC Baylors Creek

Cause Location: Baylors Creek from its headwaters to the extent of backwater of Baylors Pond.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2008 cycle, Baylors Creek was assessed as impaired of the Recreation Use due to an E.coli exceedance rate of 2/16 at the Route 17 bridge (3-BAY002.62).

Additional data was collected in the 2014 cycle. The impairment was confirmed with the following exceedance rates: 3/12 at 3-BAY002.62 3/11 at 3-BAY004.39 1/12 (FS) at 3-BAY006.66

During the 2020 cycle, the E.coli exceedance rates were 7/13 at 3-BAY002.62 and 3/13 at 3-BAY006.66.

New bacteria criteria were implemented in the 2022 cycle. No new data were collected at 3-BAY002.62, but re-analysis of the previous data confirms that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples. No new data was collected at 3-BAY004.39 and there is insufficient information to assess the criteria at 3-BAY006.66; therefore, the impairment is carried over at those stations, as well.

The Rappahannock River and Tributaries Bacteria TMDL was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. Baylors Creek was addressed and will be considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_BAY01A08 / Baylors Creek / Headwaters to extent of backwater at Baylors Pond.	4A	Escherichia coli (E. coli)	2008	L	5.89

Baylors Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			5.89

Cause Group Code: E22R-05-PH Baylors Creek

Cause Location: Baylors Creek from its headwaters to the extent of backwater of Baylors Pond.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2008 cycle, Baylors Creek was assessed as impaired of the Aquatic Life Use due to a pH exceedance rate of 6/16 at the Route 17 bridge (3-BAY002.62).

Additional monitoring was conducted during the 2014 cycle. The impairment was confirmed with the following exceedance rates: 2/13 at 3-BAY002.62 2/12 at 3-BAY004.39 11/13 at 3-BAY006.66

In the 2022 cycle, the pH exceedance rates were 2/13 at 3-BAY002.62 and 16/25 at 3-BAY006.66.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_BAY01A08 / Baylors Creek / Headwaters to extent of backwater at Baylors Pond.	$5\mathrm{C}$	рН	2008	L	5.89

Baylors Creek

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)		
	pH - Total Impaired Size by Water Type:			5.89	

Cause Group Code: E22R-06-BAC Peedee Creek

Cause Location: The mainstem of Peedee Creek from its headwaters to the extent of tide.

Cause City/County: Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2020 cycle, Peedee Creek was assessed as not supporting of the Recreation Use due to an E. coli exceedance rate of 10/49 at the Route 640 bridge (3-PEE004.46).

Due to a previous impairment, Peedee Creek was addressed in the Rappahannock River and Tributaries Bacterial TMDL, which was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. The impairment is Category 4A.

New bacteria criteria were implemented in the 2022 cycle. The segment remained impaired due to two geometric mean exceedances at 3-PEE004.96 and two or more STV exceedances in the same 90-day period with <10 samples at 3-PEE004.46.

There was insufficient information to assess the station in the 2024 cycle (no STV exceedances but insufficient data to analyze geomean.) The impairment will be carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_PEE01A08 / Peedee Creek / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2020	L	3.3

Peedee Creek

		$\operatorname{Estuary}$	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.3

Cause Group Code: E22R-06-DO Peedee Creek

Cause Location: The mainstem of Peedee Creek from its headwaters to the extent of tide.

Cause City/County: Westmoreland County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2010 cycle, Peedee Creek was assessed as not supporting of the Aquatic Life Use due to dissolved oxygen violations at the Route 640 bridge (3-PEE004.46).

Additional monitoring was conducted along the creek in the 2014 and 2018 cycles. 7/12 at 3-PEE004.11 24/48 at 3-PEE004.46 7/12 at 3-PEE004.96 0/12 (FS) at 3-PEE006.57

In the 2022 cycle, the exceedance rate was 16/36 at 3-PEE004.46 and 18/31 at 3-PEE004.96. Monitoring in the 2024 cycle at 3-PEE004.46 shows that the impairment is continuing (14/34).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_PEE01A08 / Peedee Creek / Headwaters to tidal limit	$5\mathrm{C}$	Dissolved Oxygen	2010	L	3.3

Peedee Creek

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
	Dissolved Oxygen - Total Impaired Size by Water Type:	· · · · ·		3.3

Cause Group Code: E22R-06-PH Peedee Creek

Cause Location: The mainstem of Peedee Creek from its headwaters to the extent of tide.

Cause City/County: Westmoreland County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2008 cycle, Peedee Creek was assessed as not supporting of the Aquatic Life Use due to pH exceedances at the Route 640 bridge (3-PEE004.46).

Additional monitoring was conducted along the creek in the 2014 and 2018 cycles. 1/12 (FS) at 3-PEE004.11 2/48 (FS) at 3-PEE004.46 3/12 at 3-PEE004.96 3/12 at 3-PEE006.57

Station 3-PEE004.46 remained fully supporting during the 2020 cycle (3/49), however the segment remained impaired due to the previous upstream exceedances. Continued monitoring was recommended.

3-PEE004.96 was re-sampled in the 2022 cycle and was impaired (10/31). In the 2022 and 2024 cycles, 3-PEE004.46 remained fully supporting for pH.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_PEE01A08 / Peedee Creek / Headwaters to tidal limit	$5\mathrm{C}$	рН	2008	L	3.3

Peedee Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:			3.3

Cause Group Code: E22R-07-DO XGI - Occupacia Creek, UT

Cause Location: The unnamed tributary XGI in its entirety.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/4C

Cause Description: During the 2008 cycle, tributary XGI was mistakenly included within the nontidal Occupacia Creek segment. The segment failed for dissolved oxygen with an exceedance rate of 7/22 at station 3-XGI000.44.

However, this stream actually enters below the fall line on Occupacia Creek and therefore was not reclassified as Class VII waters. The TMDL is due in 2020. As the station was addressed in the Occupacia Natural Conditions Assessment (4/4/2005), it is considered Category 4C.

Additional monitoring in the 2016 cycle confirmed the impairment (2/5 for dissolved oxygen).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_XGI01A10 / XGI - Occupacia Creek, UT / Headwaters to mouth at tidal Occupacia Creek	4C	Dissolved Oxygen	NA	NA	1.96

XGI - Occupacia Creek, UT

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
	Dissolved Oxygen - Total Impaired Size by Water Type:	(24. 1.1100)	(110100)	1.96

Cause Group Code: E22R-07-PH XGI - Occupacia Creek, UT

Cause Location: The unnamed tributary XGI in its entirety.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/4C

Cause Description: During the 2008 cycle, tributary XGI was mistakenly included within the nontidal Occupacia Creek segment. The segment failed for pH with an exceedance rate of 22/22 at station 3-XGI000.44.

However, this stream actually enters below the fall line on Occupacia Creek and therefore was not reclassified as Class VII waters. The TMDL is due in 2020. As the station was addressed in the Occupacia Natural Conditions Assessment (4/4/2005), it is considered Category 4C.

Additional monitoring in the 2016 cycle confirmed the impairment (3/5 for pH).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_XGI01A10 / XGI - Occupacia Creek, UT / Headwaters to mouth at tidal Occupacia Creek	$4\mathrm{C}$	pН	NA	NA	1.96

XGI - Occupacia Creek, UT

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
	pH - Total Impaired Size by Water Type:	, - ,		1.96

Cause Group Code: E22R-08-BAC Stillwater Creek

Cause Location: Stillwater Creek from its headwaters at Cockerel Creek downstream to its tidal limit

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Stillwater Creek was assessed as not supporting of the Recreation Use in the 2014 cycle based on an E. coli exceedance rate of 3/12 at 3-STL003.35 (Route 17 South). Monitoring at 3-STL001.54, which is located at the Route 674 bridge, was acceptable (0/12).

During the 2020 cycle, the exceedance rates were 4/13 at 3-STL003.35 and 0/13 (S) at 3-STL001.54.

New bacteria criteria were implemented in the 2022 cycle. The segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples at 3-STL003.35. There is insufficient data to assess 3-STL001.54.

The Rappahannock River and Tributaries Bacteria TMDL was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. The Stillwater Creek impairment was addressed and is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_STL01A14 / Stillwater Creek / Headwaters at Cockerel Creek to tidal limit	4A	Escherichia coli (E. coli)	2014	L	3.53

Stillwater Creek

_		•	Reservoir	
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.53

Cause Group Code: E22R-08-DO Stillwater Creek

Cause Location: Stillwater Creek from its headwaters at Cockerel Creek downstream to its tidal limit

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: Stillwater Creek was assessed as not supporting of the Aquatic Life Use in the 2014 cycle based on a dissolved oxygen exceedance rate of 4/13 at 3-STL003.35 (Route 17 South). Monitoring at 3-STL001.54, which is located at the Route 674 bridge, was acceptable (1/13).

During the 2022 cycle, the exceedance rates were 6/25 at 3-STL003.35 and 2/25 (S) at 3-STL001.54.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_STL01A14 / Stillwater Creek / Headwaters at Cockerel Creek to tidal limit	$5\mathrm{C}$	Dissolved Oxygen	2014	L	3.53

Stillwater Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Dissolved Oxygen - Total Impaired Size by Water Type:			3.53

Cause Group Code: E22R-08-PH Stillwater Creek

Cause Location: Stillwater Creek from its headwaters at Cockerel Creek downstream to its tidal limit

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: Stillwater Creek was assessed as not supporting of the Aquatic Life Use in the 2014 cycle based on pH exceedance rates of 12/13 at 3-STL003.35 (Route 17 South) and 4/13 at 3-STL001.54 (Route 674).

During the 2022 cycle, the exceedance rates were 21/25 at 3-STL003.35 and 3/25 at 3-STL001.54.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_STL01A14 / Stillwater Creek / Headwaters at Cockerel Creek to tidal limit	$5\mathrm{C}$	рН	2014	L	3.53

Stillwater Creek

Aquatic Life		Estuary (Sq. Miles)	
	pH - Total Impaired Size by Water Type:		3.53

Cause Group Code: E22R-10-PH Mill Swamp

Cause Location: Nontidal Mill Swamp below Horners Pond

Cause City/County: Westmoreland County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2014 cycle, Mill Swamp was impaired of the Aquatic Life Use due to a pH exceedance rate of 2/12 at 3-MSW000.85, which is located at Route 625 below Horners Pond.

The exceedance rate was 4/24 during the 2020 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22R_MSW01A14 / Mill Swamp / Horners Pond dam to tidal limit	5C	рН	2014	L	0.73

Mill Swamp

Aquatic Life		Estuary (Sq. Miles)		
	pH - Total Impaired Size by Water Type:		0.73	

Cause Group Code: E22R-11-BAC Smoots Mill Run, UT

Cause Location: From its headwaters to its mouth at Smoots Mill Run.

Cause City/County: Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2020 cycle, the tributary was impaired of the Recreation Use due to an E. coli exceedance rate of 5/11 at 3-SMO001.58, which is located at Route 697.

The Rappahannock River and Tributaries Bacteria TMDL was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. The creek is located within the study area and will be addressed during implementation; therefore, the impairment is considered nested (Category 4A.)

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data confirms that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_SMO01A14 / Smoots Mill Run / Headwaters to mouth.	4A	Escherichia coli (E. coli)	2020	L	1.67

Smoots Mill Run, UT				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.67	

Sources: Agriculture; Non-Point Source; On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)

Cause Group Code: E22R-11-PH Smoots Mill Run, UT

Cause Location: From its headwaters to its mouth at Smoots Mill Run.

Cause City/County: Westmoreland County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2014 cycle, the tributary was impaired of the Aquatic Life Use due to a pH exceedance rate of 7/12 at 3-SMO001.58, which is located at Route 697.

The exceedance rate was 17/24 during the 2020 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_SMO01A14 / Smoots Mill Run / Headwaters to mouth.	$5\mathrm{C}$	рН	2014	L	1.67

Smoots Mill Run, UT

		Estuary	$\operatorname{Reservoir}$	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:			1.67

Cause Group Code: E22R-12-BAC Troy Creek

Cause Location: The nontidal portion of Troy Creek.

Cause City/County: Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2020 cycle, Troy Creek was impaired of the Recreation Use due to an E. coli exceedance rate of 4/12 at 3-TRY002.08, which is located at Route 637.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The Rappahannock River and Tributaries Bacteria TMDL was approved by the SWCB on 6/27/2019 and by the EPA on 7/10/2019. Troy Creek is located within the study area and will be addressed during implementation; therefore, the impairment is considered nested (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E22R_TRY01A06 / Troy Creek / The nontidal portion of Troy Creek	4A	Escherichia coli (E. coli)	2020	L	4.29

Troy Creek			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.29

Cause Group Code: E23E-01-SF Upper Rappahannock River, Little Carter Creek, Jugs Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 025-068A, 3/15/2021

Cause City/County: Essex County; Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Non-administrative portion of VDH Shellfish Condemnation 025-068A, 3/15/2021

The Upper Rappahannock River Watershed Shellfish TMDL was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010; therefore, the impaired area is considered Category 4A. The condemnation has since shortened; the area currently open for harvest is considered Cat. 2C.

Note: a previous Little Carter Creek/Jugs Creek VDH-DSS Shellfish Condemnation (068B, 3/6/2002) remains closed but is now incorporated into this shellfish condemnation.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_LIE01A98 / Little Carter Creek, Jugs Creek / Tidal limit to mouth at the Rappahannock River. RPPMH	4A	Fecal Coliform	1998	L	0.419
VAP-E23E_PIS02A00 / Piscataway Creek / The estuarine portion of Piscataway Creek. RPPMH	4A	Fecal Coliform	1998	L	0.589
VAP-E23E_RPP02A98 / Rappahannock River / Mainstem Rappahannock as described in VDH shellfish condemnation 025A-068A, 3/15/2021, excluding administratively condemned portion. RPPMH	4A	Fecal Coliform	1998	L	8.123
VAP-E23E_ZZZ02A06 / Unsegmented estuaries in E23 / Unsegmented portion within VDH-DSS Condemnation 025A-068A, 3/15/2021. RPPMH	4A	Fecal Coliform	2006	L	0.049

Upper Rappahannock River, Little Carter Creek, Jugs Creek				
	Estuary	Reservoir	River	
Shellfishing	(Sq. Miles)	(Acres)	(Miles)	
Fecal Coliform - Total Impaired Size by Water Typ	e: 9.18			

Cause Group Code: E23E-02-BAC Cat Point Creek

Cause Location: The tidal portion of Cat Point Creek.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Tidal Cat Point Creek was impaired of the Recreation Use in the 2010 cycle due to enterococci violations at 3-CAT006.58, which is located below Rt. 624.

During the 2014 cycle, enterococci exceedance rates were 9/23 at 3-CAT006.58, as well as 3/12 at 3-CAT000.46.

Additional monitoring occurred at 3-CAT006.58 in the 2020 cycle; the exceedance rate was 5/8.

New bacteria criteria were implemented in the 2022 cycle. The segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples at 3-CAT006.58.

Cat Point Creek is located within the study area for the downstream Upper Rappahannock Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010; therefore, it is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_CAT01A02 / Cat Point Creek / The tidal portion of Cat Point Creek. RPPMH	4A	Enterococcus	2010	L	1.28

Cat Point Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	1.28		

Cause Group Code: E23E-03-BAC Hoskins Creek

Cause Location: The tidal portion of Hoskins Creek from the Tappahannock STP to its mouth at the Rappahannock River.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Tidal Hoskins Creek was initially included on the 1994 303(d) list based on excessive fecal coliform standard exceedances recorded at the Rt. 360 bridge (3-HOK000.74). The upstream limit was extended to the Town of Tappahannock STP in the 1998 cycle in recognition that the STP may be a contributing source. During the 2006 cycle, the segment remained impaired and enterococci was added as an impairing cause. TMDL monitoring was initiated in the 2008 cycle; the impairment was confirmed, extended upstream to the tidal limit, and switched to enterococci based on exceedances at multiple stations.

The entire segment remained impaired in the 2010 cycle due to the following enterococci exceedance rates: 5/13 at 3-CRC000.15 10/13 at 3-HOK000.15 24/36 at 3-HOK000.74 7/13 at 3-HOK002.74 7/13 at 3-HOK003.61

The bacterial TMDL, which was approved by the EPA on 3/27/2008 and by the SWCB on 4/28/2009 only addressed the area from the Tappahannock STP to its mouth. The extension was split off and is addressed in fact sheet E23E-03-BAC2; it is considered to be nested. Both areas are Category 4A.

During the 2020 cycle, the exceedance rates were 7/9 at 3-HOK002.74 and 4/8 at 3-HOK003.61 (in E23E-03-BAC2).

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data at 3-HOK003.61 and 3-HOK002.74 confirms that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_HOK01A98 / Hoskins Creek / Hoskins Creek from the Tappahannock STP downstream to the mouth at the Rappahannock River. RPPMH	4A	Enterococcus	2006	L	0.084

Hoskins Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Typ	be: 0.084		× ,

Cause Group Code: E23E-03-BAC2 Hoskins Creek, Church Swamp

Cause Location: The tidal portion of Hoskins Creek and Church Swamp downstream to the Tappahannock STP.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Tidal Hoskins Creek was initially included on the 1994 303(d) list based on excessive fecal coliform standard exceedances recorded at the Rt. 360 bridge (3-HOK000.74). The upstream limit was extended to the Town of Tappahannock STP in the 1998 cycle in recognition that the STP may be a contributing source. During the 2006 cycle, the segment remained impaired and enterococci was added as an impairing cause. TMDL monitoring was initiated in the 2008 cycle; the impairment was confirmed, extended upstream to the tidal limit, and switched to enterococci based on exceedances at multiple stations.

The entire segment remained impaired in the 2010 cycle due to the following enterococci exceedance rates: 5/13 at 3-CRC000.15 10/13 at 3-HOK000.15 24/36 at 3-HOK000.74 7/13 at 3-HOK002.74 7/13 at 3-HOK003.61

During the 2020 cycle, the exceedance rates were 4/8 at 3-HOK003.61 and 7/9 at 3-HOK002.74 (in E23E-03-BAC).

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data at 3-HOK003.61 and 3-HOK002.74 confirmed that the segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

The bacterial TMDL, which was approved by the EPA on 3/27/2008 and by the SWCB on 4/28/2009 only addressed the area from the Tappahannock STP to its mouth. The extension was split off. It is considered to be nested. Both areas are Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_CRC01A08 / Church Swamp / Tidal limit to mouth at Hoskins Creek RPPMH	4A	Enterococcus	2008	L	0.002
VAP-E23E_HOK02A08 / Hoskins Creek / Hoskins Creek from its tidal limit to the confluence with Church Swamp. RPPMH	4A	Enterococcus	2008	L	0.052
VAP-E23E_HOK02A10 / Hoskins Creek / Hoskins Creek from the confluence with Church Swamp downstream to the Tappahannock STP. RPPMH	4A	Enterococcus	2006	L	0.016

Hoskins Creek, Church Swamp

	(Acres)	r River (Miles)
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Cause Group Code: E23E-03-PH Hoskins Creek

Cause Location: Hoskins Creek from its tidal limit to the confluence with Church Swamp.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/4C

Cause Description: During the 2006 cycle, pH was added as an impairment because of exceedances at 3-HOK003.61, which is located at the Route 659 bridge. The violation rate was 13/36 in the 2010 cycle.

The upstream segment extent was corrected in the 2008 cycle due to acceptable pH values at three downstream stations.

A Natural Conditions Assessment was completed for Hoskins Creek during the 2012 cycle. The report recommends that tidal Hoskins Creek from its tidal limit downstream to the confluence with Church Swamp be reclassified as Class VII swampwaters. The stream is considered Category 4C.

In the 2020 cycle, pH remained impaired (2/11 at 3-HOK003.61).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_HOK02A08 / Hoskins Creek / Hoskins Creek from its tidal limit to the confluence with Church Swamp. RPPMH	4C	рН	NA	NA	0.052

Hoskins Creek

Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	
	pH - Total Impaired Size by Water Type:	0.052		

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Sediment Resuspension (Clean Sediment); Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E23E-05-BAC Little Carter Creek and Jugs Creek

Cause Location: Tidal Little Carter Creek and Jugs Creek downstream it their mouths at the Rappahannock River.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, Little Carter Creek and Jugs Creek were impaired of the Recreation Use due to an enterococci exceedance rate of 5/11 at 3-LIE003.62.

The area is within the study area for the Upper Rappahannock Watershed Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010. Implementation of the TMDL is expected to lower bacterial levels; therefore, the impairment is considered nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_LIE01A98 / Little Carter Creek, Jugs Creek / Tidal limit to mouth at the Rappahannock River. RPPMH	4A	Enterococcus	2012	L	0.419

Little Carter Creek and Jugs Creek

		$\operatorname{Estuary}$	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.419		

Cause Group Code: E23E-06-BAC Piscataway Creek

Cause Location: Tidal Piscataway Creek

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2016 cycle, the tidal Piscataway Creek was assessed not supporting of the Recreation Use based on an enterococci exceedance rate of 2/10 at the Route 17 bridge (3-PIS004.79).

Additional monitoring was conducted at 3-PIS004.67 during the 2020 cycle (7/18).

New bacteria criteria were implemented in the 2022 cycle. No additional data was collected at 3-PIS004.79. The segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples at 3-PIS004.67.

