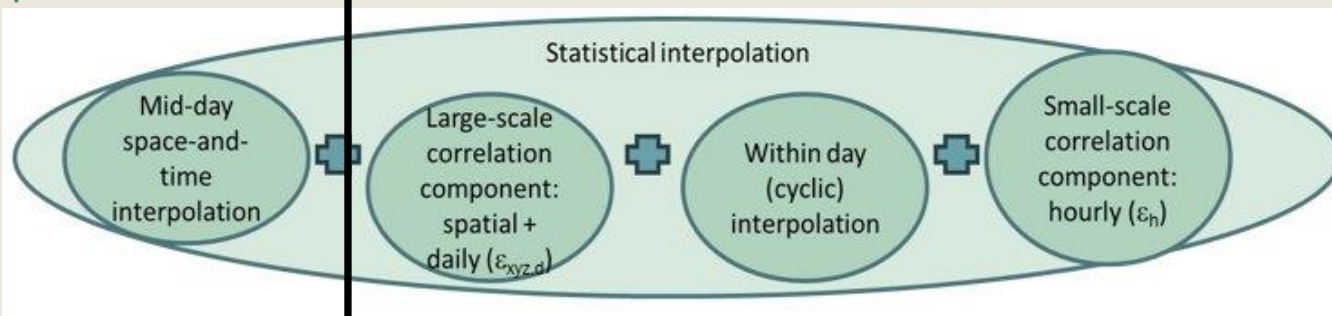


# Update on: Segment Interpolation Regions

Bay Oxygen Research Large Group  
Aug 18, 2025

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*Focus on this  
part of the tool*

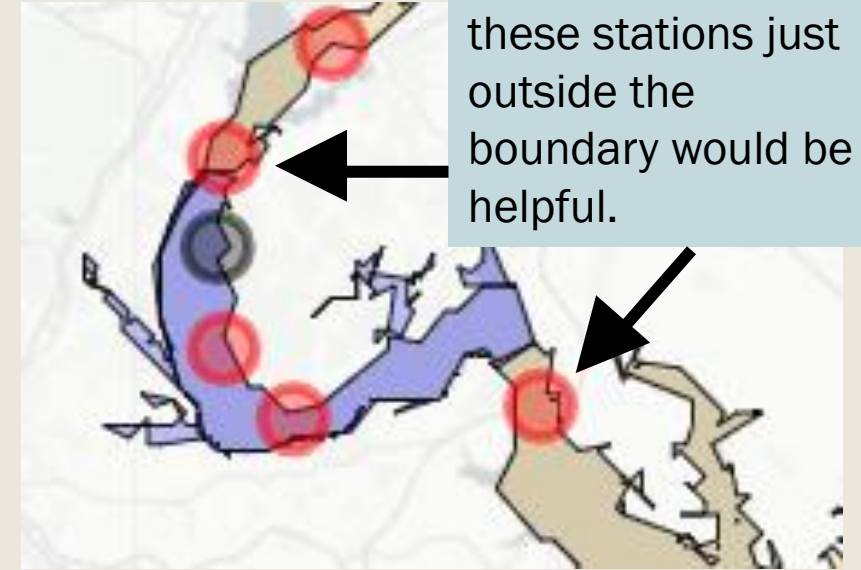


# Background

- Current 3-D interpolator uses boundary data outside the segment of interest for interpolations at the edges of segments.
- A similar need exists for the 4-D interpolation for good edge interpolations (to avoid extrapolation).

# Background

- Current 3-D interpolator uses boundary data outside the segment of interest for interpolations at the edges of segments.
- A similar need exists for the 4-D interpolation for good edge interpolations (to avoid extrapolation).
- In addition, the 4-D method requires sufficient data to build the smooth relationships between space, time, and DO. Data from one segment alone is not normally enough.



Example: If we want to interpolate in the purple segment, these stations just outside the boundary would be helpful.



For more details, see document [SegmentGrouping4D\\_8July2025](#) which includes maps for 31 currently selected regions and testing results.

# Examples:

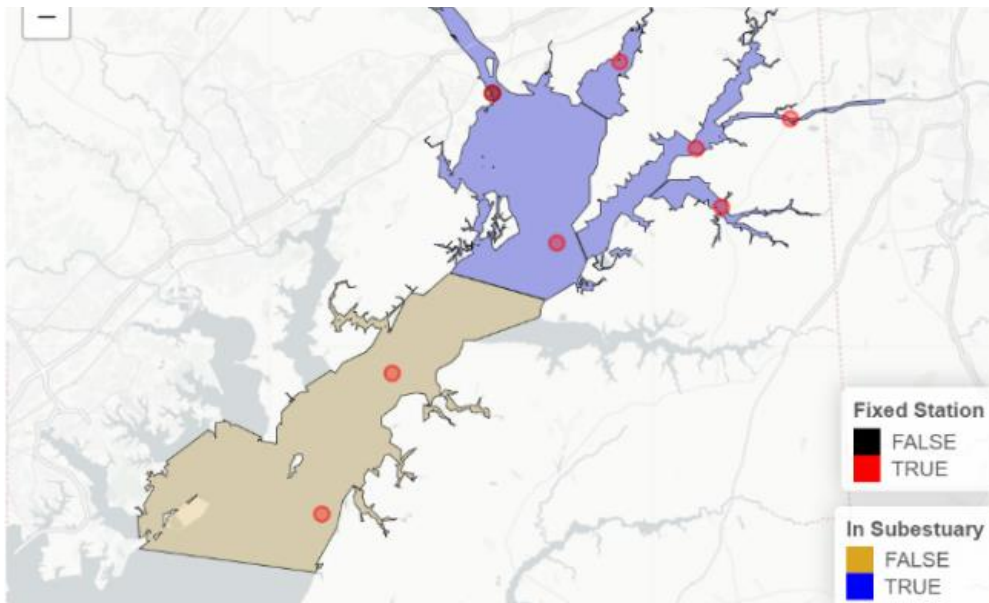
## Segment Interpolation Regions

- One or more segments are grouped together. Each region also includes all touching segments as boundary. Those boundary segments will also be in their own separate region(s).
- Including entire segments as boundary help “future proof” the approach as stations may be added or removed in coming years.