The area is within the study area for the Upper Rappahannock Watershed Shellfish TMDL, which was approved by the EPA on 8/10/2010. Implementation of the TMDL is expected to lower bacterial levels; therefore, the impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_PIS02A00 / Piscataway Creek / The estuarine portion of Piscataway Creek. RPPMH	4A	Enterococcus	2016	L	0.589

Piscataway Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.589		

Cause Group Code: E23E-07-BAC Mount Landing Creek

Cause Location: Tidal Mount Landing Creek

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2020 cycle, tidal Mount Landing Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 4/9 at 3-MTL000.12.

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data confirmed that the segment remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

The area is within the study area for the Upper Rappahannock Watershed Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010. Implementation of the TMDL is expected to lower bacterial levels; therefore, the impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_MTL01A10 / Mount Landing Creek / Tidal limit to mouth at the Rappahannock River. RPPMH	4A	Enterococcus	2020	L	0.172

Mount Landing Creek

Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	
	Enterococcus - Total Impaired Size by Water Type:	· · · · · · · · · · · · · · · · · · ·	× ,	

Cause Group Code: E23L-01-HG Chandlers Millpond

Cause Location: Chandlers Millpond in its entirety

Cause City/County: Westmoreland County

Use(s): Fish Consumption

Causes(s)/VA Category: Mercury in Fish Tissue/5A

Cause Description: On 8/31/2007, the Virginia Department of Health issued a fish consumption advisory for Chandlers Millpond based upon DEQ fish tissue monitoring at station 3-CMR001.00 in 2006. The advisory recommends consuming no more than two meals/month of largemouth bass due to the presence of mercury.

The DEQ monitoring showed mercury exceedances in both largemouth bass and black crappie.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water
VAP-E23L_CMR01A08 / Chandlers Millpond / Chandlers Millpond in its entirety	5A	Mercury in Fish	Tissue	2008	L	47.99
Chandlers Millpond						
			Estuary		ervoir	River
Fish Consumption			(Sq. Miles)) (Ac	$\operatorname{res})$	(Miles)
Mercury in Fish Tissue - Total In	npaired Size	by Water Type:		47	.99	

Sources: Atmospheric Deposition - Toxics; Source Unknown

Cause Group Code: E23R-03-PH Mill Creek and Mussell Swamp

Cause Location: Mill Creek and Mussell Swamp

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/4C

Cause Description: Piscataway Creek from Sturgeon Swamp downstream to the tidal limit was initially assessed not supporting of the Aquatic Life use support goal in 1998 based on pH standard exceedances recorded at monitoring station 3-PIS009.24. During the 2004 cycle, UT XFL was also considered impaired for pH (2004 fact sheet VAP-E23R-08).

During the 2006 cycle, additional watershed monitoring was performed and all of Piscataway Creek was impaired for pH, as well as XFL, XFM, XFN, Mussell Swamp, Sturgeon Swamp, and Mill Creek; therefore, the segment was expanded with TMDL due dates of 2018.

The "Natural Conditions Assessment for low pH, Piscataway Creek, Essex, Virginia" was completed. It recommends that Piscataway Creek and its tributaries from its headwaters to its mouth at the Rappahannock River be reclassified as Class VII swampwaters. However, only the Piscataway Creek watershed upstream of Sturgeon Swamp was reclassified as Class VII swampwaters; the reclassified portion was delisted for pH based upon acceptable exceedance rates.

The remainder includes Mill Creek and Mussell Swamp. Until the remainder of the watershed is reclassified, they are considered Cat. 4C for pH.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_MLC01A04 / Mill Creek / Headwaters to tidal limit	$4\mathrm{C}$	pH	NA	NA	5.27
VAP-E23R_MUS01A04 / Mussell Swamp / Headwaters to tidal limit.	$4\mathrm{C}$	рН	NA	NA	5.13

Mill Creek and Mussell Swamp

Aquatic Life		Estuary (Sq. Miles)	
-	pH - Total Impaired Size by Water Type:	· - /	10.4

Cause Group Code: E23R-04-BAC Hoskins Creek

Cause Location: Headwaters to tidal limit

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Hoskins Creek was assessed as impaired of the Recreation Use during the 2014 cycle due to E. coli exceedances at 3-HOK011.45.

The exceedance rate was 9/36 during the 2020 cycle.

New bacteria criteria were implemented in the 2022 cycle. There is insufficient information to assess the criteria in either the 2022 or 2024 cycles; therefore, the impairment is carried over.

The impairment is nested within the tidal Hoskins Creek TMDL, which was approved by the EPA on 3/27/2008 and by the SWCB on 4/28/2009; therefore, it is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E23R_HOK01A04 / Hoskins Creek / Headwaters to the tidal limit	4A	Escherichia coli (E. coli)	2014	L	13.17

Hoskins Creek

		Estuary		
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			13.17

Cause Group Code: E23R-06-BAC Cat Point Creek and Tributaries

Cause Location: Nontidal Cat Point Creek and all tributaries draining to that segment.

Cause City/County: Richmond County; Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2010 cycle, Cat Point Creek from Ruin Branch (river mile 14.1) downstream to the tidal limit near Canal Swamp (river mile 10.54) was assessed as not supporting of the Recreation Use due to E. coli violations at 3-CAT011.62, which is located at the Route 637 bridge.

During the 2012 cycle, Nanny Sanford Swamp above Chandlers Millpond was assessed as not supporting of the Recreation Use due to an E. coli exceedance rate of 3/12 at 3-NSS000.77, which is located at the Route 622 bridge. It was addressed in 2012 fact sheet E23R-01-BAC.

Additional monitoring was conducted during the 2014 cycle. The E. coli exceedances were widespread (see below); therefore, the impairments were combined and expanded to include all tributaries to nontidal Cat Point Creek.

6/30 at 3-CAT011.62 (2018 cycle) 3/12 at 3-NSS000.77 2/12 at 3-BLA002.31 4/12 at 3-CAT015.44 4/12 at 3-BRL000.15 3/12 at 3-CMR000.50 2/12 at 3-PAN003.00 5/12 at 3-RUN000.13 3/12 at 2-SYN000.42 2/12 at 3-TBS001.08 3/12 at 3-TBS003.39

In the 2020 cycle, the E. coli exceedance rate was 8/30 at 3-CAT011.62 and 4/9 at 3-TBS003.39.

The watershed is located within the study area for the downstream Upper Rappahannock Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010; therefore, it is considered nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. Station 3-TBS003.39 remained impaired due to two or more STV exceedances in the same 90-day period with <10 samples. The remaining stations were either not sampled or there was insufficient data to assess the new criteria; therefore, the impairment was carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_BLA01A06 / Black Swamp / Black Swamp from its headwaters downstream to Chandlers Millpond	4A	Escherichia coli (E. coli)	2014	L	4.19
VAP-E23R_CAT01A98 / Cat Point Creek / Cat Point Creek from Ruin Branch downstream to tidal limit near Canal Swamp (river mile 10.54)	4A	Escherichia coli (E. coli)	2010	L	5.34
VAP-E23R_CAT02A02 / Cat Point Creek / Cat Point Creek from The Big Swamp to Ruin Branch.	4A	Escherichia coli (E. coli)	2014	L	1.20
VAP-E23R_CAT03A04 / Cat Point Creek tributaries / Cat Point Creek tributaries above the tidal limit, excluding Black Swamp, The Big Swamp, Ruin Branch, and Nanny Sanford Swamp above Chandlers Millpond	4A	Escherichia coli (E. coli)	2014	L	94.76
VAP-E23R_NSS01A12 / Nanny Sanford Swamp / Mainstem above Chandlers Millpond	$4\mathrm{A}$	Escherichia coli (E. coli)	2012	L	3.58
VAP-E23R_RUN01A14 / Ruin Branch / Headwaters to mouth at Cat Point Creek	4A	Escherichia coli (E. coli)	2014	L	2.54
VAP-E23R_TBS01A06 / The Big Swamp / Headwaters to mouth at Cat Point Creek	4A	Escherichia coli (E. coli)	2014	L	6.75

Cat Point Creek and Tributaries

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			118.36

Cause Group Code: E23R-07-BEN Ruin Branch

Cause Location: Ruin Branch in its entirety

Cause City/County: Richmond County; Westmoreland County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: During the 2014 cycle, Ruin Branch was assessed as not supporting the Aquatic Life Use due to impairment of the benthic community at 3-RUN000.13.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_RUN01A14 / Ruin Branch / Headwaters to mouth at Cat Point Creek	5A	Benthic Macroinvertebrates Bioassessments	2014	L	2.54

Ruin Branch				
Η	Estuary	Reservoir	River	
Aquatic Life (Sector)	Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			2.54	

Sources: Source Unknown

Cause Group Code: E23R-08-BAC Muddy Run

Cause Location: Nontidal Muddy Run

Cause City/County: Richmond County; Westmoreland County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2014 cycle, Muddy Run was assessed as not supporting of the Recreation Use due to an E. coli exceedance rate of 2/12 at 3-MUR001.19, which is located at the Route 690 bridge.

The impairment is nested within the downstream Upper Rappahannock Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010; therefore, it is considered Category 4A.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_MUR01A04 / Muddy Run / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2014	L	4.66

Muddy Run

Estuary	Reservoir	River
Sq. Miles)	(Acres)	(Miles)
		4.66
	~	Estuary Reservoir q. Miles) (Acres)

Cause Group Code: E23R-12-DO Mussell Swamp

Cause Location: Headwaters to mouth at Piscataway Creek

Cause City/County: Essex County

Use(s): Aquatic Life

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Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2006 cycle, Mussell Swamp was assessed as impaired of the Aquatic Life Use based on dissolved oxygen exceedances at 3-MUS001.23, located at the Route 615 bridge.

Natural conditions are suspected; therefore, the segment is assessed as Cat. 5C until the natural conditions assessment can be performed.

During the 2008 cycle, the exceedance rate was 3/26. No additional monitoring has been conducted.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E23R_MUS01A04 / Mussell Swamp / Headwaters to tidal limit.	$5\mathrm{C}$	Dissolved Oxygen	2006	L	5.13

Mussell Swamp					
		Estuary	Reservoir	River	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	Dissolved Oxygen - Total Impaired Size by Water Type:			5.13	

Cause Group Code: E23R-16-BEN Church Swamp

Cause Location: Church Swamp from its headwaters to its tidal limit at Hoskins Creek

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: Benthic Macroinvertebrates Bioassessments/5A

Cause Description: During the 2008 cycle, Church Swamp was assessed as not supporting the Aquatic Life Use due to impairment of the benthic community at freshwater probabilistic monitoring station 3-CRC001.38.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_CRC01A06 / Church Swamp / Headwaters to tidal limit	5A	Benthic Macroinvertebrates Bioassessments	2008	L	3.24

Church Swamp				
	Estuary	Reservoir	River	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)	
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water				
Type:			3.24	

Sources: Source Unknown

Cause Group Code: E23R-19-BAC Clarks Run

Cause Location: Nontidal Clarks Run

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2014 cycle, Clarks Run was assessed as not supporting of the Recreation Use due to an E. coli exceedance rate of 3/12 at 3-CLK000.27, which is located at the Route 621 bridge.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The impairment is considered nested within the downstream Upper Rappahannock Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010; therefore, it is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E23R_CLK01A14 / Clarks Run / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2014	L	3.83

Clarks Run

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	,		3.83

Cause Group Code: E23R-20-DO Scates Millstream

Cause Location: Nontidal Scates Millstream

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2014 cycle, Scates Millstream was impaired of the Aquatic Life Use due to a dissolved oxygen exceedance rate of 2/12 at station 3-SMS000.77, which is located at Route 635.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_SMS01A14 / Scates Millstream / Headwaters to tidal limit	$5\mathrm{C}$	Dissolved Oxygen	2014	L	2.89

Scates Millstream				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Dissolved Oxygen - Total Impaired Size by Water Type:			2.89

Cause Group Code: E23R-20-PH Scates Millstream

Cause Location: Nontidal Scates Millstream

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2014 cycle, Scates Millstream was impaired of the Aquatic Life Use due to a pH exceedance rate of 6/12 at station 3-SMS000.77, which is located at Route 635.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	$\tilde{\mathrm{First}}$	TMDL Dev. Priority	Water Size
VAP-E23R_SMS01A14 / Scates Millstream / Headwaters to tidal limit	$5\mathrm{C}$	pH	2014	L	2.89

Scates Millstream				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:			2.89

Cause Group Code: E23R-22-BAC Mount Landing Creek

Cause Location: Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit at approximately river mile 4.44.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2020 cycle, Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit was impaired of the Recreation Use due to an E. coli exceedance rate of 5/23 at 3-MTL004.82, which is located at the Route 716 bridge.

The stream is located within the study area for the Upper Rappahannock River Watershed (growing areas 25 and 26) Bacterial TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010. The impairment is considered nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_MTL01A98 / Mount Landing Creek / Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit at approximately river mile 4.44.	4A	Escherichia coli (E. coli)	2020	L	1.16

Mount Landing Creek			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.16

Cause Group Code: E23R-22-PCB Mount Landing Creek

Cause Location: Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit at approximately river mile 4.44.

Cause City/County: Essex County

Use(s): Fish Consumption

Causes(s)/VA Category: PCBs in Fish Tissue/5A

Cause Description: During the 2020 cycle, Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit was impaired of the Fish Consumption Use due to PCBs over the fish tissue level in gizzard shad and blue catfish at 3-MTL004.82, which is located at the Route 716 bridge.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23R_MTL01A98 / Mount Landing Creek / Mount Landing Creek from the first tributary upstream of the Route 716 bridge downstream to the tidal limit at approximately river mile 4.44.	$5\mathrm{A}$	PCBs in Fish Tissue	2020	L	1.16

Mount Landing Creek

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
PCBs in Fish Tissue - Total Impaired Size by Water Type:			1.16

Sources: Source Unknown

Cause Group Code: E24E-01-SF Richardson Creek / Rappahannock River / Totuskey Creek

Cause Location: Portion of VDH Notice and Description of Shellfish Condemnation 025-071A, 3/15/2021

Cause City/County: Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH Shellfish Condemnation 025-071A, 3/15/2021

Previous shellfish condemnations have included Totuskey and Richardson Creeks (separately or combined). The streams have been impaired since the 1998 cycle. However, in 2006 the segments were recombined and extended into the Rappahannock mainstem. The condemnation was further extended in the 2008 cycle.

During the 2010 cycle, the condemnation was shortened and it was determined that the entire portion of the condemnation located within Totuskey Creek and portions of the Rappahannock River and Richardson Creek were considered administrative (VDH-DSS SFC 025-071A, 4/2/2008.) Those areas were partially delisted. The upstream portion of Richardson Creek remained listed.

The Totuskey and Richardson Creeks Bacterial TMDL was approved by the EPA on 2/19/2010. The TMDL was based on the maximum extent of the condemnation, which occurred in condemnation 025-071A, 3/16/2007.

The condemnation was shortened and split in the 2012 cycle and the Rappahannock River and the mouth of Richardson Creek were re-opened for harvest; those portions were partially delisted. The closed portion is considered Category 4A. Open areas are Category 2C.

The condemnations expanded and merged again in the 2018 cycle.

The condemnation expanded further in the 2022 and 2024 cycles and incorporated the previously open portion of Richardson Creek and Totuskey Creek.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RIC01A04 / Richardson Creek / Richardson Creek within VDH-DSS Condemnation 025-071A, 3/15/2021 (non-administrative.) RPPMH	4A	Fecal Coliform	1998	L	0.321
VAP-E24E_RPP01E18 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/15/2021 (non-admin) Expanded in the 2024 cycle. RPPMH	4A	Fecal Coliform	2018	L	0.378
VAP-E24E_TOT02B10 / Totuskey Creek / Portion of Totuskey Creek within the non-administrative portion of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	4A	Fecal Coliform	2024	L	0.064

Richardson Creek / Rappahannock River / Totuskey Creek

ShellfishingEstuary
(Sq. Miles)Reservoir
(Acres)River
(Miles)Fecal Coliform - Total Impaired Size by Water Type:0.763

Cause Group Code: E24E-02-BAC Totuskey Creek

Cause Location: The tidal portions of Totuskey Creek.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Totuskey Creek was previously assessed as not supporting of the Recreation Use because of fecal coliform exceedances at the Route 3 bridge (3-TOT005.11). During the 2006 cycle, the segment remained impaired for fecal coliform and enterococci was added as an impairment. During the 2008 cycle, the impairment converted solely to enterococci. The bacteria TMDL was due in 2014.

Additional monitoring has been conducted. During the 2016 cycle, the enterococci exceedance rates were as follows: 17/37 at 3-TOT005.11 (2020 cycle) 6/12 at 3-TOT006.34 6/12 at 3-LIK000.15 2/11 at 2-MAY000.12

New bacteria criteria were implemented. The segment remains impaired in the 2024 cycle due to two or more STV exceedances in the same 90-day period with <10 samples at 3-TOT005.11. There is insufficient information to assess the criteria at 3-TOT006.34. The remaining stations were not re-sampled.

The bacterial TMDL was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. Totuskey Creek is considered a Category 4A water.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_LIK01A12 / Little Totuskey Creek / Tidal limit to mouth at Totuskey Creek RPPMH	4A	Enterococcus	2006	L	0.055
VAP-E24E_TOT01A00 / Totuskey Creek / The segment boundary is delineated in VDH condemnation 025-071B, 3/15/2021 excluding Little Totuskey Creek. RPPMH	4A	Enterococcus	2006	L	0.302
VAP-E24E_TOT02A00 / Totuskey Creek / Portion of VDH shellfish condemnation 025-071A, 4/15/2020 within Totuskey Creek. RPPMH	4A	Enterococcus	2006	L	0.647
VAP-E24E_TOT02B10 / Totuskey Creek / Portion of Totuskey Creek within the non-administrative portion of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	4A	Enterococcus	2006	L	0.064

Totuskey Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	1.068		

Cause Group Code: E24E-02-EBTOX Totuskey Creek

Cause Location: The tidal portions of Totuskey Creek.

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: Sediment Bioassay/5A

Cause Description: During the 2006 cycle, estuarine probabilistic monitoring was conducted through the Coastal 2000 program at 3-TOT007.84 and 3-TOT004.92. The data was assessed by DEQ-CO through the Weight of Evidence approach. The alteration at station 3-TOT007.84 was assessed as Category 5A.

Note: The impairment cause was changed from toxics to estuarine bioassessments in the 2020 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_LIK01A12 / Little Totuskey Creek / Tidal limit to mouth at Totuskey Creek RPPMH	5A	Sediment Bioassay	2006	L	0.055
VAP-E24E_TOT01A00 / Totuskey Creek / The segment boundary is delineated in VDH condemnation 025-071B, 3/15/2021 excluding Little Totuskey Creek. RPPMH	5A	Sediment Bioassay	2006	L	0.302
VAP-E24E_TOT02A00 / Totuskey Creek / Portion of VDH shellfish condemnation 025-071A, 4/15/2020 within Totuskey Creek. RPPMH	$5\mathrm{A}$	Sediment Bioassay	2006	L	0.647
VAP-E24E_TOT02B10 / Totuskey Creek / Portion of Totuskey Creek within the non-administrative portion of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	5A	Sediment Bioassay	2006	L	0.064

Totuskey Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Sediment Bioassay - Total Impaired Size by Water Type:	1.068		

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E24E-06-SF Garrett's Marina

Cause Location: Described in VDH-DSS Condemnation 026-181A, 3/15/2021

Cause City/County: Essex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 026-181A, 3/15/2021

Garrett's Marina was impaired of the Shellfish Consumption Use in the 1998 cycle (E24E-03-SF). During the 2008 cycle, the condemnation expanded and incorporated previous condemnation M271, which had been seasonally condemned (observed effects).

Garrett's Marina was included in the Upper Rappahannock Watershed Shellfish TMDL, which was approved by the EPA on 8/10/2010; therefore, this portion was considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP01B14 / Garrett's Marina / As delineated in VDH shellfish condemnation 026-181A, 3/15/2021. RPPMH	4A	Fecal Coliform	2022	L	0.003

Garrett's Marina

Sh - 116 - h to -		•	Reservoir	
Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	$\begin{array}{c} \text{(Sq. Miles)} \\ 0.003 \end{array}$	(Acres)	(Miles)

Cause Group Code: E24E-07-SF Rappahannock River at Brown's Point

Cause Location: Described in VDH shellfish condemnation 026-270A, 5/25/2022.

Cause City/County: Essex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 026-270A, 5/25/2022

The area closed for harvest in the 2024 cycle. It is proposed for nesting in the upstream Upper Rappahannock River Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP03D24 / Rappahannock River / Described in VDH shellfish condemnation 026-270A, 5/25/2022. RPPMH	4A	Fecal Coliform	2024	L	0.061

Rappahannock River at Brown's Point

		Estuary	Reservoir	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.061			

Cause Group Code: E24R-01-BAC Bookers Mill Stream

Cause Location: Bookers Mill Stream from its headwaters to its mouth at the confluence with Totuskey Creek.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: Bookers Mill Stream was assessed not supporting of the Recreation Use support goal in 2002 based on fecal coliform exceedances recorded at the Route 612 bridge (3-BMS002.00). Monitoring was discontinued in 2001; therefore, the previous assessment was carried over.

Additional monitoring was conducted during the 2012 cycle; the E. coli exceedance rates were as follows: 3/12 at 3-BMS000.37 2/12 at 3-BMS002.00 5/12 at 3-BMS004.42

New bacteria criteria were implemented in the 2022 cycle. There has been no additional monitoring at 3-BMS000.37 and 3-BMS004.42 and there is insufficient information to assess the criteria at 3-BMS002.00; therefore, the impairment is carried over.

The bacterial TMDL for the tidal Recreation Use and Shellfish Use impairments on Totuskey Creek was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. The impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E24R_BMS01A98 / Bookers Mill Stream / Bookers Mill Stream in its entirety.	4A	Escherichia coli (E. coli)	2012	L	6.54

Bookers Mill Stream

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			6.54

Cause Group Code: E24R-01-DO Bookers Mill Stream

Cause Location: Bookers Mill Stream from its headwaters to its mouth at the confluence with Totuskey Creek.

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2012 cycle, Bookers Mill Stream was impaired of the Aquatic Life Use due to the following dissolved oxygen exceedance rates:

2/12 at 3-BMS000.37 0/14 at 3-BMS002.00 (FS) 3/12 at 3-BMS004.42

Station 3-BMS002.00 remained fully supporting in the 2022 cycle (0/7). However, the impairment will be carried over because the other stations were not re-sampled.

Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_BMS01A98 / Bookers Mill Stream / Bookers Mill Stream in its entirety.	$5\mathrm{C}$	Dissolved Oxygen	2012	L	6.54

Bookers Mill Stream	1			
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Dissolved Oxygen - Total Impaired Size by Water Type:			6.54

Cause Group Code: E24R-02-BAC Totuskey Creek

Cause Location: The free flowing portion of Totuskey Creek.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2006 cycle, the nontidal portion of Totuskey Creek was assessed as not supporting the Recreation Use due to E. coli exceedances at 3-TOT009.95, which is located at the Route 619 bridge.

During the 2012 cycle, the exceedance rates were as follows: 5/25 at 3-TOT009.95 3/12 at 3-TOT012.53 4/12 at 3-TOT014.49

The exceedance rate was 3/11 at 3-TOT009.95 in the 2020 cycle.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected at 3-TOT012.53 and 3-TOT014.49 and there is insufficient information to assess the criteria at 3-TOT009.95; therefore, the impairment is carried over.

The bacterial TMDL for the tidal Recreation Use and Shellfish Use impairments was completed during the 2010 cycle. It was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. The nontidal Recreation Use impairment is considered to be nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	$\tilde{\mathrm{First}}$	TMDL Dev. Priority	Water Size
VAP-E24R_TOT01A06 / Totuskey Creek / The nontidal portion of Totuskey Creek	4A	Escherichia coli (E. coli)	2006	L	8.05

Totuskey Creek

	Estuary	$\operatorname{Reservoir}$	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			8.05

Cause Group Code: E24R-03-BAC Muddy Gut

Cause Location: Headwaters to mouth at Rappahannock River.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2008 cycle, Muddy Gut was assessed as impaired of the Recreation Use based on an E. coli violation rate of 5/10 at the Route 607 bridge (3-MUG000.96).

New bacteria criteria were implemented in the 2022 cycle. There is insufficient information to assess the criteria; therefore, the impairment is carried over.

Muddy Gut is located within the study area for the Upper Rappahannock Shellfish TMDL, which was approved by the EPA on 2/10/2010 and by the SWCB on 9/30/2010. Muddy Gut is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_MUG01A08 / Muddy Gut / Headwaters to mouth at the Rappahannock River.	4A	Escherichia coli (E. coli)	2008	L	2.64

Muddy Gut

		Estuary		
Recreation	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	(Sq. Miles)	(Acres)	$\frac{\text{(Miles)}}{2.64}$

Cause Group Code: E24R-03-PH Muddy Gut

Cause Location: Headwaters to mouth at Rappahannock River.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2008 cycle, Muddy Gut was assessed as impaired of the Aquatic Life Use based on a pH exceedance rate of 4/10 at the Route 607 bridge (3-MUG000.96).

The station was re-sampled in the 2022 cycle and remains impaired (4/10).

Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_MUG01A08 / Muddy Gut / Headwaters to mouth at the Rappahannock River.	$5\mathrm{C}$	pH	2008	L	2.64

Muddy Gut

Aquatic Life pH - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)		
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Cause Group Code: E24R-04-BAC Little Totuskey Creek

Cause Location: Headwaters to the tidal limit

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, Little Totuskey Creek was assessed as not supporting the Recreation Use due to an E. coli exceedance rate of 2/12 at LIK002.21, which is located at the Route 360 bridge.

New bacteria criteria were implemented in the 2022 cycle. That station has not been re-sampled. Additional monitoring was conducted at 3-LIK002.12. There is insufficient information to assess the criteria; therefore, the impairment is carried over.

The bacterial TMDL for the tidal Totuskey Creek Recreation Use and Shellfish Use impairments was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. The Recreation Use impairment is considered to be nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_LIK01A08 / Little Totuskey Creek / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2012	L	1.91

Little Totuskey Creek				
	Estuary	Reservoir	River	
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, _		1.91	

Cause Group Code: E24R-05-PH Branham Mill Swamp

Cause Location: Branham Mill Swamp from its headwaters to its mouth at Marshy Swamp

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2012 cycle, Branham Mill Swamp was impaired of the Aquatic Life Use due to a pH exceedance rate of 2/12 at 3-BRA000.85.

The site is on private property and cannot be re-sampled.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E24R_BRA01A08 / Branham Mill Swamp / Headwaters to mouth at Marshy Swamp	$5\mathrm{C}$	pH	2012	L	3.66

Branham Mill Swamp				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:			3.66

Cause Group Code: E24R-06-BAC Richardson Creek and Tributaries

Cause Location: Headwaters to the tidal limit

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, the streams were assessed as impaired of the Recreation Use due to E. coli exceedances.

4/23 at 2-RIC003.85 4/12 at 3-RIC005.00 5/12 at 3-RIC006.43 3/12 at 3-RNF002.04 1/12 at 3-XHJ000.04 (FS)

The exceedance rate was 3/11 in the 2020 cycle at 3-RIC003.85.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The bacterial TMDL for the tidal Totuskey and Richardson Creeks Recreation Use and Shellfish Use impairments was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. The Recreation Use impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_RIC01A12 / Richardson Creek and Tributaries / The nontidal streams in the Richardson Creek watershed.	4A	Escherichia coli (E. coli)	2012	L	17.22

Richardson Creek and Tributaries				
	· · · · · · · · · · · · · · · · · · ·	Reservoir		
Recreation	(Sq. Miles)	(Acres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			17.22	

Cause Group Code: E24R-06-DO Richardson Creek and Tributaries

Cause Location: Headwaters to the tidal limit

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2012 cycle, Richardson Creek and its tributaries were impaired of the Aquatic Life Use due to dissolved oxygen exceedances.

During the 2016 cycle, the exceedance rates were as follows: 11/24 at 3-RIC003.85 0/12 (FS) at 3-RIC005.00 4/12 at 3-RIC006.43 1/12 (FS) at 3-RNF002.04 7/12 at 3-XHJ000.04

Additional monitoring was conducted at 3-RIC003.85 during the 2022 cycle; the station remained impaired (5/11).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_RIC01A12 / Richardson Creek and Tributaries / The nontidal streams in the Richardson Creek watershed.	$5\mathrm{C}$	Dissolved Oxygen	2012	L	17.22

Richardson	Creek and	Tributaries
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Aquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E24R-06-PH Richardson Creek and Tributaries

Cause Location: Headwaters to the tidal limit

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2012 cycle, Richardson Creek and its tributaries were impaired of the Aquatic Life Use due to pH exceedances.

The pH exceedance rates were as follows in the 2016 cycle: 16/24 at 3-RIC003.85 3/12 at 3-RIC005.00 11/12 at 3-RIC006.43 2/12 at 3-RNF002.04 7/12 at 3-XHJ000.04

Additional monitoring was conducted at 3-RIC003.85 during the 2022 cycle; the station remained impaired (2/11).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_RIC01A12 / Richardson Creek and Tributaries / The nontidal streams in the Richardson Creek watershed.	$5\mathrm{C}$	рН	2012	L	17.22

Richardson Creek and Tributaries

Aquatic Life	pH - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E24R-07-BAC Totuskey Creek Tributaries

Cause Location: The tributaries of Totuskey Creek above the confluence with Little Totuskey Creek, excluding Bookers Mill Swamp

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, the tributaries were impaired of the Recreation Use due to widespread E. coli exceedances.

3/11 at 3-MIL000.15 4/12 at 3-DRK001.35 4/12 at 3-XHK000.65 4/11 at 3-XHL000.96 6/11 at 3-XHM000.27

The bacterial TMDL for the tidal Totuskey and Richardson Creeks Recreation Use and Shellfish Use impairments was approved by the EPA on 2/19/2010 and by the SWCB on 9/30/2010. The Recreation Use impairment is considered to be nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_TOT02B12 / Totuskey Creek Tributaries / The nontidal tributaries of Totuskey Creek above the confluence with Little Totuskey, unless otherwise segmented.	4A	Escherichia coli (E. coli)	2012	L	73.27
VAP-E24R_XHL01A12 / XHL - Bookers Mill Stream, UT / Headwaters to mouth at Bookers Mill Stream	4A	Escherichia coli (E. coli)	2012	L	2.01

Totuskey Creek Tributaries

		Estuary	$\operatorname{Reservoir}$	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, - ,		75.28

Cause Group Code: E24R-08-PH XHL - Bookers Mill Stream, UT

Cause Location: Headwaters to mouth at Bookers Mill Stream

Cause City/County: Richmond County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2012 cycle, tributary XHL was impaired of the Aquatic Life Use due to a pH exceedance rate of 2/11 at 3-XHL000.96, which is located at the Route 603 bridge.

Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_XHL01A12 / XHL - Bookers Mill Stream, UT / Headwaters to mouth at Bookers Mill Stream	$5\mathrm{C}$	рН	2012	L	2.01
Stream					

XHL - Bookers Mill Stream, UT

		Estuary	Reservoir	River	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	pH - Total Impaired Size by Water Type:			2.01	

Cause Group Code: E24R-10-BAC Bellview Creek

Cause Location: Bellview Creek from its headwaters to its mouth at the Rappahannock River.

Cause City/County: Essex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2020 cycle, Bellview Creek was impaired of the Recreation Use due to an E.coli exceedance rate of 4/10 at 3-BLV002.94, which is located at Route 611.

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data confirmed that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

The stream is located within the study area for the Upper Rappahannock Shellfish TMDL, which was approved by the EPA on $\frac{8}{10}/2010$ and by the EPA on $\frac{12}{13}/2010$. The impairment is considered nested (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_BLV01A20 / Bellview Creek / Headwaters to mouth at the Rappahannock River	4A	Escherichia coli (E. coli)	2020	L	3.31

Bellview Creek			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, – , ,		3.31

Cause Group Code: E24R-10-DO Bellview Creek

Cause Location: Bellview Creek from its headwaters to its mouth at the Rappahannock River.

Cause City/County: Essex County

Use(s): Aquatic Life

Bellview Creek

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2020 cycle, Bellview Creek was impaired of the Aquatic Life Use to a dissolved oxygen exceedance rate of 2/10 at 3-BLV002.94, which is located at Route 611.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_BLV01A20 / Bellview Creek / Headwaters to mouth at the Rappahannock River	$5\mathrm{C}$	Dissolved Oxygen	2020	L	3.31

Aquatic Life		Estuary (Sq. Miles)	Reservoir		
fiquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	(bq. miles)	(Hereb)	3.31	

Cause Group Code: E24R-10-PH Bellview Creek

Cause Location: Bellview Creek from its headwaters to its mouth at the Rappahannock River.

Cause City/County: Essex County

Use(s): Aquatic Life

Causes(s)/VA Category: pH/5C

Cause Description: During the 2020 cycle, Bellview Creek was impaired of the Aquatic Life Use to a pH exceedance rate of 10/10 at 3-BLV002.94, which is located at Route 611.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24R_BLV01A20 / Bellview Creek / Headwaters to mouth at the Rappahannock River	$5\mathrm{C}$	pH	2020	L	3.31

Bellview Creek				
		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	pH - Total Impaired Size by Water Type:			3.31

Cause Group Code: E25E-01-BAC Lagrange Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 127, 6/11/1996

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, Lagrange Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 2/12 at 3-LGG001.92, which is located at the end of Route 656.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; the impairment is carried over.

The Lagrange Creek Shellfish Bacterial TMDL was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. Implementation of that TMDL is expected to bring the stream into compliance with the Recreation WQS; therefore, the impairment is considered nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LGG01A98 / Lagrange Creek / As described in VDH Shellfish Condemnation 028-127A, 1/15/2022. Expanded in the 2024 cycle. RPPMH	4A	Enterococcus	2012	L	0.562
VAP-E25E_LGG01B18 / Lagrange Creek / Portion of VDH Shellfish Condemnation 127, 6/11/1996 open on 028-127, 1/15/2022. RPPMH	4A	Enterococcus	2012	L	0.037

Lagrange Creek

Recreation		Estuary (Sq. Miles)		
	Enterococcus - Total Impaired Size by Water Type:	0.598		

Cause Group Code: E25E-01-SF Lagrange Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 127, 6/11/1996.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 028-127A,1/15/2022

A portion of Lagrange Creek was assessed in 1998 as not supporting the Shellfish Consumption Use based on VDH-DSS Condemnation 127, 6/11/1996. The TMDL for this portion was approved by the EPA on 11/15/2005 and by the SWCB on 11/15/2005. The segment is classified as Cat. 4A.

The condemnation has expanded and contracted several times and is currently smaller than the 1998 impairment. The condemned area is Category 4A. The re-opened area which was addressed in the TMDL was partially delisted (Category 2C.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LGG01A98 / Lagrange Creek / As described in VDH Shellfish Condemnation 028-127A, 1/15/2022. Expanded in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.562

Lagrange Creek

			Reservoir	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.561		

Cause Group Code: E25E-02-BAC Robinson Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 177, 5/28/1997

Cause City/County: Middlesex County

Use(s): Recreation; Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: In 2002, the segment was assessed as not supporting the Recreation Use due to fecal coliform exceedances at the end of Route 680 (3-ROS001.35). The violation rate in the 2004 cycle was 4/20.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The area was addressed in the "Rappahannock River: Lagrange and Robinson Creeks TMDL Report for Shellfish Condemnation Areas Listed due to Bacteria Contamination" which was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. Because the bacteria standard for the Shellfish Use is more stringent than the standard for the Recreation Use, the impairment is considered nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS01A00 / Robinson Creek / Described in VDH shellfish condemnation 028-177A and -177E, 1/15/2022. Expanded and merged in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.168
VAP-E25E_ROS01C20 / Robinson Creek / Portion of VDH shellfish condemnation 177, 5/28/1997 open in 028-177, 1/15/2022. RPPMH	4A	Fecal Coliform	2002	L	0.039

Robinson Creek

Recreation	Fecal Coliform - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 0.207	Reservoir (Acres)	River (Miles)	
Robinson Creek		Estuary	Reservoir	River	
Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	(Sq. Miles) 0.207	(Acres)	(Miles)	

Cause Group Code: E25E-02-SF Robinson Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 177, 5/28/1997

Cause City/County: Middlesex County

Use(s): Recreation; Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 028-177A and -177E, 1/15/2022

The upstream portion of Robinson Creek was assessed in 1998 as not supporting the Shellfish Consumption Use based on VDH-DSS Condemnation 177, 5/28/1997. The TMDL was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. The impairment is classified as Category 4A.

In the 2020 cycle, the condemnation shrank and split. The open and seasonally condemned areas were partially delisted (Category 2C and 2C/2B, respectively.)

The condemnations adjusted in the 2022 and 2024 cycles but remain smaller than the TMDL extent.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS01A00 / Robinson Creek / Described in VDH shellfish condemnation 028-177A and -177E, 1/15/2022. Expanded and merged in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.168

Robinson Creek

Recreation	Fecal Coliform - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 0.168	Reservoir (Acres)	River (Miles)
Robinson Creek				
Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 0.168	Reservoir (Acres)	$\operatorname{River}(\operatorname{Miles})$
	recar comorni rotar impaired size sy trater rype.	0.100		

Cause Group Code: E25E-05-BAC Farnham Creek

Cause Location: Farnham Creek from its tidal limit to its mouth at the Rappahannock River.

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: In 2002, Farnham Creek was assessed as not supporting of the Recreation Use due to fecal coliform exceedances at 3-FAM002.62, which is located at the Route 608 bridge.

The bacteria TMDL for shellfish condemnations in Farnham Creek was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2008. The Recreation Use impairment is considered to be nested.

The impairment converted to enterococci in the 2010 cycle.

During the 2016 cycle, the exceedance rate was 8/12.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_FAM01A98 / Farnham Creek / Portion of VDH shellfish condemnation 070, 10/22/1996 not administratively condemned in 024-070A, 11/15/2022. RPPMH	4A	Enterococcus	2010	L	0.350
VAP-E25E_FAM01B22 / Farnham Creek / Portion of VDH shellfish condemnation 070, 10/22/1996 administratively condemned in 024-070A, 11/15/2022. RPPMH	4A	Enterococcus	2010	L	0.074

Farnham Creek					
		Estuary	Reservoir	River	
Recreation		(Sq. Miles)	(Acres)	(Miles)	
	Enterococcus - Total Impaired Size by Water Type:	0.425			

Cause Group Code: E25E-05-SF Farnham Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 024-070A, 10/22/1996.

Cause City/County: Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS Condemnation 024-070A, 11/15/2022

Farnham Creek has been assessed as not supporting the Shellfish Use since 1998. The TMDL was due in 2010.

The bacteria TMDL for shellfish condemnations in Farnham Creek was approved by the EPA on 8/2/2006. The TMDL was based on the extent of the 1998 condemnation, which extended to the mouth of Farnham Creek (070, 10/22/1996).

During the 2010 cycle, the condemnation size was reduced; the lower portion now open for harvest was partially delisted (Category 2C). The condemned area is considered a Category 4A water for the Shellfish Consumption Use.

The condemnation expanded in the 2020 cycle and matches the extent of the completed TMDL.

However, in the 2022 cycle, a portion of the condemnation converted to administratively condemned and was partially delisted.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_FAM01A98 / Farnham Creek / Portion of VDH shellfish condemnation 070, 10/22/1996 not administratively condemned in 024-070A, 11/15/2022. RPPMH	4A	Fecal Coliform	1998	L	0.35

Farnham Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.35	. ,	

Cause Group Code: E25E-07-SF Parrotts Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 090, 4/27/1989

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 027-090A, 1/15/2022

The Shellfish TMDL report for "Rappahannock River: Mud and Parrotts Creeks" was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. The TMDL addressed the 1998 portion of the current condemnation; therefore, the impairment is considered Cat. 4A. The downstream portion of the Parrotts Creek condemnation is addressed in fact sheet E25E-27-SF.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDI Dev. Priorit	Water Size
VAP-E25E_PRR01A02 / Parrotts Creek / Delineated in VDH shellfish condemnation 090, 4/27/1989. RPPMH	4A	Fecal Coliform		1998	L	0.153
Parrotts Creek			Estuary	Rese	ervoir	River
Shellfishing			(Sq. Miles)	(Ac	res)	(Miles)

Sources: Agriculture; Non-Point Source; On-site Treatment Systems (Septic Systems and Similar Decentralized Systems); Urban Runoff/Storm Sewers

0.153

Fecal Coliform - Total Impaired Size by Water Type:

Cause Group Code: E25E-10-SF Deep Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 121, 11/16/1994

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH-DSS Shellfish Condemnation 023-121B, 12/15/2021

A 0.0491 sq. mi. portion of Deep Creek was assessed as impaired of the Shellfish Consumption Use on the 1998 303(d) list due to VDH condemnation 121, 11/16/1994.

The condemnation began expanding in the 2002 cycle. However, the shellfish TMDL, which was approved by the EPA on 8/2/2006, only addressed the 1998 impairment. The original area is considered a Category 4A water; the TMDL for the downstream portion is addressed in fact sheet E25E-10-SF2.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_DEE01A04 / Deep Creek / Described in VDH shellfish condemnation 121, 11/16/1994. RPPMH	4A	Fecal Coliform	1998	L	0.049

Deep Cre	ek
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Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 0.049	Reservoir (Acres)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.049		

Cause Group Code: E25E-10-SF2 Deep Creek

Cause Location: Portions of VDH Shellfish Condemnation 023-121B, 12/15/2021 not included in the 11/16/1994 condemnation

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH-DSS condemnation 023-121B, 12/15/2021

A 0.0491 sq. mi. upstream portion of Deep Creek was assessed as impaired of the Shellfish Consumption Use on the 1998 303(d) list due to VDH condemnation 121, 11/16/1994. The condemnation began expanding in the 2002 cycle; however, the TMDL was completed only for the original impairment (see fact sheet E25E-10-SF). The TMDL for this downstream portion was due in 2014.

The expanded portion is nested within the upstream Deep Creek Shellfish TMDL, which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2006.

The condemnations shrank in the 2022 cycle. They expanded and merged in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_DEE01B08 / Deep Creek / Portions of VDH-DSS condemnations 023-121B, 12/15/2021 not included in the 11/16/1994 condemnation. Merged and adjusted in the 2024 cycle. RPPMH	4A	Fecal Coliform	2002	L	0.182
Deep Creek					

Shellfishing		Estuary (Sq. Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.182	

Cause Group Code: E25E-11-SF Lancaster Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 023-120A, 8/14/1995

Cause City/County: Lancaster County; Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 020-120A, 9/22/2022

A portion of Lancaster Creek was assessed as impaired of the Shellfish Use in the 1998 cycle due to VDH Shellfish Condemnation 120A, 8/14/1995.

The TMDL Report for Shellfish Areas Listed due to Bacterial Contamination for Lancaster Creek was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007. Although the condemnation on Lancaster Creek has extended downstream since the 1998 cycle, only the original impairment was included in the TMDL. The expansion is addressed in fact sheet E25E-11-SF2. This segment is considered Category 4A for the Shellfish Use.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LAN01A98 / Lancaster Creek / As delineated in VDH-DSS Shellfish Condemnation 023-120A, 8/14/1995. RPPMH	4A	Fecal Coliform	1998	L	0.27

Lancaster Creek

Shellfishing		Estuary (Sq. Miles)	Reservoir (Acres)	
U U	Fecal Coliform - Total Impaired Size by Water Type:	0.27	× ,	

Cause Group Code: E25E-11-SF2 Lancaster Creek

Cause Location: Portion of VDH Notice and Description of Shellfish Condemnation 023-120A, 9/22/2022 not included in condemnation 023-120A, 8/14/1995

Cause City/County: Lancaster County; Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 020-120A, 9/22/2022

A portion of Lancaster Creek was assessed as impaired of the Shellfish Use in the 1998 cycle due to VDH Shellfish Condemnation 120A, 8/14/1995. Although the condemnation on Lancaster Creek has extended downstream since the 1998 cycle, only the original impairment was included when the TMDL was developed. Since the segment was first expanded downstream in the 2002 cycle, the TMDL for this downstream segment was due in 2014.

It is considered nested in the upstream "TMDL Report for Shellfish Areas Listed due to Bacterial Contamination for Lancaster Creek," which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

The condemnation expanded slightly in the 2018 cycle (portion of VDH shellfish condemnation 020-120A, 12/19/2016).

The condemned area shrank in the 2020 cycle and the downstream portion became seasonally condemned and was partially delisted. It shrank further in the 2022 cycle.

It expanded again in the 2024 cycle and re-incorporated the seasonally condemned area.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LAN01B08 / Lancaster Creek / The portion of VDH Shellfish Condemnation 023-120A, 9/22/2022 not included in 120A, 8/14/1995. Merged and expanded in the 2024 cycle. RPPMH	4A	Fecal Coliform	2002	L	0.489

Lancaster Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.489		

Cause Group Code: E25E-12-SF Morattico Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 023-120B, 9/22/2022

Cause City/County: Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 023-120B, 9/22/2022

The Morattico Creek shellfish impairment is nested in the neighboring "TMDL Report for Shellfish Areas Listed due to Bacterial Contamination for Lancaster Creek," which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

The condemnation expanded in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_MTT01A00 / Morattico Creek / Delineated in VDH Shellfish Condemnation 023-120B, 9/22/2022. Expanded in the 2024 cycle. RPPMH	4A	Fecal Coliform	2002	L	0.345

Morattico Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.345		

Cause Group Code: E25E-13-SF Mulberry Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 120B, 8/14/1995.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 023-121A, 12/15/2021

A portion of Mulberry Creek was included on the 1998 303(d) list due to VDH Shellfish condemnation 120B, 8/14/1995. The TMDL for Shellfish Areas Listed due to Bacterial Contamination for Mulberry Creek was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

The TMDL only addressed the fecal coliform impairment within the 1998 portion of Mulberry Creek. The segment has shrunk and extended several times. The condemnation became smaller than the original impairment. The closed area remained Category 4A and the opened area was partially delisted (Category 2C/2B).

During the 2024 cycle, the condemnation expanded past the TMDL extent. The expansion will be considered nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_MUB01A02 / Mulberry Creek / Described in VDH shellfish condemnation 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.148
VAP-E25E_MUB02A06 / Mulberry Creek / Portion of VDH shellfish condemnation 023-121A, 12/15/2021 not included in 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Fecal Coliform	2024	L	0.100

Mulberry Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.248		

Cause Group Code: E25E-15-BAC Greenvale Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 094, 11/7/1994

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, Greenvale Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 4/5 at 3-GEE001.44, which is located at Route 624.

As the area is within the Greenvale Creek Shellfish TMDL which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007, the impairment is considered nested.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_GEE01A98 / Greenvale Creek / As delineated in VDH shellfish condemnation 094, 11/7/1994. RPPMH	4A	Enterococcus	2012	L	0.087

Greenvale Creek

Recreation		Estuary (Sq. Miles)	
	Enterococcus - Total Impaired Size by Water Type:	0.087	

Cause Group Code: E25E-15-SF Greenvale Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 094, 11/7/1994

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH-DSS condemnation 022-094A, 11/15/2022

A portion of Greenvale Creek was included on the 1998 303(d) list due to VDH condemnation 94, 11/7/1994.

The bacteria TMDL for the Shellfish Impairment on Greenvale Creek was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

The impairment has periodically expanded; however, the TMDL only addressed this 1998 portion, which is considered Category 4A.

The condemnation shrank in the 2020 cycle and became co-incident with the TMDL segment. It expanded again in the 2022 cycle (see E25E-32-SF)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_GEE01A98 / Greenvale Creek / As delineated in VDH shellfish condemnation 094, 11/7/1994. RPPMH	4A	Fecal Coliform	1998	L	0.087

Greenvale Creek				
		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
Feca	Coliform - Total Impaired Size by Water Type:	0.087		

Cause Group Code: E25E-22-SF Robinson Creek / Perkins Creek

Cause Location: As described in VDH Shellfish Condemnation 028-177B and -177C, 1/15/2022

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 028-177B and -177C, 1/15/2022

The upstream portion of Robinson Creek was assessed in 1998 as not supporting the Shellfish Consumption Use based on VDH-DSS Condemnation 177, 5/28/1997. The TMDL for that original portion was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

During the 2006 cycle, however, the condemnation extended downstream. The expansion is considered nested in the upstream Robinson Creek Shellfish TMDL.

The Perkins Creek condemnation expanded further in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS02A04 / Robinson Creek, UT and Perkins Creek / Described in VDH Shellfish Condemnation 028-177C, 1/15/2022 and the portion of 028-177B, 1/15/2022 that is within Perkins Creek. RPPMH	4A	Fecal Coliform	2006	L	0.039
VAP-E25E_ROS02D24 / Robinson Creek / Portion of VDH-DSS Condemnation 028-177B, 1/15/2022 within Robinson Creek. RPPMH	4A	Fecal Coliform	2024	L	0.039

Robinson Creek / Perkins Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
_	Fecal Coliform - Total Impaired Size by Water Type:	0.078	. ,	

Cause Group Code: E25E-27-SF Parrotts Creek

Cause Location: Portion of VDH Notice and Description of Shellfish Condemnation 027-090A, 1/15/2022 not included in 90, 4/27/1989

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 027-090A, 1/15/2022

A portion of Parrotts Creek was listed in the 1998 cycle due to VDH condemnation 027-090A, 8/18/2009. The Shellfish TMDL report for "Rappahannock River: Mud and Parrotts Creeks" was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

The condemnation subsequently expanded. The expanded area is considered nested in the upstream Parrotts Creek TMDL.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_PRR02A08 / Parrotts Creek / Portion of VDH-DSS Condemnation 027-090A, 1/15/2022 downstream of condemnation 090, 4/27/1989. RPPMH	4A	Fecal Coliform	2008	L	0.011

Parrotts Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
-	Fecal Coliform - Total Impaired Size by Water Type:		× ,	

Cause Group Code: E25E-30-BAC Town Bridge Swamp

Cause Location: Town Bridge Swamp from its tidal limit to its mouth at tidal Urbanna Creek

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, sampling on Town Bridge Swamp at 3-TWN000.35 upstream of Urbanna Creek indicated that a portion of the creek is tidally influenced. Town Bridge Swamp is impaired of the Recreation Use due to an enterococci exceedance rate of 5/11.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The impairment is considered nested due to the downstream Urbanna Creek Shellfish TMDL, which was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_TWN01A12 / Town Bridge Swamp / Tidal limit to mouth at Urbanna Creek RPPMH	4A	Enterococcus	2012	L	0.002

Town Bridge Swamp				
		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.002		

Cause Group Code: E25E-31-PCB Urbanna Creek

Cause Location: Urbanna Creek from its tidal limit to its mouth at the Rappahannock River.

Cause City/County: Middlesex County

Use(s): Fish Consumption

Causes(s)/VA Category: PCBs in Fish Tissue/5A

Cause Description: During the 2020 cycle, tidal Urbanna Creek was impaired of the Fish Consumption Use due to exceedance of the fish tissue level in croaker and gizzard shad during monitoring in 2018 at station 2-URB000.00.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_URB01A00 / Urbanna Creek / As described in VDH-DSS condemnation 029-042B, 2/14/2006. RPPMH	5A	PCBs in Fish Tissue	2020	L	0.215
VAP-E25E_URB02A00 / Urbanna Creek / As delineated in VDH shellfish condemnation 029-042A, 2/14/2006. RPPMH	$5\mathrm{A}$	PCBs in Fish Tissue	2020	L	0.238

Urbanna Creek

	Estuary	Reservoir	River
Fish Consumption	(Sq. Miles)	(Acres)	(Miles)
PCBs in Fish Tissue - Total Impaired Size by Water Type:	0.452		

Sources: Source Unknown

Cause Group Code: E25E-32-SF Greenvale Creek

Cause Location: Portion of VDH-DSS condemnation 022-094A,11/15/2022 not included in 94, 11/7/1994

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH-DSS condemnation 022-094A,11/15/2022

A portion of Greenvale Creek was included on the 1998 303(d) list due to VDH condemnation 94, 11/7/1994.

The bacteria TMDL for the Shellfish Impairment on Greenvale Creek was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

The condemnation has expanded and contracted several times; however, the TMDL only addressed the 1998 portion. This expansion is considered nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_GEE02A06 / Greenvale Creek / Portion of VDH-DSS condemnation 022-094A, 11/15/2022 not included in 94, 11/7/1994. RPPMH	4A	Fecal Coliform	2022	L	0.012

Greenvale Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.012		

Cause Group Code: E25E-33-SF Rappahannock River

Cause Location: As described in VDH shellfish condemnation 026-181B, 3/15/2021.

Cause City/County: Essex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS Condemnation 026-181B, 3/15/2021

The area closed for shellfish harvest in the 2024 cycle. It is proposed for nesting in the Mark Haven Beach Shellfish TMDL, which was approved by the EPA on 8/10/2010 and by the SWCB on 12/13/2010.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAP-E25E_RPP03B16 / Rappahannock River Run Bluffs / As described in VDH shellfish condemnation 026-181B, 3/15/2021. RPPMH	4A	Fecal Coliform		2024	L	0.003
Rappahannock River Shellfishing Fecal Coliform - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 0.003		ervoir res)	River (Miles)

Cause Group Code: E25E-34-SF Lancaster Creek

Cause Location: Described in VDH Shellfish Condemnation 023-120C, 9/22/2022.

Cause City/County: Richmond County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 023-120C, 9/22/2022

In the 2024 cycle, the area was closed for harvest and is not considered an administrative condemnation. It is proposed for nesting in the upstream Lancaster Creek Shellfish TMDL, which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LAN03A06 / Lancaster Creek / Described in VDH Shellfish Condemnation 023-120C, 9/22/2022. RPPMH	4A	Fecal Coliform		2024	L	0.023
Lancaster Creek			Fetuary	Boso	rvoir B	iver

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
_	Fecal Coliform - Total Impaired Size by Water Type:	0.023		

Cause Group Code: E25E-35-EBTOX Parrotts Creek

Cause Location: Parrotts Creek within VDH shellfish condemnation 090, 4/27/1989

Cause City/County: Middlesex County

Use(s): Aquatic Life

Causes(s)/VA Category: Sediment Bioassay/5A

Cause Description: During the 2024 cycle, the area around estuarine probabilistic monitoring station 3-PRR000.60 was impaired of the Aquatic Life Use based on weight-of-evidence analysis. The 2021 sample was assigned Category 5A, scenario 1.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Prior	Water
VAP-E25E_PRR01A02 / Parrotts Creek / Delineated in VDH shellfish condemnation 090, 4/27/1989. RPPMH	$5\mathrm{A}$	Sediment Bioassa	ıy	2024	L	0.153
Parrotts Creek Aquatic Life Sediment Bioassay - Total I	mpaired Size	by Water Type:	Estuary (Sq. Miles) 0.153		ervoir eres)	River (Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E25E-36-SF Weeks Creek

Cause Location: Delineated in VDH shellfish condemnation 202, 10/8/1996

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS condemnation 027-202A, 1/15/2021

Weeks Creek was assessed as not supporting of the Shellfish Use during the 1998 cycle due to VDH shellfish condemnation 202, 10/8/1996. The TMDL was completed for the original 1998 segment and was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

The condemnation was later delisted. It was relisted in the 2024 cycle (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water
VAP-E25E_WEE01A00 / Weeks Creek / Delineated in VDH shellfish condemnation 202, 10/8/1996. RPPMH	4A	Fecal Coliform		2024	L	0.123
Weeks Creek Shellfishing Fecal Coliform - Total In	npaired Size	by Water Type:	Estuary (Sq. Miles) 0.123		rvoir res)	River (Miles)

Cause Group Code: E25E-37-SF Robinson Creek, UT

Cause Location: Described in VDH Shellfish Condemnation 028-177D, 1/15/2022.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 028-177D, 1/15/2022

The area closed for harvest in the 2024 cycle. The impairment is proposed for nesting in the upstream Robinson Creek Shellfish TMDL, which was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS02C16 / Robinson Creek, UT / Described in VDH Shellfish Condemnation 028-177D, 1/15/2022. RPPMH	4A	Fecal Coliform	2024	L	0.013
Robinson Creek, UT			D	in Dia	

		Estuary	Reservoir	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.013			

Cause Group Code: E25R-01-BAC Laton Swamp

Cause Location: Laton Swamp from its headwaters to its mouth at Farnham Creek

Cause City/County: Richmond County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2014 cycle, Laton Swamp was assessed as impaired of the Recreation Use due to an E. coli exceedance rate of 3/12 at 3-LAT002.34, which is located at Route 3.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The impairment is nested in the downstream Farnham Creek Shellfish TMDL, which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25R_LAT01A14 / Laton Swamp / Headwaters to mouth at Farnham Creek.	4A	Escherichia coli (E. coli)	2014	L	4.87

	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.87

Cause Group Code: E25R-02-DO Lagrange Creek

Cause Location: Lagrange Creek from the headwaters to the extent of tide at approximately river mile 3.75.

Cause City/County: Middlesex County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: Lagrange Creek was assessed in 2010 as not supporting of the Aquatic Life Use support goal based on dissolved oxygen exceedances recorded at the Route 610 bridge (3-LGG004.54).

The exceedance rate was 7/24 during the 2012 cycle.

Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25R_LGG01A98 / Lagrange Creek / Lagrange Creek from its headwaters to the limit of tidal influence.	$5\mathrm{C}$	Dissolved Oxygen	2010	L	2.5

Lagrange Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Dissolved Oxygen - Total Impaired Size by Water Type:	, – , ,		2.5

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E25R-03-BAC Nickleberry Swamp

Cause Location: Nickleberry Swamp from its headwaters to its mouth at Hilliard Pond

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, Nickleberry Swamp was impaired of the Recreation Use due to an E. coli exceedance rate of 3/12 at 3-NIC000.38, which is located at Route 17.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The stream is located within the Lagrange Creek watershed, which has a completed shellfish TMDL. The TMDL was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. The impairment is considered nested (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E25R_NIC01A12 / Nickleberry Swamp / Headwaters to mouth at Hilliard Pond	4A	Escherichia coli (E. coli)	2012	L	1.86

Nickleberry Swamp			
	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:			1.86

Cause Group Code: E25R-04-BAC South Branch Lagrange Creek

Cause Location: The nontidal portion of South Branch Lagrange Creek.

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, the nontidal portion of South Branch Lagrange Creek was impaired of the Recreation Use due to an E. coli exceedance rate of 3/12 at 3-LSB002.17, which is located at Route 602.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The stream is located within the Lagrange Creek watershed, which has a completed shellfish TMDL. The TMDL was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. The impairment is considered nested (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	First	TMDL Dev. Priority	Water Size
VAP-E25R_LSB01A12 / South Branch Lagrange Creek / Hilliard Pond dam downstream to tidal limit	4A	Escherichia coli (E. coli)	2012	L	0.4

South Branch Lagrange Creek

Recreation		Estuary (Sq. Miles)	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:		0.4

Sources: Agriculture; Non-Point Source; On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)

Cause Group Code: E25R-04-DO South Branch Lagrange Creek

Cause Location: The nontidal portion of South Branch Lagrange Creek.

Cause City/County: Middlesex County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5A

Cause Description: South Branch Lagrange Creek was impaired of the Aquatic Life Use during the 2012 cycle due to a dissolved oxygen exceedance rate of 2/12 at 3-LSB002.17.

The low dissolved oxygen ($\sim 2 \text{ mg/L}$) occurred during the summer months. Natural conditions are possible, however it is unknown if Hilliard Pond contributes to the impairment; therefore, it will be considered Category 5A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25R_LSB01A12 / South Branch Lagrange Creek / Hilliard Pond dam downstream to tidal limit	5A	Dissolved Oxygen	2012	L	0.4

South Branch Lagrange Creek

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Dissolved Oxygen - Total Impaired Size by Water Type:			0.4

Sources: Dam or Impoundment; Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Source Unknown

Cause Group Code: E25R-17-DO Masons Mill Swamp

Cause Location: Masons Mill Swamp from its headwaters downstream to its tidal limit.

Cause City/County: Middlesex County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During older cycles, Masons Mill Swamp was mistakenly assessed as a tidal water. The creek was assessed as not supporting of the Aquatic Life Use for dissolved oxygen since the 2006 cycle because it was thought to be a part of the mesohaline portion of the Rappahannock; the TMDL had a 2010 due date because of the Bay Overlist.

However, during the 2008 cycle, it was determined that station 3-MAO000.62 is on the free flowing section of Masons Mill Swamp. The stream remained impaired for dissolved oxygen due to an exceedance rate of 4/13. The dissolved oxygen TMDL due date was changed to 2018.

Additional monitoring during the 2012 cycle confirmed the dissolved oxygen impairment (6/14).

A Natural Conditions Assessment was developed during the 2014 cycle. The report states "Based on the above information, a change in the water quality standards classification to Class VII Swampwater due to natural conditions, rather than a TMDL, is NOT indicated for Masons Mill Swamp because of TP elevated almost twice the USGS background level, for a total of 3.36 rivermiles. Masons Mill Swamp will be assessed as Category 4C, Impaired due to natural condition, no TMDL needed, although natural TP was greater than that recommended in USGS 1999." However, based on the elevated nutrients, the stream will remain Category 5C.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25R_MAO01A00 / Masons Mill Swamp / Masons Mill Swamp from its headwaters to its tidal limit near Route 604.	$5\mathrm{C}$	Dissolved Oxygen	2008	L	3.37

Masons Mill Swamp

		Estuary	Reservoir	River	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	Dissolved Oxygen - Total Impaired Size by Water Type:			3.37	

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E26E-01-SF Meachim Creek

Cause Location: Described in VDH Notice and Description of Shellfish Condemnation 179A, 12/9/1996.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 030-179A, and -179C, 9/15/2022

Two portions of Meachim Creek were included on the 1998 303(d) list due to 179A and 179B, 12/9/1996. The Shellfish TMDL was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006.

The condemnations have expanded and shrunk several times. In the 2018 cycle, the condemnations became smaller than the TMDL study areas and re-opened areas were delisted.

Condemnation A expanded slightly in the 2022 cycle, but remains smaller than the TMDL extent.

Condemnation C closed in the 2024 cycle. A portion of the closure is located within the TMDL area and will be considered Category 4A. The area outside of the TMDL will be nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MEA01A00 / Meachim Creek / Described in VDH shellfish condemnation 030-179A, 9/15/2022. RPPMH	4A	Fecal Coliform	1998	L	0.077
VAP-E26E_MEA01D24 / Meachim Creek, UT / Portions of VDH shellfish condemnation 030179C, 9/15/2022 within 179A, 12/9/1996. RPPMH	4A	Fecal Coliform	2024	L	0.013
VAP-E26E_MEA02C24 / Meachim Creek, UT / Portion of VDH Shellfish Condemnation 030-179C, 9/15/2022 not closed in 030-179A, 12/9/1996. RPPMH	4A	Fecal Coliform	2024	L	0.024

Meachim Creek				
		•	Reservoir	
Shellfishing		(Sq. Miles)	(Acres)	(Mlles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.114		

Cause Group Code: E26E-02-SF Meachim Creek

Cause Location: Described in VDH Notice and Description of Shellfish Condemnation 030-179B, 9/15/2022.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 030-179B, 9/16/2022

This area was included on the 1998 303(d) list due to VDH condemnation 179B, 12/9/1996. The impairment was addressed in the Meachim and Whiting Creek Shellfish TMDL, which was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. The impairment has subsequently expanded and contracted in multiple cycles.

During the 2012 cycle, the condemnation shrank considerably and became smaller than the TMDL study area. The open area within the TMDL study area was partially delisted (Category 2C.). The condemnation remains Category 4A.

In the 2022 cycle, the condemnation grew and is now larger than the TMDL extent; the expansion is nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MEA01B00 / Meachim Creek / Described in VDH shellfish condemnation 179B, 12/9/1996. RPPMH	4A	Fecal Coliform	1998	L	0.032
VAP-E26E_MEA02B22 / Meachim Creek, UT / Portion of VDH Shellfish Condemnation 030-179B, 9/15/2022 downstream of 030-179, 12/9/1996. RPPMH	4A	Fecal Coliform	2022	L	0.004
Meachim Creek					

Shellfishing		Estuary (Sq. Miles)	$\frac{\text{Reservoir}}{(\text{Acres})}$	
	Fecal Coliform - Total Impaired Size by Water T	ype: 0.036	, , ,	· · · ·

Cause Group Code: E26E-04-EBEN Corrotoman River

Cause Location: The mainstem Corrotoman River and its large branches within segment CRRMH.

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Estuarine Bioassessments/5A

Cause Description: During the 2014 cycle, the mainstem Corrotoman River and its large tributaries were impaired of the Aquatic Life Use due to an insufficient Chesapeake Bay Index of Biological Integrity (B-IBI).

The smaller tributaries were added to the impairment in the 2022 cycle. The impairment continued in the 2024 cycle.

Also, an impaired benthic community was noted at estuarine probabilistic monitoring station 3-CTM000.38 during monitoring in 2015. It was attributed to water quality.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BES01A98 / Bells Creek / As described in VDH shellfish condemnation 58B, 4/28/1997. CRRMH	5A	Estuarine Bioassessments	2022	L	0.055
VAP-E26E_BLD01A98 / Belwood Swamp / Tidal limit to its mouth at the Western Branch Corrotoman River. CRRMH	5A	Estuarine Bioassessments	2022	L	0.009
VAP-E26E_CRR01A00 / Corrotoman River / The mainstem of the Corrotoman River within segment CRRMH.	5A	Estuarine Bioassessments	2014	L	3.769
VAP-E26E_CTM01A00 / Eastern Branch Corrotoman River / Non-administrative portion of VDH-DSS Condemnations 021U-058B and condemnations 021U-058C & -058D, 11/15/2022. Shortened and split in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2014	L	0.429
VAP-E26E_CTM01B24 / Eastern Branch Corrotoman River, UT / Administratively condemned portion of VDH shellfish condemnation 021U-058B, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2014	L	0.007
VAP-E26E_CTM01C20 / Eastern Branch Corrotoman River / Portion of VDH shellfish condemnation 058C, 4/28/1997 that is seasonally condemned. Expanded and merged in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2014	L	0.186
VAP-E26E_CTM02A08 / Eastern Branch Corrotoman River, UT / Described in VDH Shellfish Condemnation 021U-058E, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.011

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(continued)

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM03A08 / Eastern Branch Corrotoman River / Downstream boundary of VDH condemnation 021-058C, 4/28/1997 to rivermile 0.88. CRRMH	5A	Estuarine Bioassessments	2014	L	0.348
VAP-E26E_CTM03B24 / Eastern Branch Corrotoman River / Rivermile 0.88 to mouth. CRRMH	5A	Estuarine Bioassessments	2014	L	0.410
VAP-E26E_CTO01A02 / Western Branch Corrotoman River / As described in VDH shellfish condemnation 021U-132A, 11/15/2022, not otherwise segmented. CRRMH	5A	Estuarine Bioassessments	2014	L	0.452
VAP-E26E_CTO01B12 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 included in 021U-132S64, 11/15/20202. CRRMH	5A	Estuarine Bioassessments	2014	L	0.046
VAP-E26E_CTO01C22 / Western Branch Corrotoman River / Portion of SFC 132, 4/28/1997 open in VDH-DSS condemnation 021U-132, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2014	L	0.100
VAP-E26E_CTO02A06 / Western Branch Corrotoman River / Mainstem Western Corrotoman River from rivermile 2.21 to the mouth. Split in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2014	L	0.795
VAP-E26E_CTO02B24 / Western Branch Corrotoman River / Mainstem from the downstream boundary of VDH Shellfish Condemnation 132A, 4/28/1997 to rivermile 2.21. Split in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2014	L	0.414
VAP-E26E_DAS01A02 / Davis Creek / As described in VDH-DSS condemnation 021-132S63, 11/15/2022. Segment expanded in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.043
VAP-E26E_EWE01B20 / Ewells Prong / As described in VDH shellfish condemnation 187A, 4/28/1997. Merged in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.036
VAP-E26E_EWE02A08 / Ewells Prong / Portion of VDH Shellfish Condemnation 021L-187S53, 11/15/2022 not included on 187A or 187B, 4/28/1997. Expanded in the 2024 cycle. CRRMH	$5\mathrm{A}$	Estuarine Bioassessments	2022	L	0.032

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(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_HLS01A00 / Hills Creek / As described in VDH shellfish condemnation 021U-058A, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.038
VAP-E26E_HLS01B20 / Hills Creek / The portion of VDH shellfish condemnation 58A, 4/28/1997 seasonally condemned/open (021U-058S59, 11/15/2022). CRRMH	5A	Estuarine Bioassessments	2022	L	0.024
VAP-E26E_JON01A08 / John Creek / Described in VDH-DSS Condemnation 021U-132210, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.036
VAP-E26E_JON02A08 / John Creek / Downstream of VDH-DSS condemnation 021U-132S210, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.016
VAP-E26E_LIT01A06 / Little Branch / Tidal limit to mouth at Western Branch Corrotoman River CRRMH	5A	Estuarine Bioassessments	2022	L	0.114
VAP-E26E_LOW01A08 / Lowrey Creek / Described in VDH Shellfish Condemnation 021U-132S62, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.028
VAP-E26E_MIP01A00 / Millenbeck Prong / Portion of VDH shellfish condemnation 187B, 4/28/1997 seasonally condemned in 021L-187S53, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.004
VAP-E26E_MIP01B20 / Millenbeck Prong / Described in VDH shellfish condemnation 021L-187C, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.037
VAP-E26E_MOR01B12 / Moran Creek / Described in VDH-DSS condemnation 021L-198S57, 11/15/2021. Expanded in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.045
VAP-E26E_MOR01C22 / Moran Creek, UT / Described in VDH-DSS condemnation 021L-198D, 11/15/2021. Merged in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.049
VAP-E26E_MOR02A08 / Moran Creek / Downstream of VD-DSS condemnation 021L-198, 11/15/2021. CRRMH	5A	Estuarine Bioassessments	2022	L	0.060
VAP-E26E_MYE01A00 / Myer Creek / As described in VDH shellfish condemnation 198, 4/28/1997. CRRMH	5A	Estuarine Bioassessments	2022	L	0.081

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(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MYE01B02 / Myer Creek, UT / As described in VDH-DSS condemnation 021L-198S58, 11/15/2022. Expanded in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.042
VAP-E26E_MYE01D18 / Myer Creek / Portion of VDH-DSS condemnation 021L-198A, 11/15/2022 open in 198, 4/28/1997. CRRMH	5A	Estuarine Bioassessments	2022	L	0.095
VAP-E26E_MYE03A08 / Myer Creek / Downstream of condemnations to mouth at Corrotoman River. Split in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.442
VAP-E26E_MYE03B24 / Myer Creek / Described in VDH-DSS condemnation 021L-198S56, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.028
VAP-E26E_SEN01A00 / Senior Creek / As described in VDH shellfish condemnation 021U-132B and -132C, 11/15/2022. Expanded in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.053
VAP-E26E_SEN01B20 / Senior Creek / As described in VDH shellfish condemnation 021U-132S105, 11/15/2022. Shrank in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.017
VAP-E26E_TAY01A00 / Taylor Creek / As described in VDH-DSS condemnations 021L-198B and -C, 11/15/2021. CRRMH	$5\mathrm{A}$	Estuarine Bioassessments	2022	L	0.068
VAP-E26E_TAY02A08 / Taylor Creek / Portion of VDH-DSS condemnation 205, 4/28/1997 within 021L-198S55, 11/15/2021. CRRMH	$5\mathrm{A}$	Estuarine Bioassessments	2022	L	0.123
VAP-E26E_TAY03A24 / Taylor Creek / Portion of VDH-DSS Shellfish Condemnation 021L-198S55, 11/15/2021 not included in VDH condemnation 205, 4/28/1997. CRRMH	5A	Estuarine Bioassessments	2022	L	0.011
VAP-E26E_TON01A00 / Town Creek / Described in VDH shellfish condemnation 021L-187S54, 11/15/2022. Shrank in the 2024 cycle. CRRMH	$5\mathrm{A}$	Estuarine Bioassessments	2022	L	0.018

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(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHR01A00 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187SS52, 11/15/2022. Expanded and split in the 2024 cycle. CRRMH	5A	Estuarine Bioassessments	2022	L	0.052
VAP-E26E_WHR01B24 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187A, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.045
VAP-E26E_WHR01C24 / Whitehouse Creek, UT / As described in VDH shellfish condemnation 021L-187B, 11/15/2022. CRRMH	5A	Estuarine Bioassessments	2022	L	0.005
VAP-E26E_ZZZ02A14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA70 CRRMH	5A	Estuarine Bioassessments	2022	L	0.091
VAP-E26E_ZZZ02C14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA72 CRRMH	5A	Estuarine Bioassessments	2022	L	0.445

Corrotoman River			
A		Reservoir	
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Estuarine Bioassessments - Total Impaired Size by Water Type:	9.204		

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E26E-05-SF Myer Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 021L-198A, 11/15/2022.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 021L-198A, 11/15/2022

A portion of Myer Creek was included on the 1998 303(d) list due to VDH-DSS Condemnation 198, 4/28/1997. The TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

The condemnation has since varied in size.

In the 2022 cycle, the area expanded and is larger than the 1997 condemnation. The expansion is nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MYE01A00 / Myer Creek / As described in VDH shellfish condemnation 198, 4/28/1997. CRRMH	4A	Fecal Coliform	1998	L	0.081
VAP-E26E_MYE01D18 / Myer Creek / Portion of VDH-DSS condemnation 021L-198A, 11/15/2022 open in 198, 4/28/1997. CRRMH	4A	Fecal Coliform	2022	L	0.095

Myer Creek				
		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.176		

Cause Group Code: E26E-08-SF Senior Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 021U-132B and -132C, 11/15/2022.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnations 021U-132B and -132C, 11/15/2022

Senior Creek was included on the 1998 303(d) list due to VDH condemnation 132B, 4/28/1997. The TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

The condemnation shrank in the 2020 cycle and a portion is now seasonally condemned (021-132S105, 10/31/2018) and was partially delisted. A second cove closed again in the 2024 cycle.

The condemned portion is considered Category 4A; the seasonal closure is Category 2C/2B.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_SEN01A00 / Senior Creek / As described in VDH shellfish condemnation 021U-132B and -132C, 11/15/2022. Expanded in the 2024 cycle. CRRMH	4A	Fecal Coliform	1998	L	0.053

Senior Creek

		Estuary	$\operatorname{Reservoir}$	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.053			

Cause Group Code: E26E-09-SF Western Branch Corrotoman River

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 132A, 4/28/1997.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 021U-132A, 11/15/2022

A portion of the Western Branch Corrotoman River was included on the 1998 303(d) list due to VDH condemnation 132A, 4/28/1997. The condemnation has subsequently shortened several times. In the 2020 cycle, the closure contracted again and split (021-132A and -C, 10/31/2018). It re-merged in the 2022 cycle.

The TMDL was completed for the 1998 boundary; it was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. The condemned portion is considered Category 4A; the open portion is considered Category 2C.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BLD01A98 / Belwood Swamp / Tidal limit to its mouth at the Western Branch Corrotoman River. CRRMH	4A	Fecal Coliform	2002	L	0.009
VAP-E26E_CTO01A02 / Western Branch Corrotoman River / As described in VDH shellfish condemnation 021U-132A, 11/15/2022, not otherwise segmented. CRRMH	4A	Fecal Coliform	1998	L	0.452
VAP-E26E_LIT01A06 / Little Branch / Tidal limit to mouth at Western Branch Corrotoman River CRRMH	4A	Fecal Coliform	1998	L	0.114

Western Branch Corrotoman River

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.574		

Cause Group Code: E26E-10-SF Bush Park Creek / Rappahannock River

Cause Location: Portion of VDH Notice and Description of Shellfish Condemnation 032-109A, 12/10/2009 that is not administratively condemned.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 032-109A, 9/15/2022

Bush Park Creek was included on the 1998 303(d) list as impaired of the Shellfish Consumption Use due to VDH condemnation 109, 4/27/1989. The TMDL for this area was approved by the EPA on 6/7/2006 and by the SWCB on 6/27/2007.

In the 2022 cycle, the condemnation merged with the condemnation on neighboring Woods Creek (administrative) and expanded into a portion of the Rappahannock River. The expansion is nested.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BPC01A98 / Bush Park Creek / As delineated in VDH shellfish condemnation 109, 4/27/1989. RPPMH	4A	Fecal Coliform	1998	L	0.103
VAP-E26E_RPP08A22 / Rappahannock River / Portion of VDH-DSS Condemnation 032-109A, 9/15/2022 that is within the mainstem Rappahannock River. Shrank in the 2024 cycle. RPPMH	4A	Fecal Coliform	2022	L	0.002
Bush Park Creek / Rappahannock River				·	

Shellfishing		Estuary (Sq. Miles)	Reservoir (Acres)	
_	Fecal Coliform - Total Impaired Size by Water Type:	0.104		

Cause Group Code: E26E-11-SF Mill Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 103, 12/10/1991.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 031-102A, 9/15/2021

A portion of Mills Creek was impaired in the 1998 cycle due to VDH condemnation 103, 12/10/1991. The TMDL for this segment was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007. The segment is considered Category 4A.

However, during the 2012 cycle, the condemnation retracted and became smaller than the TMDL study area. The open area within the TMDL study area was partially delisted (Category 2C.)

The condemnation has continued to vary in size, but remains smaller than the TMDL extent.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MLL01A98 / Mill Creek / Described in VDH shellfish condemnation 031-102A, 9/15/2021. Expanded slightly in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.111

Mill Creek

Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	(Sq. Miles)	Reservoir (Acres)	

Cause Group Code: E26E-12-SF Sturgeon Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 104, 11/28/1994

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH condemnation 032-104A, 9/15/2022

A portion of Sturgeon Creek was included on the 1998 303(d) list due to VDH shellfish condemnation 104, 11/28/1994. The TMDL was approved by the EPA on 6/7/2006 and by the SWCB on 6/27/2007.

The condemnation has subsequently shrunk and open/seasonally condemned portions were partially delisted. In the 2022 cycle, the condemnation grew but remained smaller than the TMDL extent; the residual was seasonally condemned (Cat 2C/2B).

The condemnation expanded to the TMDL extent again in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_STE01A98 / Sturgeon Creek / As delineated in VDH shellfish condemnation 104, 11/28/1994. Merged in the 2024 cycle. RPPMH	4A	Fecal Coliform	1998	L	0.085

Sturgeon Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.085		

Cause Group Code: E26E-13-BAC Locklies Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 102, 10/31/1994

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, Locklies Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 2/12 at 3-LOL000.77.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

As this impairment is within the study area for the Locklies and Mill Creek Shellfish TMDL, which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007, the impairment is considered nested (Category 4A.)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_LOL01A02 / Locklies Creek / Delineated in VDH shellfish condemnation 102, 10/31/1994. Merged in the 2024 cycle. RPPMH	4A	Enterococcus	2012	L	0.101

Locklies Creek

Recreation	Enterococcus - Total Impaired Size by Water Type:	(Sq. Miles)	Reservoir (Acres)	
	I J J J I			

Cause Group Code: E26E-13-SF Locklies Creek

Cause Location: Described in VDH Notice and Description of Shellfish Condemnation 102,10/31/1994

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 031-102B, 9/15/2021

Locklies Creek was included on the 1998 303(d) list due to VDH condemnation 102, 4/13/1993. The Locklies Creek Shellfish TMDL was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007; the TMDL was based on the extent of condemnation 102, 10/31/1994.

During the 2012 cycle, the condemnation retracted and a portion of the TMDL study area was included in seasonal condemnation 031-102M1. The seasonally condemned segment was partially delisted (Category 2C); the condemned area is considered a Category 4A water.

The condemnation grew slightly during the 2022 cycle, but remained smaller than the TMDL study area.

It expanded to the TMDL extent again in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water
VAP-E26E_LOL01A02 / Locklies Creek / Delineated in VDH shellfish condemnation 102, 10/31/1994. Merged in the 2024 cycle. RPPMH	4A	Fecal Coliform		1998	L	0.101
Locklies Creek Shellfishing Fecal Coliform - Total I	mpaired Size	by Water Type:	Estuary (Sq. Miles) 0.101	Rese (Ac		River (Miles)

Cause Group Code: E26E-14-SF Hills Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 58A, 4/25/1997

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 021U-058A, 11/15/2022

Hills Creek was included on the 1998 303(d) list due to VDH-DSS Condemnation 58A, 4/28/1997. The TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

During the 2020 cycle, the condemnation shrank and a portion is now seasonally condemned and was partially delisted (021-058S59, 10/31/2018). The closed segment is considered Category 4A and the seasonal area is Category 2C/2B

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_HLS01A00 / Hills Creek / As described in VDH shellfish condemnation 021U-058A, 11/15/2022. CRRMH	4A	Fecal Coliform	1998	L	0.038

Hills Creek

Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 0.038	
	recar comorni - rotar imparted size by water rype.	0.000	

Cause Group Code: E26E-16-SF Eastern Branch Corrotoman River

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 58C, 4/28/1997.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnations 021U-058B, -C, and - D, 11/15/2022

The Eastern Branch Corrotoman River was included on the 1998 303(d) list due to VDH condemnation 58C, 4/28/1997. The TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. The condemnation subsequently shortened. The condemned area is considered Category 4A waters; the open area was previously partially delisted and is Category 2C.

The condemned area has expanded and contracted several times.

During the 2024 cycle, a portion of condemnation 021U-132B converted to administratively condemned and will be partially delisted.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM01A00 / Eastern Branch Corrotoman River / Non-administrative portion of VDH-DSS Condemnations 021U-058B and condemnations 021U-058C & -058D, 11/15/2022. Shortened and split in the 2024 cycle. CRRMH	4A	Fecal Coliform	1998	L	0.429

Eastern Branch Corrot	oman River			
		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.429		

Cause Group Code: E26E-17-SF Eastern Branch Carter Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 041C, 11/1/1996.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 020-041A, 10/15/2022

A portion of Eastern Branch Carters Creek was assessed as impaired of the Shellfish Use during the 1998 303(d) cycle due to VDH condemnation 41C, 11/1/1996. Although the segment has expanded several times, the TMDL was completed only for the original segment. It was approved by the EPA on 9/20/2007 and by the SWCB on 7/31/2008. The original segment is considered Category 4A; the expansion is addressed in fact sheet E26E-46-SF.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDI Dev. Priori	Water
VAP-E26E_CEB01A00 / Eastern Branch Carter Creek / Described in VDH shellfish condemnation 041C, 11/1/1996. RPPMH	4A	Fecal Coliform		1998	L	0.084
Eastern Branch Carter Creek Shellfishing Fecal Coliform - Total I	mpaired Size	by Water Type:	Estuary (Sq. Miles) 0.084	Rese (Ac		River (Miles)

Cause Group Code: E26E-21-SF XII - Windmill Point, UT (aka White Marsh)

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 018-053B, 7/23/2018

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS condemnation 018-053B, 7/23/2018

The impairment is nested in the shellfish TMDL for Oyster Creek, which was approved by the EPA on 4/15/2009 and by the SWCB on 7/27/2009. It is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_XII01A18 / XII - Windmill Point Creek, UT (aka White Marsh) / Described in VDH-DSS condemnation 018-053B, 7/23/2018 RPPMH	4A	Fecal Coliform	2018	L	0.034

XII - Windmill Point, UT (aka White Marsh)

		Estuary	Reservoir	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.034			

Cause Group Code: E26E-24-BAC Whiting Creek

Cause Location: Tidal Whiting Creek as described in VDH Shellfish Condemnation 029-051A, 9/1/2015

Cause City/County: Middlesex County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/5A

Cause Description: During the 2012 cycle, Whiting Creek was impaired of the Recreation Use due to an enterococci exceedance rate of 3/19 at 3-WHS000.89.

Although Whiting Creek is administratively condemned by VDH and the Shellfish Use is considered removed, the TMDL was completed and was approved by the EPA on 11/15/2005 and by the SWCB on 9/27/2006. However, the TMDL did not include a nearby VPDES discharger; therefore, the Recreation Use cannot be considered nested.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHS01B00 / Whiting Creek / As delineated in VDH shellfish condemnation 029-051A, 9/1/2015. RPPMH	5A	Enterococcus	2012	L	0.196

Whiting Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.196		

Sources: Source Unknown

Cause Group Code: E26E-26-BAC Little Branch

Cause Location: Little Branch from its tidal limit to its mouth at the Western Branch Corrotoman River

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Little Branch was assessed as not supporting of the Recreation Use during the 2006 cycle due to enterococci exceedances at 3-LIT000.85, which is located at a private dock off Route 620. The segment remained impaired during the 2010 cycle; the violation rate was 3/11.

The area was addressed in the Western Branch Corrotoman River Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. Because the bacterial standard for the Shellfish Use is more stringent than the standard for the Recreation Use, the impairment is considered to be nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_LIT01A06 / Little Branch / Tidal limit to mouth at Western Branch Corrotoman River CRRMH	4A	Enterococcus	2006	L	0.114

Little Branch

Reservoir	River
s) (Acres)	(Miles)
	es) (Acres)

Cause Group Code: E26E-27-BAC Belwood Swamp

Cause Location: Tidal Belwood Swamp

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: Riverine Belwood Swamp was initially assessed in 1998 as fully supporting but threatened of the Recreation Use based on exceedances of the fecal coliform standard at monitoring station 3-BLD000.58, located at the Route 3 bridge. During the year 2002 cycle, the segment was downgraded to impaired.

However, in the 2006 cycle, it was determined that the station is tidally influenced. The station remained impaired for fecal coliform and the fact sheet and AU were renamed. The TMDL was due in 2014. There had been no enterococci monitoring at this site; therefore, the fecal coliform impairment was carried over.

The area was addressed in the Western Branch Corrotoman River Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. Because the bacterial standard for the Shellfish Use is more stringent than the standard for the Recreation Use, the impairment is considered to be nested (Category 4A).

Additional monitoring was conducted during the 2012 cycle. The impairment converted to enterococci due to an exceedance rate of 8/12 at 3-BLD000.58.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_BLD01A98 / Belwood Swamp / Tidal limit to its mouth at the Western Branch Corrotoman River. CRRMH	4A	Enterococcus	2012	L	0.009

Belwood Swamp

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.009		

Cause Group Code: E26E-28-BAC Western Branch Corrotoman River

Cause Location: The Western Branch Corrotoman River from its tidal limit to the downstream extent of VDH-DSS condemnation 021U-132A, 11/15/2022

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Enterococcus/4A

Cause Description: During the 2012 cycle, the upper portion of the Western Branch Corrotoman River was impaired of the Recreation Use due to an enterococci exceedance rate of 7/12 at 3-CTO007.51, which is located off of Route 3.

The area was already addressed in the Corrotoman River Watershed Shellfish Bacterial TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. The condemned portion is considered nested (Category 4A).

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTO01A02 / Western Branch Corrotoman River / As described in VDH shellfish condemnation 021U-132A, 11/15/2022, not otherwise segmented. CRRMH	4A	Enterococcus	2012	L	0.452

Western Branch Corrotoman River

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Enterococcus - Total Impaired Size by Water Type:	0.452		

Cause Group Code: E26E-42-SF Hunting Creek

Cause Location: Described in VDH Notice and Description of Shellfish Condemnation 032-104B, 9/15/2022.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 032-104B, 9/15/2022

The Hunting Creek shellfish impairment is nested in the nearby Sturgeon Creek Shellfish TMDL, which was approved by the EPA on 6/7/2006 and by the SWCB on 6/27/2007.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Prior	Water
VAP-E26E_HNU01A08 / Hunting Creek / Described in VDH-DSS Condemnation 032-104B, 9/15/2022. RPPMH	4A	Fecal Coliform		2008	L	0.021
Hunting Creek Shellfishing Fecal Coliform - Total 1	Impaired Size	by Water Type:	Estuary (Sq. Miles) 0.02		ervoir eres)	River (Miles)

Cause Group Code: E26E-46-SF Eastern Branch Carter Creek

Cause Location: Portion of VDH condemnation 020-041A, 10/15/2022 not included on condemnation 41, 11/1/1996

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: Portion of VDH shellfish condemnation 020-041A, 10/15/2022

A portion of Eastern Branch Carter Creek was assessed as impaired of the Shellfish Use during the 1998 303(d) cycle due to VDH condemnation 41C, 11/1/1996. Although the segment has expanded several times, the TMDL was completed only for the original segment. The TMDL due date for this downstream portion was 2014 since it first expanded during the 2002 cycle.

It expanded further in the 2018 cycle.

The closure shrank and split in the 2020 cycle (portion of 020-041A, 1/8/2018 and 020-041C, 11/8/2018).

In the 2022 cycle, condemnation A expanded, however condemnation C converted to administratively condemned on 11/15/2020 and was partially delisted.

The condemnation shrank again in the 2024 cycle; Jacks Cove converted to seasonally condemned and will be partially delisted.

It is considered nested in the upstream Eastern Branch Carter Creek Shellfish TMDL, which was approved by the EPA on 9/20/2007.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CEB01B08 / Eastern Branch Carter Creek / Portion of VDH shellfish condemnation 020-041A, 10/15/2022 not included in 041C, 11/1/1996. Shrank and split in the 2024 cycle. RPPMH	4A	Fecal Coliform	2002	L	0.072

Eastern Branch Carter	Creek				
		· · · · · ·	Reservoir		
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.072			

Cause Group Code: E26E-55-SF Mosquito Creek

Cause Location: As described in VDH Notice and Description of Shellfish Condemnation 018-203A, 1/6/2005

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS Condemnation 018-203A, 11/15/2020

Mosquito Creek was included on the 1998 303(d) list due to VDH Condemnation 203, 11/22/1996. The Mosquito Creek Shellfish TMDL was approved by the EPA on 4/15/2009 and by the SWCB on 7/27/2009. The TMDL was based on the 1/6/2005 condemnation, as that had been the largest condemnation to date.

The condemnation was subsequently rescinded several times.

A portion was relisted in the 2020 cycle (018-203A, 10/28/2014).

In the 2022 cycle, the condemned area expanded to the TMDL extent. It is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MOS01C20 / Mosquito Creek / As delineated in VDH shellfish condemnation 018-203, 1/6/2005. RPPMH	4A	Fecal Coliform	2020	L	0.069

Mosquito Creek

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
_	Fecal Coliform - Total Impaired Size by Water Type:	0.069		

Cause Group Code: E26E-56-SF Roane Cove of Locklies Creek

Cause Location: Described in VDH Shellfish Condemnation 031-102C, 9/15/2021.

Cause City/County: Middlesex County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH Shellfish Condemnation 031-102C, 9/15/2021

Roane Cove was listed in the 2022 cycle. The impairment is nested in the Locklies and Mill Creeks Shellfish TMDL, which was approved by the EPA on 8/2/2006 and by the SWCB on 6/27/2007.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_LOL03A08 / Roane Cove of Locklies Creek / Described in VDH-DSS Shellfish Condemnation 031-102C, 9/15/2021. RPPMH	4A	Fecal Coliform	2022	L	0.034

Roane Cove of Locklies Creek

		$\operatorname{Estuary}$	$\operatorname{Reservoir}$	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.034			

Cause Group Code: E26E-57-EBTOX Eastern Branch Carter Creek

Cause Location: Described in VDH shellfish condemnation 041C, 11/1/1996.

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Sediment Bioassay/5A

Cause Description: During the 2022 cycle, upper Eastern Branch Carter Creek and its tidal tributaries were assessed as impaired of the Aquatic Life Use due to the weight-of-evidence assessment at estuarine probabilistic monitoring station 3-DUN000.08 (Category 5A, scenario 1). The station is located on Dunton Cove.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMD Dev. Priori	Water Size
VAP-E26E_CEB01A00 / Eastern Branch Carter Creek / Described in VDH shellfish condemnation 041C, 11/1/1996. RPPMH	5A	Sediment Bioass	ay	2022	L	0.084
Eastern Branch Carter Creek Aquatic Life Sediment Bioassay - Total I	Impaired Size	by Water Type:	Estuary (Sq. Miles) 0.084		rvoir res)	River (Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E26E-60-SF Eastern Branch Corrotoman River, UT

Cause Location: Described in VDH-DSS Condemnation 021U-058E, 11/15/2022.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS Condemnation 021U-058E, 11/15/2022

The area was listed in the 2022 cycle. It is considered nested in the nearby Eastern Branch Corrotoman River Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM02A08 / Eastern Branch Corrotoman River, UT / Described in VDH Shellfish Condemnation 021U-058E, 11/15/2022. CRRMH	4A	Fecal Coliform	2022	L	0.011

Eastern Branch Corrotoman River, UT

		Estuary	$\operatorname{Reservoir}$	River	
Shellfishing		(Sq. Miles)	(Acres)	(Miles)	
	Fecal Coliform - Total Impaired Size by Water Type:	0.011			

Cause Group Code: E26E-61-SF Moran Creek, UT

Cause Location: Described in VDH-DSS condemnation 021L-198D, 11/15/2021.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS condemnation 021L-198D, 11/15/2021

Moran Creek was impaired of the Shellfish Use in the 2022 cycle. It is nested within the nearby Taylor Creek Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. It expanded in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDI Dev. Priorit	Water
VAP-E26E_MOR01C22 / Moran Creek, UT / Described in VDH-DSS condemnation 021L-198D, 11/15/2021. Merged in the 2024 cycle. CRRMH	4A	Fecal Coliform		2022	L	0.049
Moran Creek, UT Shellfishing Fecal Coliform - Total In	mpaired Size	by Water Type:	Estuary (Sq. Miles) 0.049	Rese (Ac		River (Miles)

Cause Group Code: E26E-62-SF Taylor Creek

Cause Location: Described in VDH-DSS Condemnations 021L-198B and -C, 11/15/2021

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS Condemnations 021L-198B and -C, 11/15/2021

A large portion of Taylor Creek was included on the 1998 303(d) list due to VDH condemnation 205, 4/28/1997. The Taylors Creek TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

The condemnation has been listed and re-listed in several cycles. Currently, two coves of Taylor Creek are impaired (Category 4A). The remainder is Category 2C or 2C/2B.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_TAY01A00 / Taylor Creek / As described in VDH-DSS condemnations 021L-198B and -C, 11/15/2021. CRRMH	4A	Fecal Coliform	2022	L	0.068

Taylor	Creek
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Shellfishing	Fecal Coliform - Total Impaired Size by Water Type:	(Sq. Miles)	Reservoir (Acres)	
	Fecal Coliform - Total Impaired Size by Water Type:		()	(

Cause Group Code: E26E-63-EBTOX Western Branch Corrotoman River

Cause Location: Approximately 1/2 mile upstream and downstream of station 3-CTO002.71

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Sediment Bioassay/5A

Cause Description: During the 2024 cycle, 2021 estuarine probabilistic monitoring station 3-CTO002.71 was impaired of the Aquatic Life Use due to weight of evidence analysis (scenario 8).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTO02B24 / Western Branch Corrotoman River / Mainstem from the downstream boundary of VDH Shellfish Condemnation 132A, 4/28/1997 to rivermile 2.21. Split in the 2024 cycle. CRRMH	5A	Sediment Bioassay	2024	L	0.414
Western Branch Corrotoman River			D	·	

		Estuary	Reservoir	River	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	Sediment Bioassay - Total Impaired Size by Water Type:	0.414	. ,	· · · ·	

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E26E-64-EBTOX Eastern Branch Corrotoman River

Cause Location: Eastern Branch Corrotoman River from rivermile 0.88 to the mouth.

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Sediment Bioassay/5A

Cause Description: An impaired benchic community was noted at estuarine probabilistic monitoring station 3-CTM000.38 during monitoring in 2015. Weight of Evidence analysis assigned it to Category 5A, scenario 8. In previous cycles, it was included within the Chesapeake Bay B-IBI impairment, but it is being split out in the 2024 cycle as its own impairments since scenario 8 indicates possible chemical causes. The TMDL due date will be 2030 to reflect the earliest impairment.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CTM03B24 / Eastern Branch Corrotoman River / Rivermile 0.88 to mouth. CRRMH	5A	Sediment Bioassay	2024	L	0.41

Eastern Branch Corrotoman River

		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Sediment Bioassay - Total Impaired Size by Water Type:	0.41		

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Cause Group Code: E26E-65-SF Millenbeck Prong

Cause Location: Described in VDH shellfish condemnation 021L-187C, 11/15/2021.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH-DSS condemnation 021L-187C, 11/15/2022

Millenbeck Prong was included on the 1998 303(d) list due to VDH condemnation 187B, 4/28/1997. The TMDL was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

During the 2016 cycle, Millenbeck Prong was entirely re-opened to shellfish harvest and was delisted (Category 2C).

A portion re-closed to harvest in the 2024 cycle.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MIP01B20 / Millenbeck Prong / Described in VDH shellfish condemnation 021L-187C, 11/15/2022. CRRMH	4A	Fecal Coliform	2024	L	0.037

Millenbeck Prong				
		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.037		

Cause Group Code: E26E-66-SF Whitehouse Creek

Cause Location: As described in VDH shellfish condemnation 021L-187A, 11/15/2022.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 021L-187A, 11/15/2022

Whitehouse Creek was impaired of the Shellfish Consumption Use in the 2024 cycle. The condemnation is considered nested in the nearby Ewells Prong Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	TMDL Dev. Priorit	Water
VAP-E26E_WHR01B24 / Whitehouse Creek / As described in VDH shellfish condemnation 021L-187A, 11/15/2022. CRRMH	4A	Fecal Coliform		2024	L	0.045
Whitehouse Creek Shellfishing			Estuary (Sq. Miles)			River (Miles)

Sources: Agriculture; Non-Point Source; On-site Treatment Systems (Septic Systems and Similar Decentralized Systems); Urban Runoff/Storm Sewers

0.045

Fecal Coliform - Total Impaired Size by Water Type:

Cause Group Code: E26E-67-SF Whitehouse Creek, UT

Cause Location: As described in VDH shellfish condemnation 021L-187B, 11/15/2022.

Cause City/County: Lancaster County

Use(s): Shellfishing

Causes(s)/VA Category: Fecal Coliform/4A

Cause Description: VDH shellfish condemnation 021L-187B, 11/15/2022

The tributary was impaired of the Shellfish Consumption Use in the 2024 cycle. The condemnation is considered nested in the nearby Ewells Prong Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

Assessment Unit / Water Name / Location Desc.	cause 1			Dev. Priority	Water Size
VAP-E26E_WHR01C24 / Whitehouse Creek, UT / As described in VDH shellfish condemnation 021L-187B, 11/15/2022. CRRMH 4A	Fecal C	oliform	2024	L	0.005

Whitehouse Creek, UT

		Estuary	Reservoir	River
Shellfishing		(Sq. Miles)	(Acres)	(Miles)
	Fecal Coliform - Total Impaired Size by Water Type:	0.005		

Cause Group Code: E26R-01-BAC Belwood Swamp and Tributaries

Cause Location: Belwood Swamp and tributaries from its headwaters to its tidal limit.

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2012 cycle, the nontidal Belwood Swamp watershed was impaired of the Recreation Use due to E. coli exceedances at Belwood Swamp at station 3-BLD001.54 and McMahon Swamp at 3-MCM000.96. In the 2014 cycle, the exceedance rates were 9/27 and 6/12, respectively.

Additional monitoring was conducted at 3-BLD001.54 in the 2020 cycle; the E. coli exceedance rate was 5/11.

New bacteria criteria were implemented in the 2022 cycle. No new data were collected at 3-BLD001.54, but re-analysis of the previous data confirms that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples. No additional data have been collected at 3-MCM000.96 either; therefore, the impairment is carried over at that station.

The area drains to tidal Belwood Swamp, which was included in the Corrotoman River Shellfish Bacterial TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. Implementation of the TMDL is expected to address the nontidal area; therefore, the impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26R_BLD01A08 / Belwood Swamp and Tributaries / Watershed from its headwaters to tidal limit	4A	Escherichia coli (E. coli)	2012	L	24.54

Belwood Swamp and Tributaries

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	· - /	· · · · ·	24.54

Cause Group Code: E26R-03-DO Norris Prong

Cause Location: Norris Prong from its headwaters to its tidal limit.

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: During the 2008 cycle, Norris Prong was considered impaired of the Aquatic Life Use based on a dissolved oxygen exceedance rate of 4/10 at the Route 3 bridge (3-NOR001.00).

No additional data has been collected.

Natural causes are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26R_NOR01A08 / Norris Prong / Headwaters to tidal limit	$5\mathrm{C}$	Dissolved Oxygen	2008	L	2.47

Norris Prong					
		Estuary	Reservoir	River	
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)	
	Dissolved Oxygen - Total Impaired Size by Water Type:			2.47	

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E26R-04-BAC Browns Creek

Cause Location: Browns Creek from its headwaters to its tidal limit.

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2008 cycle, Browns Creek was considered impaired of the Recreation Use based on E. coli exceedances at the Route 614 bridge (3-BON001.65).

The exceedance rate was 5/24 in the 2014 cycle.

Additional monitoring was conduced in the 2020 cycle; the exceedance rate was 4/11.

New bacteria criteria were implemented in the 2022 cycle. No new data were collected, but re-analysis of the previous data confirms that the segment remains impaired due to two or more STV exceedances in the same 90-day period with <10 samples.

The impairment is considered nested (Category 4A) because it is located within the watershed study area for the Corrotoman River Shellfish TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26R_BON01A08 / Browns Creek / Headwaters to tidal limit	4A	Escherichia coli (E. coli)	2008	L	2.59

Browns Creek

		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:	, <u> </u>	. ,	2.59

Cause Group Code: E26R-04-DO Browns Creek

Cause Location: Browns Creek from its headwaters to its tidal limit.

Cause City/County: Lancaster County

Use(s): Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/5C

Cause Description: Browns Creek has been impaired of the Aquatic Life Use since the 2008 cycle based on dissolved oxygen exceedances at the Route 614 bridge (3-BON001.65).

The violation rate was 3/22 during the 2024 cycle.

Natural conditions are suspected.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26R_BON01A08 / Browns Creek / Headwaters to tidal limit	$5\mathrm{C}$	Dissolved Oxygen	2008	L	2.59

Browns Creek

Aquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles)	Reservoir (Acres)		
	Dissolved Oxygen - Total Impaired Size by Water Type:			2.59	

Sources: Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

Cause Group Code: E26R-05-BAC Little Branch

Cause Location: Nontidal Little Branch below Blakemore Millpond

Cause City/County: Lancaster County

Use(s): Recreation

Causes(s)/VA Category: Escherichia coli (E. coli)/4A

Cause Description: During the 2014 cycle, the segment was impaired of the Recreation Use due to an E. coli exceedance rate of 5/12 at station 3-LIT001.89, which is located on Little Branch at Route 201.

New bacteria criteria were implemented in the 2022 cycle. No additional data have been collected; therefore, the impairment is carried over.

The watershed was addressed in the Western Branch Corrotoman River Shellfish Bacterial TMDL, which was approved by the EPA on 1/23/2008 and by the SWCB on 7/31/2008. Implementation of the TMDL is expected to address the nontidal area; therefore, the impairment is considered nested (Category 4A).

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26R_LIT01A14 / Little Branch / Blakemore Millpond dam downstream to its tidal limit	4A	Escherichia coli (E. coli)	2014	L	0.64

Recreation		Estuary (Sq. Miles)	$\begin{array}{c} \text{Reservoir} \\ \text{(Acres)} \end{array}$	
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			0.64

Sources: Agriculture; Non-Point Source; On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)

Rappahannock River Basin

Cause Group Code: RPPMH-DO-BAY Rappahannock River

Cause Location: The mesohaline Rappahannock River and tidal tributaries.

Cause City/County: Essex County; Lancaster County; Middlesex County; Richmond County

Use(s): Aquatic Life; Deep-Channel Seasonal Refuge; Deep-Water Aquatic Life; Migratory Fish Spawning and Nursery; Open-Water Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/4A; Dissolved Oxygen/4D

Cause Description: The mainstem of the Rappahannock River from Myrtle Swamp to its mouth was originally listed in 1998 by DEQ due to dissolved oxygen exceedances and nutrient overenrichment. The EPA extended the segment upstream to the confluence with Totuskey Creek. In the 2004 cycle, dissolved oxygen exceedances were noted in deep water and deep channel stations downstream of the confluence with Lancaster Creek (Morattico), which is further downstream.

The Chesapeake Bay Water Quality Standards were implemented during the 2006 cycle. During the 2014 cycle, the mesohaline portion of the Rappahannock failed the Chesapeake Bay Open Water Subuse's summer 30-day mean dissolved oxygen criterion. Applicable areas also failed the Deep Water 30-day mean dissolved oxygen criteria and the Deep Channel Subuse's instantaneous minimum dissolved oxygen criteria.

RPPMH passed both Open Water Subuse 30-day mean criterion as well as the Deep Water summer 30-day mean criterion in the 2016 cycle; these areas were delisted in the tributaries (Category 2C). However, due to EPA rules, areas included on the 1998 EPA overlist for dissolved oxygen must remain listed until all dissolved oxygen criteria can be assessed. This includes the Rappahannock River mainstem from Totuskey Creek to the mouth as well as the tidal Corrotoman River. These areas were considered Category 4D.

In the 2018 cycle, Deep Water areas also were impaired.

In the 2020 cycle, the Deep Water and Deep Channel Subuses continued to be impaired in applicable areas. In addition, the Open Water Use also failed. The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, all areas are considered Category 4A.

The Open Water summer-, Deep Water-, and Deep Channel criteria continued to fail in the 2022 cycle. The Open Water rest-of-year criteria was met and there was insufficient information to assess the other frequencies.

In the 2024 cycle, both Open Water 30-day mean criteria passed, as well as the Deep Channel Subuse. Areas outside the Consent Decree extent will be partially delisted. The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, it is considered Category 4D/Category 2C as appropriate.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP01B98 / Rappahannock River: Garrett's Marina / As delineated in VDH shellfish condemnation 026-181M1, 3/15/2021. RPPMH	4D	Dissolved Oxygen	1998	L	0.025
VAP-E24E_RPP01C06 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/16/2007 (non-admin) that is open 3/15/2021. Shrank in the 2024 cycle. RPPMH	4D	Dissolved Oxygen	1998	L	0.327
VAP-E24E_RPP01D10 / Rappahannock River / The portion of the Rappahannock River within the administratively closed area of VDH shellfish condemnation 025-071A, 3/15/2021.	4D	Dissolved Oxygen	1998	L	0.137

RPPMH

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP01E18 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/15/2021 (non-admin) Expanded in the 2024 cycle. RPPMH	4D	Dissolved Oxygen	1998	L	0.378
VAP-E24E_RPP03A00 / Rappahannock River / The Rappahannock River from the limit of VDH shellfish condemnation 068A, 11/14/2005 downstream to end of MSN (Sharps/0.7 mi DS of Mark Haven Beach) unless otherwise segmented RPPMH	4D	Dissolved Oxygen	1998	L	10.858
VAP-E24E_RPP03D24 / Rappahannock River / Described in VDH shellfish condemnation 026-270A, 5/25/2022. RPPMH	4D	Dissolved Oxygen	1998	L	0.061
VAP-E25E_DEE01A04 / Deep Creek / Described in VDH shellfish condemnation 121, 11/16/1994. RPPMH	4A	Dissolved Oxygen	2020	NA	0.049
VAP-E25E_DEE01B08 / Deep Creek / Portions of VDH-DSS condemnations 023-121B, 12/15/2021 not included in the 11/16/1994 condemnation. Merged and adjusted in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.182
VAP-E25E_GEE01A98 / Greenvale Creek / As delineated in VDH shellfish condemnation 094, 11/7/1994. RPPMH	4A	Dissolved Oxygen	2020	NA	0.087
VAP-E25E_GEE02A06 / Greenvale Creek / Portion of VDH-DSS condemnation 022-094A, 11/15/2022 not included in 94, 11/7/1994. RPPMH	4A	Dissolved Oxygen	2020	NA	0.012
VAP-E25E_GEE02B10 / Belmont Creek / Described in VDH-DSS Shellfish Condemnation 022-094S108, 11/15/2022. Shrank in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.024
VAP-E25E_HRY01A06 / Harry George Creek / Tidal limit to mouth at the Rappahannock River RPPMH	4A	Dissolved Oxygen	2020	NA	0.095
VAP-E25E_LAN01A98 / Lancaster Creek / As delineated in VDH-DSS Shellfish Condemnation 023-120A, 8/14/1995. RPPMH	4A	Dissolved Oxygen	2020	NA	0.270
VAP-E25E_LAN01B08 / Lancaster Creek / The portion of VDH Shellfish Condemnation 023-120A, 9/22/2022 not included in 120A, 8/14/1995. Merged and expanded in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.489

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LAN02A02 / Lancaster Creek / Lancaster Creek downstream of VDH Shellfish Condemnation 023-120, 9/22/2022, not otherwise segmented. Size reduced in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.823
VAP-E25E_LAN03A06 / Lancaster Creek / Described in VDH Shellfish Condemnation 023-120C, 9/22/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.023
VAP-E25E_LGG01A98 / Lagrange Creek / As described in VDH Shellfish Condemnation 028-127A, 1/15/2022. Expanded in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.562
VAP-E25E_LGG01B18 / Lagrange Creek / Portion of VDH Shellfish Condemnation 127, 6/11/1996 open on 028-127, 1/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.037
VAP-E25E_LGG02A06 / Lagrange Creek / Lagrange Creek downstream of VDH Shellfish Condemnation 127, 6/11/1996 RPPMH	4A	Dissolved Oxygen	2020	NA	0.040
VAP-E25E_MTT01A00 / Morattico Creek / Delineated in VDH Shellfish Condemnation 023-120B, 9/22/2022. Expanded in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.345
VAP-E25E_MUB01A02 / Mulberry Creek / Described in VDH shellfish condemnation 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.148
VAP-E25E_MUB02A06 / Mulberry Creek / Portion of VDH shellfish condemnation 023-121A, 12/15/2021 not included in 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.100
VAP-E25E_MUC01A04 / Mud Creek / Described in VDH-DSS Condemnation 027-090B, 8/26/2008. RPPMH	4A	Dissolved Oxygen	2020	NA	0.204
VAP-E25E_PRR01A02 / Parrotts Creek / Delineated in VDH shellfish condemnation 090, 4/27/1989. RPPMH	4A	Dissolved Oxygen	2020	NA	0.153
VAP-E25E_PRR02A08 / Parrotts Creek / Portion of VDH-DSS Condemnation 027-090A, 1/15/2022 downstream of condemnation 090, 4/27/1989. RPPMH	4A	Dissolved Oxygen	2020	NA	0.011

(continued)					
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS01A00 / Robinson Creek / Described in VDH shellfish condemnation 028-177A and -177E, 1/15/2022. Expanded and merged in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.168
VAP-E25E_ROS01C20 / Robinson Creek / Portion of VDH shellfish condemnation 177, 5/28/1997 open in 028-177, 1/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.039
VAP-E25E_ROS02A04 / Robinson Creek, UT and Perkins Creek / Described in VDH Shellfish Condemnation 028-177C, 1/15/2022 and the portion of 028-177B, 1/15/2022 that is within Perkins Creek. RPPMH	4A	Dissolved Oxygen	2020	NA	0.039
VAP-E25E_ROS02B12 / Robinson Creek / Described in VDH Shellfish Condemnation 028-177M1, 1/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.007
VAP-E25E_ROS02C16 / Robinson Creek, UT / Described in VDH Shellfish Condemnation 028-177D, 1/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.013
VAP-E25E_ROS02D24 / Robinson Creek / Portion of VDH-DSS Condemnation 028-177B, 1/15/2022 within Robinson Creek. RPPMH	4A	Dissolved Oxygen	2020	NA	0.039
VAP-E25E_RPP01C10 / Rappahannock River: Mark Haven Beach Basin / The portion of VDH shellfish condemnation 026-181B, 1/20/2006 not administratively closed. RPPMH	4A	Dissolved Oxygen	1998	L	0.010
VAP-E25E_RPP01C98 / Mark Haven Beach Basin / As delineated in VDH shellfish condemnation 026-181A, 4/3/2012. RPPMH	4A	Dissolved Oxygen	2020	NA	0.004
VAP-E25E_RPP02A02 / Rappahannock River / The mainstem of the Rappahannock River from the end of MSN (Sharps/0.7 mi DS of Mark Haven Beach to the mouth, excluding area in SFC 051A. RPPMH	4A	Dissolved Oxygen	1998	L	81.443
VAP-E25E_RPP03A06 / Rappahannock River / Described in VDH Shellfish Condemnation 024-070B, 11/15/2022. RPPMH	4D	Dissolved Oxygen	1998	L	0.008
VAP-E25E_RPP03B16 / Rappahannock River Run Bluffs / As described in VDH shellfish condemnation 026-181B, 3/15/2021. RPPMH	4A	Dissolved Oxygen	1998	L	0.003

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_URB01A00 / Urbanna Creek / As described in VDH-DSS condemnation 029-042B, 2/14/2006. RPPMH	4A	Dissolved Oxygen	2020	NA	0.215
VAP-E25E_URB02A00 / Urbanna Creek / As delineated in VDH shellfish condemnation 029-042A, 2/14/2006. RPPMH	4A	Dissolved Oxygen	2020	NA	0.238
VAP-E25E_WEE01A00 / Weeks Creek / Delineated in VDH shellfish condemnation 202, 10/8/1996. RPPMH	4A	Dissolved Oxygen	2020	NA	0.123
VAP-E25E_WEE02A04 / Weeks Creek / The portion of VDH shellfish condemnation 027-202A, 1/27/2015 not included in the 1989 closure. RPPMH	4A	Dissolved Oxygen	2020	NA	0.013
VAP-E25E_XDV01A02 / XDV - Rappahannock River, UT (aka Beach Creek) / The segment boundaries are delineated in VDH shellfish condemnation 022-116S107, 11/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.084
VAP-E25E_ZZZ01C14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA68. RPPMH	4A	Dissolved Oxygen	2020	NA	0.219
VAP-E25E_ZZZ01D14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA69. RPPMH	4A	Dissolved Oxygen	2020	NA	0.239
VAP-E26E_BPC01A98 / Bush Park Creek / As delineated in VDH shellfish condemnation 109, 4/27/1989. RPPMH	4A	Dissolved Oxygen	2020	NA	0.103
VAP-E26E_BRD01A00 / Broad Creek / As delineated in VDH shellfish condemnation 033-038B, 11/21/2013. RPPMH	4A	Dissolved Oxygen	2020	NA	0.084
VAP-E26E_BRD02A00 / Broad Creek / As delineated in VDH shellfish condemnation 033-038A, 11/21/2013. RPPMH	4A	Dissolved Oxygen	2020	NA	0.040
VAP-E26E_BRD04A00 / Broad Creek / Described in VDH-DSS condemnation 033-038M1, 11/21/2013. RPPMH	4A	Dissolved Oxygen	2020	NA	0.037
VAP-E26E_BRI01A18 / Bridge Cove / Described in VDH-DSS condemnation 020-041D, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.005
VAP-E26E_BRI01B22 / Bridge Cove / Described in VDH-DSS Condemnation 020-041S50, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.035

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CAR01A02 / Carter Cove / Described in VDH-DSS Condemnation 020-041E, 11/15/2020. RPPMH	4A	Dissolved Oxygen	2020	NA	0.040
VAP-E26E_CAR01B14 / Carter Cove / Portion of VDH-DSS SFC 020-041A, 11/1/1996 included in 020-041M1, 11/15/2020. RPPMH	4A	Dissolved Oxygen	2020	NA	0.016
VAP-E26E_CEB01A00 / Eastern Branch Carter Creek / Described in VDH shellfish condemnation 041C, 11/1/1996. RPPMH	4A	Dissolved Oxygen	2020	NA	0.084
VAP-E26E_CEB01B08 / Eastern Branch Carter Creek / Portion of VDH shellfish condemnation 020-041A, 10/15/2022 not included in 041C, 11/1/1996. Shrank and split in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.072
VAP-E26E_CEB01C22 / Eastern Branch Carter Creek / Described in VDH-DSS condemnation 020-041C, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.012
VAP-E26E_CRR02A08 / Corrotoman River / The portion of the Corrotoman River that is within CB segment RPPMH.	4A	Dissolved Oxygen	1998	L	1.039
VAP-E26E_CTR01A00 / Carter Creek / As delineated in VDH shellfish condemnation 020-041E, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.204
VAP-E26E_CTR02A00 / Carter Creek / Delineated in VDH shellfish condemnation 020-041B, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.058
VAP-E26E_CTR03A00 / Carter Creek / Portion of VDH-DSS SFC 020-041M1, $10/15/2022$ not included in 020-041A, $11/1/1996$. RPPMH	4A	Dissolved Oxygen	2020	NA	0.114
VAP-E26E_CTR03B16 / Carter Creek / Portion of Carter Creek open in VDH-DSS Condemnation 020-041, 10/15/2022. Expanded slightly in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.268
VAP-E26E_HNU01A08 / Hunting Creek / Described in VDH-DSS Condemnation 032-104B, 9/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.021
VAP-E26E_HNU02A20 / Hunting Creek / Described in VDH-DSS condemnation 032-104S36, 9/15/2019. RPPMH	4A	Dissolved Oxygen	2020	NA	0.017

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_JAC01A24 / Jacks Cove / Described in VDH shellfish condemnation 020-041S49, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.016
VAP-E26E_LOL01A02 / Locklies Creek / Delineated in VDH shellfish condemnation 102, 10/31/1994. Merged in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.101
VAP-E26E_LOL02A06 / Locklies Creek / Described in VDH-DSS Shellfish Condemnation 031-102M1, 9/15/2021. RPPMH	4A	Dissolved Oxygen	2020	NA	0.054
VAP-E26E_LOL03A08 / Roane Cove of Locklies Creek / Described in VDH-DSS Shellfish Condemnation 031-102C, 9/15/2021. RPPMH	4A	Dissolved Oxygen	2020	NA	0.034
VAP-E26E_MEA01A00 / Meachim Creek / Described in VDH shellfish condemnation 030-179A, 9/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.077
VAP-E26E_MEA01B00 / Meachim Creek / Described in VDH shellfish condemnation 179B, 12/9/1996. RPPMH	4A	Dissolved Oxygen	2020	NA	0.032
VAP-E26E_MEA01C06 / Meachim Creek / Described in VDH Shellfish Condemnation 030-179M1, 9/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.034
VAP-E26E_MEA01D24 / Meachim Creek, UT / Portions of VDH shellfish condemnation 030179C, 9/15/2022 within 179A, 12/9/1996. RPPMH	4A	Dissolved Oxygen	2020	NA	0.013
VAP-E26E_MEA02A00 / Meachim Creek / Downstream of VDH Shellfish Condemnation 030-179, 12/9/1996 not otherwise segmented. Split in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.108
VAP-E26E_MEA02C24 / Meachim Creek, UT / Portion of VDH Shellfish Condemnation 030-179C, 9/15/2022 not closed in 030-179A, 12/9/1996. RPPMH	4A	Dissolved Oxygen	2020	NA	0.024
VAP-E26E_MEA03A10 / Meachim Creek / Portions of VDH shellfish condemnation 179A, 12/9/1996 open on 030-179, 9/15/2022. Split in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.040

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MLL01A98 / Mill Creek / Described in VDH shellfish condemnation 031-102A, 9/15/2021. Expanded slightly in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.111
VAP-E26E_MLL01B12 / Mill Creek / Portion of VDH shellfish condemnation 103, 12/10/1991 open in 031-102, 9/15/2021. Shrank in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.013
VAP-E26E_MLL02A06 / Mill Creek / Downstream of VDH shellfish condemnation 103, 12/10/1991 RPPMH	4A	Dissolved Oxygen	2020	NA	0.358
VAP-E26E_MOS01C20 / Mosquito Creek / As delineated in VDH shellfish condemnation 018-203, 1/6/2005. RPPMH	4A	Dissolved Oxygen	2020	NA	0.069
VAP-E26E_OYS01B24 / Oyster Creek / Portion of VDH condemnation 018-053A, 1/4/2005 within RPPMH. RPPMH	4A	Dissolved Oxygen	2020	NA	0.071
VAP-E26E_RPP04A00 / Rappahannock River / Described in VDH Shellfish Condemnation 029-051B, 9/1/2015. RPPMH	4A	Dissolved Oxygen	1998	L	0.131
VAP-E26E_RPP05A00 / Rappahannock River / Delineated in VDH-DSS condemnation 029-051C, 9/1/2015. RPPMH	4A	Dissolved Oxygen	1998	L	0.049
VAP-E26E_RPP07A02 / Rappahannock River / As delineated in VDH-DSS condemnation 018-053A, 7/23/2018. RPPMH	4D	Dissolved Oxygen	1998	L	0.139
VAP-E26E_RPP08A22 / Rappahannock River / Portion of VDH-DSS Condemnation 032-109A, 9/15/2022 that is within the mainstem Rappahannock River. Shrank in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	1998	L	0.002
VAP-E26E_STE01A98 / Sturgeon Creek / As delineated in VDH shellfish condemnation 104, 11/28/1994. Merged in the 2024 cycle. RPPMH	4A	Dissolved Oxygen	2020	NA	0.085
VAP-E26E_STE02A08 / Sturgeon Creek / Sturgeon Creek downstream of condemnation 104, 11/28/1994. RPPMH	4A	Dissolved Oxygen	2020	NA	0.192

(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHS01B00 / Whiting Creek / As delineated in VDH shellfish condemnation 029-051A, 9/1/2015. RPPMH	4A	Dissolved Oxygen	2020	NA	0.196
VAP-E26E_WOO01A08 / Woods Creek / Tidal Woods Creek RPPMH	4A	Dissolved Oxygen	2020	NA	0.037
VAP-E26E_XEV01A02 / Windmill Point Yacht Harbor / As delineated in VDH-DSS Shellfish Condemnation 018-053C, 7/23/2018. RPPMH	4A	Dissolved Oxygen	2020	NA	0.015
VAP-E26E_XII01A18 / XII - Windmill Point Creek, UT (aka White Marsh) / Described in VDH-DSS condemnation 018-053B, 7/23/2018 RPPMH	4A	Dissolved Oxygen	2020	NA	0.034
VAP-E26E_YOP01A18 / Yopps Cove / Described in VDH-DSS condemnation 020-041S51, 10/15/2022. RPPMH	4A	Dissolved Oxygen	2020	NA	0.022
VAP-E26E_ZZZ01D14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA73 RPPMH	4A	Dissolved Oxygen	2020	NA	0.028
VAP-E26E_ZZZ01E14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA74 RPPMH	4A	Dissolved Oxygen	2020	NA	0.613

Rappahannock River

Aquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 103.285	Reservoir (Acres)	River (Miles)
Rappahannock Rive	r			
Deep-Channel Se	easonal Refuge Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 97.468	Reservoir (Acres)	River (Miles)
Rappahannock Rive	r			
Deep-Water Aqu	atic Life Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 102.41	Reservoir (Acres)	River (Miles)
Rappahannock Rive	r			
Migratory Fish S	Description Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 11.81	Reservoir (Acres)	River (Miles)

Rappahannock River

	Estuary	Reservoir	River
Open-Water Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Dissolved Oxygen - Total Impaired Size by Water Type:	103.285	`	

Sources: Agriculture; Atmospheric Deposition; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Rappahannock River Basin

Cause Group Code: RPPMH-SAV-BAY Rappahannock River

Cause Location: The Rappahannock River mesohaline estuary (RPPMH)

Cause City/County: Essex County; Lancaster County; Middlesex County; Richmond County

Use(s): Aquatic Life; Shallow-Water Submerged Aquatic Vegetation

Causes(s)/VA Category: Aquatic Plants (Macrophytes)/4A

Cause Description: In the 2024 cycle, the mesohaline portion of the mainstem Rappahannock does not meet the Chesapeake Bay Shallow Water Subuse's submerged aquatic vegetation acreage criterion. There was insufficient information to assess the water clarity criteria.

The Chesapeake Bay TMDL was approved by the EPA on 12/29/2010; therefore, it is considered Category 4A

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E22E_RPP05A02 / Rappahannock River / The oligohaline/mesohaline boundary at river mile 48.51 to the downstream boundary of VDH shellfish condemnation area 025A-068B, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	6.958
VAP-E22E_WAR01A18 / Waterview Creek / Tidal portion of Waterview Creek RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.038
VAP-E22E_ZZZ02A06 / Unsegmented estuaries in E22 / Unsegmented portion of watershed. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.014
VAP-E23E_CAT01A02 / Cat Point Creek / The tidal portion of Cat Point Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	1.280
VAP-E23E_CRC01A08 / Church Swamp / Tidal limit to mouth at Hoskins Creek RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.002
VAP-E23E_HOK01A98 / Hoskins Creek / Hoskins Creek from the Tappahannock STP downstream to the mouth at the Rappahannock River. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.084
VAP-E23E_HOK02A08 / Hoskins Creek / Hoskins Creek from its tidal limit to the confluence with Church Swamp. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.052
VAP-E23E_HOK02A10 / Hoskins Creek / Hoskins Creek from the confluence with Church Swamp downstream to the Tappahannock STP. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.016
VAP-E23E_LIE01A98 / Little Carter Creek, Jugs Creek / Tidal limit to mouth at the Rappahannock River. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.419
VAP-E23E_MTL01A10 / Mount Landing Creek / Tidal limit to mouth at the Rappahannock River. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.172

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E23E_PIS02A00 / Piscataway Creek / The estuarine portion of Piscataway Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.589
VAP-E23E_RPP02A98 / Rappahannock River / Mainstem Rappahannock as described in VDH shellfish condemnation 025A-068A, 3/15/2021, excluding administratively condemned portion. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	8.123
VAP-E23E_RPP02B10 / Rappahannock River / Portion of mainstem Rappahannock River that is administratively condemned within condemnation 025A-068A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.158
VAP-E23E_RPP02C12 / Rappahannock River / Portion of VDH shellfish condemnation 025A-068A, 11/14/2005 not included in 025A-068A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.387
VAP-E23E_ZZZ02A06 / Unsegmented estuaries in E23 / Unsegmented portion within VDH-DSS Condemnation 025A-068A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.049
VAP-E23E_ZZZ02B10 / Unsegmented estuaries in E23 / Administrative portion within SFC 025A-068A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.007
VAP-E24E_LIK01A12 / Little Totuskey Creek / Tidal limit to mouth at Totuskey Creek RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.055
VAP-E24E_RIC01A04 / Richardson Creek / Richardson Creek within VDH-DSS Condemnation 025-071A, 3/15/2021 (non-administrative.) RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.321
VAP-E24E_RIC01C10 / Richardson Creek / Portion of Richardson Creek within the administratively condemned portion of VDH Condemnation 025-071A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.024
VAP-E24E_RPP01B14 / Garrett's Marina / As delineated in VDH shellfish condemnation 026-181A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.003
VAP-E24E_RPP01B98 / Rappahannock River: Garrett's Marina / As delineated in VDH shellfish condemnation 026-181M1, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.025

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E24E_RPP01C06 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/16/2007 (non-admin) that is open 3/15/2021. Shrank in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.327
VAP-E24E_RPP01D10 / Rappahannock River / The portion of the Rappahannock River within the administratively closed area of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.137
VAP-E24E_RPP01E18 / Rappahannock River / The Rappahannock River mainstem within VDH shellfish condemnation 025-071A, 3/15/2021 (non-admin) Expanded in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.378
VAP-E24E_RPP03A00 / Rappahannock River / The Rappahannock River from the limit of VDH shellfish condemnation 068A, 11/14/2005 downstream to end of MSN (Sharps/0.7 mi DS of Mark Haven Beach) unless otherwise segmented RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	10.858
VAP-E24E_RPP03D24 / Rappahannock River / Described in VDH shellfish condemnation 026-270A, 5/25/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.061
VAP-E24E_TOT01A00 / Totuskey Creek / The segment boundary is delineated in VDH condemnation 025-071B, 3/15/2021 excluding Little Totuskey Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.302
VAP-E24E_TOT02A00 / Totuskey Creek / Portion of VDH shellfish condemnation 025-071A, 4/15/2020 within Totuskey Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.647
VAP-E24E_TOT02B10 / Totuskey Creek / Portion of Totuskey Creek within the non-administrative portion of VDH shellfish condemnation 025-071A, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.064
VAP-E25E_DEE01A04 / Deep Creek / Described in VDH shellfish condemnation 121, 11/16/1994. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.049
VAP-E25E_DEE01B08 / Deep Creek / Portions of VDH-DSS condemnations 023-121B, 12/15/2021 not included in the 11/16/1994 condemnation. Merged and adjusted in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.182

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_FAM01A98 / Farnham Creek / Portion of VDH shellfish condemnation 070, 10/22/1996 not administratively condemned in 024-070A, 11/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.350
VAP-E25E_FAM01B22 / Farnham Creek / Portion of VDH shellfish condemnation 070, 10/22/1996 administratively condemned in 024-070A, 11/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.074
VAP-E25E_GEE01A98 / Greenvale Creek / As delineated in VDH shellfish condemnation 094, 11/7/1994. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.087
VAP-E25E_GEE02A06 / Greenvale Creek / Portion of VDH-DSS condemnation 022-094A, 11/15/2022 not included in 94, 11/7/1994. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.012
VAP-E25E_GEE02B10 / Belmont Creek / Described in VDH-DSS Shellfish Condemnation 022-094S108, 11/15/2022. Shrank in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.024
VAP-E25E_HRY01A06 / Harry George Creek / Tidal limit to mouth at the Rappahannock River RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.095
VAP-E25E_LAN01A98 / Lancaster Creek / As delineated in VDH-DSS Shellfish Condemnation 023-120A, 8/14/1995. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.270
VAP-E25E_LAN01B08 / Lancaster Creek / The portion of VDH Shellfish Condemnation 023-120A, 9/22/2022 not included in 120A, 8/14/1995. Merged and expanded in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.489
VAP-E25E_LAN02A02 / Lancaster Creek / Lancaster Creek downstream of VDH Shellfish Condemnation 023-120, 9/22/2022, not otherwise segmented. Size reduced in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.823
VAP-E25E_LAN03A06 / Lancaster Creek / Described in VDH Shellfish Condemnation 023-120C, 9/22/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.023
VAP-E25E_LGG01A98 / Lagrange Creek / As described in VDH Shellfish Condemnation 028-127A, 1/15/2022. Expanded in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.562

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_LGG01B18 / Lagrange Creek / Portion of VDH Shellfish Condemnation 127, 6/11/1996 open on 028-127, 1/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.037
VAP-E25E_LGG02A06 / Lagrange Creek / Lagrange Creek downstream of VDH Shellfish Condemnation 127, 6/11/1996 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.040
VAP-E25E_MTT01A00 / Morattico Creek / Delineated in VDH Shellfish Condemnation 023-120B, 9/22/2022. Expanded in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.345
VAP-E25E_MUB01A02 / Mulberry Creek / Described in VDH shellfish condemnation 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.148
VAP-E25E_MUB02A06 / Mulberry Creek / Portion of VDH shellfish condemnation 023-121A, 12/15/2021 not included in 120B, 8/14/1995. Merged in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.100
VAP-E25E_MUC01A04 / Mud Creek / Described in VDH-DSS Condemnation 027-090B, 8/26/2008. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.204
VAP-E25E_PAY01A02 / Paynes Creek / As delineated in VDH-DSS Shellfish Condemnation 022-094S211, 11/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.049
VAP-E25E_PRR01A02 / Parrotts Creek / Delineated in VDH shellfish condemnation 090, 4/27/1989. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.153
VAP-E25E_PRR02A08 / Parrotts Creek / Portion of VDH-DSS Condemnation 027-090A, 1/15/2022 downstream of condemnation 090, 4/27/1989. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.011
VAP-E25E_ROS01A00 / Robinson Creek / Described in VDH shellfish condemnation 028-177A and -177E, 1/15/2022. Expanded and merged in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.168
VAP-E25E_ROS01C20 / Robinson Creek / Portion of VDH shellfish condemnation 177, 5/28/1997 open in 028-177, 1/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.039

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_ROS02A04 / Robinson Creek, UT and Perkins Creek / Described in VDH Shellfish Condemnation 028-177C, 1/15/2022 and the portion of 028-177B, 1/15/2022 that is within Perkins Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.039
VAP-E25E_ROS02B12 / Robinson Creek / Described in VDH Shellfish Condemnation 028-177M1, 1/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.007
VAP-E25E_ROS02C16 / Robinson Creek, UT / Described in VDH Shellfish Condemnation 028-177D, 1/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.013
VAP-E25E_ROS02D24 / Robinson Creek / Portion of VDH-DSS Condemnation 028-177B, 1/15/2022 within Robinson Creek. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.039
VAP-E25E_RPP01C10 / Rappahannock River: Mark Haven Beach Basin / The portion of VDH shellfish condemnation 026-181B, 1/20/2006 not administratively closed. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.010
VAP-E25E_RPP01C98 / Mark Haven Beach Basin / As delineated in VDH shellfish condemnation 026-181A, 4/3/2012. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.004
VAP-E25E_RPP02A02 / Rappahannock River / The mainstem of the Rappahannock River from the end of MSN (Sharps/0.7 mi DS of Mark Haven Beach to the mouth, excluding area in SFC 051A. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	81.443
VAP-E25E_RPP03A06 / Rappahannock River / Described in VDH Shellfish Condemnation 024-070B, 11/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.008
VAP-E25E_RPP03B16 / Rappahannock River Run Bluffs / As described in VDH shellfish condemnation 026-181B, 3/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.003
VAP-E25E_TWN01A12 / Town Bridge Swamp / Tidal limit to mouth at Urbanna Creek RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.002
VAP-E25E_URB01A00 / Urbanna Creek / As described in VDH-DSS condemnation 029-042B, 2/14/2006. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.215
VAP-E25E_URB02A00 / Urbanna Creek / As delineated in VDH shellfish condemnation 029-042A, 2/14/2006. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.238

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E25E_WEE01A00 / Weeks Creek / Delineated in VDH shellfish condemnation 202, 10/8/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.123
VAP-E25E_WEE02A04 / Weeks Creek / The portion of VDH shellfish condemnation 027-202A, 1/27/2015 not included in the 1989 closure. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.013
VAP-E25E_XDV01A02 / XDV - Rappahannock River, UT (aka Beach Creek) / The segment boundaries are delineated in VDH shellfish condemnation 022-116S107, 11/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.084
VAP-E25E_ZZZ01A14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA65 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.077
VAP-E25E_ZZZ01C14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA68. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.219
VAP-E25E_ZZZ01D14 / Unsegmented estuaries in E25 / Unsegmented portion of watershed RA69. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.239
VAP-E26E_BPC01A98 / Bush Park Creek / As delineated in VDH shellfish condemnation 109, 4/27/1989. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.103
VAP-E26E_BRD01A00 / Broad Creek / As delineated in VDH shellfish condemnation 033-038B, 11/21/2013. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.084
VAP-E26E_BRD02A00 / Broad Creek / As delineated in VDH shellfish condemnation 033-038A, 11/21/2013. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.040
VAP-E26E_BRD04A00 / Broad Creek / Described in VDH-DSS condemnation 033-038M1, 11/21/2013. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.037
VAP-E26E_BRI01A18 / Bridge Cove / Described in VDH-DSS condemnation 020-041D, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.005
VAP-E26E_BRI01B22 / Bridge Cove / Described in VDH-DSS Condemnation 020-041S50, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.035
VAP-E26E_CAR01A02 / Carter Cove / Described in VDH-DSS Condemnation 020-041E, 11/15/2020. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.040
VAP-E26E_CAR01B14 / Carter Cove / Portion of VDH-DSS SFC 020-041A, 11/1/1996 included in 020-041M1, 11/15/2020. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.016

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_CEB01A00 / Eastern Branch Carter Creek / Described in VDH shellfish condemnation 041C, 11/1/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.084
VAP-E26E_CEB01B08 / Eastern Branch Carter Creek / Portion of VDH shellfish condemnation 020-041A, 10/15/2022 not included in 041C, 11/1/1996. Shrank and split in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.072
VAP-E26E_CEB01C22 / Eastern Branch Carter Creek / Described in VDH-DSS condemnation 020-041C, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.012
VAP-E26E_CRR02A08 / Corrotoman River / The portion of the Corrotoman River that is within CB segment RPPMH.	4A	Aquatic Plants (Macrophytes)	2024	L	1.039
VAP-E26E_CTR01A00 / Carter Creek / As delineated in VDH shellfish condemnation 020-041E, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.204
VAP-E26E_CTR02A00 / Carter Creek / Delineated in VDH shellfish condemnation 020-041B, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.058
VAP-E26E_CTR03A00 / Carter Creek / Portion of VDH-DSS SFC 020-041M1, 10/15/2022 not included in 020-041A, 11/1/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.114
VAP-E26E_CTR03B16 / Carter Creek / Portion of Carter Creek open in VDH-DSS Condemnation 020-041, 10/15/2022. Expanded slightly in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.268
VAP-E26E_HNU01A08 / Hunting Creek / Described in VDH-DSS Condemnation 032-104B, 9/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.021
VAP-E26E_HNU02A20 / Hunting Creek / Described in VDH-DSS condemnation 032-104S36, 9/15/2019. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.017
VAP-E26E_JAC01A24 / Jacks Cove / Described in VDH shellfish condemnation 020-041S49, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.016

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_LOL01A02 / Locklies Creek / Delineated in VDH shellfish condemnation 102, 10/31/1994. Merged in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.101
VAP-E26E_LOL02A06 / Locklies Creek / Described in VDH-DSS Shellfish Condemnation 031-102M1, 9/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.054
VAP-E26E_LOL03A08 / Roane Cove of Locklies Creek / Described in VDH-DSS Shellfish Condemnation 031-102C, 9/15/2021. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.034
VAP-E26E_MEA01A00 / Meachim Creek / Described in VDH shellfish condemnation 030-179A, 9/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.077
VAP-E26E_MEA01B00 / Meachim Creek / Described in VDH shellfish condemnation 179B, 12/9/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.032
VAP-E26E_MEA01C06 / Meachim Creek / Described in VDH Shellfish Condemnation 030-179M1, 9/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.034
VAP-E26E_MEA01D24 / Meachim Creek, UT / Portions of VDH shellfish condemnation 030179C, 9/15/2022 within 179A, 12/9/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.013
VAP-E26E_MEA02A00 / Meachim Creek / Downstream of VDH Shellfish Condemnation 030-179, 12/9/1996 not otherwise segmented. Split in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.108
VAP-E26E_MEA02B22 / Meachim Creek, UT / Portion of VDH Shellfish Condemnation 030-179B, 9/15/2022 downstream of 030-179, 12/9/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.004
VAP-E26E_MEA02C24 / Meachim Creek, UT / Portion of VDH Shellfish Condemnation 030-179C, 9/15/2022 not closed in 030-179A, 12/9/1996. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.024
VAP-E26E_MEA03A10 / Meachim Creek / Portions of VDH shellfish condemnation 179A, 12/9/1996 open on 030-179, 9/15/2022. Split in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.040

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Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_MLL01A98 / Mill Creek / Described in VDH shellfish condemnation 031-102A, 9/15/2021. Expanded slightly in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.111
VAP-E26E_MLL01B12 / Mill Creek / Portion of VDH shellfish condemnation 103, 12/10/1991 open in 031-102, 9/15/2021. Shrank in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.013
VAP-E26E_MLL02A06 / Mill Creek / Downstream of VDH shellfish condemnation 103, 12/10/1991 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.358
VAP-E26E_MOS01C20 / Mosquito Creek / As delineated in VDH shellfish condemnation 018-203, 1/6/2005. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.069
VAP-E26E_OYS01B24 / Oyster Creek / Portion of VDH condemnation 018-053A, 1/4/2005 within RPPMH. RPPMH	4A	Aquatic Plants (Macrophytes)	2006	L	0.071
VAP-E26E_RPP04A00 / Rappahannock River / Described in VDH Shellfish Condemnation 029-051B, 9/1/2015. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.131
VAP-E26E_RPP05A00 / Rappahannock River / Delineated in VDH-DSS condemnation 029-051C, 9/1/2015. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.049
VAP-E26E_RPP07A02 / Rappahannock River / As delineated in VDH-DSS condemnation 018-053A, 7/23/2018. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.139
VAP-E26E_RPP08A22 / Rappahannock River / Portion of VDH-DSS Condemnation 032-109A, 9/15/2022 that is within the mainstem Rappahannock River. Shrank in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.002
VAP-E26E_STE01A98 / Sturgeon Creek / As delineated in VDH shellfish condemnation 104, 11/28/1994. Merged in the 2024 cycle. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.085
VAP-E26E_STE02A08 / Sturgeon Creek / Sturgeon Creek downstream of condemnation 104, 11/28/1994. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.192

(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAP-E26E_WHS01B00 / Whiting Creek / As delineated in VDH shellfish condemnation 029-051A, 9/1/2015. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.196
VAP-E26E_WID01A12 / Windmill Point Creek / Described in VDH-DSS condemnation 018-053B, 11/2/2010. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.082
VAP-E26E_WOO01A08 / Woods Creek / Tidal Woods Creek RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.037
VAP-E26E_XEV01A02 / Windmill Point Yacht Harbor / As delineated in VDH-DSS Shellfish Condemnation 018-053C, 7/23/2018. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.015
VAP-E26E_XII01A18 / XII - Windmill Point Creek, UT (aka White Marsh) / Described in VDH-DSS condemnation 018-053B, 7/23/2018 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.034
VAP-E26E_YOP01A18 / Yopps Cove / Described in VDH-DSS condemnation 020-041S51, 10/15/2022. RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.022
VAP-E26E_ZZZ01D14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA73 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.028
VAP-E26E_ZZZ01E14 / Unsegmented estuaries in E26 / Unsegmented portion of watershed RA74 RPPMH	4A	Aquatic Plants (Macrophytes)	2024	L	0.613

Rappahannock River			
Aquatic Life Aquatic Plants (Macrophytes) - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 123.686	Reservoir (Acres)	River (Miles)
Rappahannock River			
	Estuary	Reservoir	River
Shallow-Water Submerged Aquatic Vegetation	(Sq. Miles)	(Acres)	(Miles)
Aquatic Plants (Macrophytes) - Total Impaired Size by Water Type:	123.686		

Sources: Agriculture; Atmospheric Deposition; Atmospheric Deposition - Nitrogen; Clean Sediments; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Natural Conditions - Water Quality Standards Use Attainability Analyses Needed; Sediment Resuspension (Clean Sediment); Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)

Rappahannock River Basin

Cause Group Code: RPPTF-DO-BAY Rappahannock Tidal Freshwater Estuary

Cause Location: The tidal freshwater Rappahannock River and its tributaries to the segment.

Cause City/County: Caroline County; Essex County; Fredericksburg; King George County; Spotsylvania County; Stafford County; Westmoreland County

Use(s): Aquatic Life; Migratory Fish Spawning and Nursery; Open-Water Aquatic Life

Causes(s)/VA Category: Dissolved Oxygen/4A

Cause Description: The Chesapeake Bay standards were implemented in the 2006 cycle.

During the 2018 - 2024 cycles, the tidal freshwater Rappahannock estuary (RPPTF) failed the Open Water Subuse 30-day mean summer dissolved oxygen criterion. The Open Water rest-of-year criterion was met. There was insufficient data to assess the other Open Water dissolved oxygen criteria and the Migratory Spawning and Nursery subuse.

The Chesapeake Bay TMDL was approved by the EPA on 12/19/2010; therefore, RPPTF is considered Category 4A.

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E20E_RPP01A02 / Rappahannock River / Segment begins at the confluence with Little Falls Run and continues downstream until the outlet of waterbody VAN-E20E. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.285
VAN-E20E_RPP02A02 / Rappahannock River / Segment begins at the confluence with Deep Run and continues downstream until the confluence with Little Falls Run. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.135
VAN-E20E_RPP03A02 / Rappahannock River / Segment begins at the fall line at Route 1 and continues downstream until the confluence with Deep Run. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.195
VAN-E21E_MIC01A06 / Mill Creek / Segment includes all tidal waters of Mill Creek. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.203
VAN-E21E_RPP01A02 / Rappahannock River / Segment begins at the confluence with Mill Creek, at rivermile 78.94, and continues downstream until immediately upstream of Devils Elbow, at rivermile 70.52. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	4.547
VAN-E21E_RPP03A02 / Rappahannock River / Segment begins at the confluence with Mount Creek and continues downstream until the confluence with Mill Creek. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	1.366
VAN-E21E_RPP04A02 / Rappahannock River / Segment begins at the confluence with Ware Creek and continues downstream until the confluence with Mount Creek. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	1.206

(continued)

Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAN-E21E_RPP05A02 / Rappahannock River / Segment begins at the confluence with Snow Creek (at the start of E21) and continues downstream until the confluence with Ware Creek. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.579
VAN-E21E_RPP48A02 / Rappahannock River-Muddy Creek / Segment includes all tidal waters in watershed RA48 not included in other delineated stream segments. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.006
VAN-E21E_RPP49A02 / Rappahannock River-Mount Creek / Segment includes all tidal waters in watershed RA49 not included in other delineated stream segments. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.147
VAN-E21E_RPP51A02 / Rappahannock River-Goldenvale Creek / Segment includes all tidal waters in watershed RA51 not included in other delineated stream segments. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.192
VAN-E21E_RPP52A02 / Rappahannock River-Portobago Creek / Segment includes all tidal waters in watershed RA52 not included in other delineated stream segments. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.079
VAN-E22E_ZZZ01A08 / Unnamed Rappahannock River Embayments / Segment includes all tidal waters in watershed not included in other delineated stream segments. Portion of CBP segment RPPTF.	4A	Dissolved Oxygen	2018	L	0.073
VAP-E22E_ELM01A10 / Elmwood Creek / Tidal limit to mouth at the Rappahannock River. RPPTF	4A	Dissolved Oxygen	2018	L	0.047
VAP-E22E_RPP01A02 / Rappahannock River / The Rappahannock River from Devils Elbow at Toby Point and Green Bay (river mile 70.52) downstream to the tidal freshwater/oligohaline boundary at river mile 57.85. RPPTF	4A	Dissolved Oxygen	2018	L	5.133
VAP-E22E_ZZZ01A00 / Unsegmented estuaries in E22 / Unsegmented portion of watershed within RPPTF.	4A	Dissolved Oxygen	2018	L	0.164

Rappahannock Tidal Freshwater Estuary

Aquatic Life	Dissolved Oxygen - Total Impaired Size by Water Type:	(Sq. Miles)	Reservoir (Acres)	
	Dissolved Oxygen - Total Impaired Size by water Type:	14.330		

Rappahannock Tidal Freshwater Estuary

Migratory Fish Spawning and Nursery Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 14.356	Reservoir (Acres)	River (Miles)
Rappahannock Tidal Freshwater Estuary		D .	D
Open-Water Aquatic Life Dissolved Oxygen - Total Impaired Size by Water Type:	Estuary (Sq. Miles) 14.356	Reservoir (Acres)	River (Miles)

Sources: Agriculture; Atmospheric Deposition - Nitrogen; Industrial Point Source Discharge; Internal Nutrient Recycling; Loss of Riparian Habitat; Municipal Point Source Discharges; Source Unknown; Sources Outside State Jurisdiction or Borders; Wet Weather Discharges (Point Source and Combination of Stormwater, SSO or CSO)