### CB1\_upE:

BOHOH, C&DOH\_DE, C&DOH\_MD, CB1TF, ELKOH, NORTF

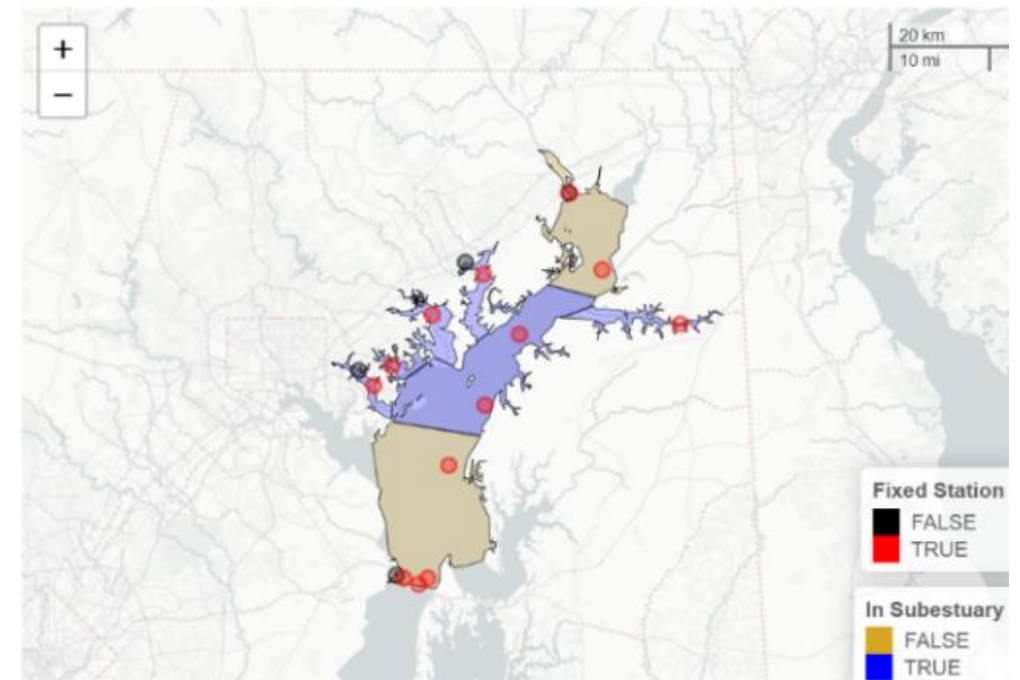
Boundary: CB20H



### CB2\_tribs:

BACOH, BSHOH, CB20H, GUNOH, MIDOH, SASOH

Boundary: CB1TF and CB3MH



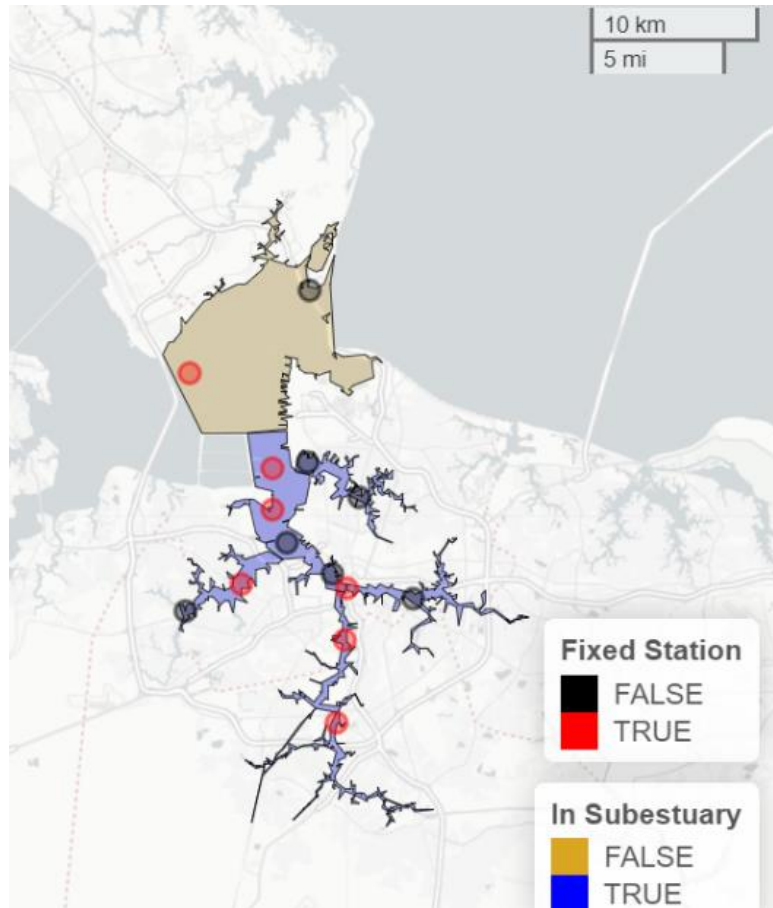
# Examples:

## Segment Interpolation Regions

### ELZ

ELIPH, LAFMH, EBEMH, SBEMH, WBEMH

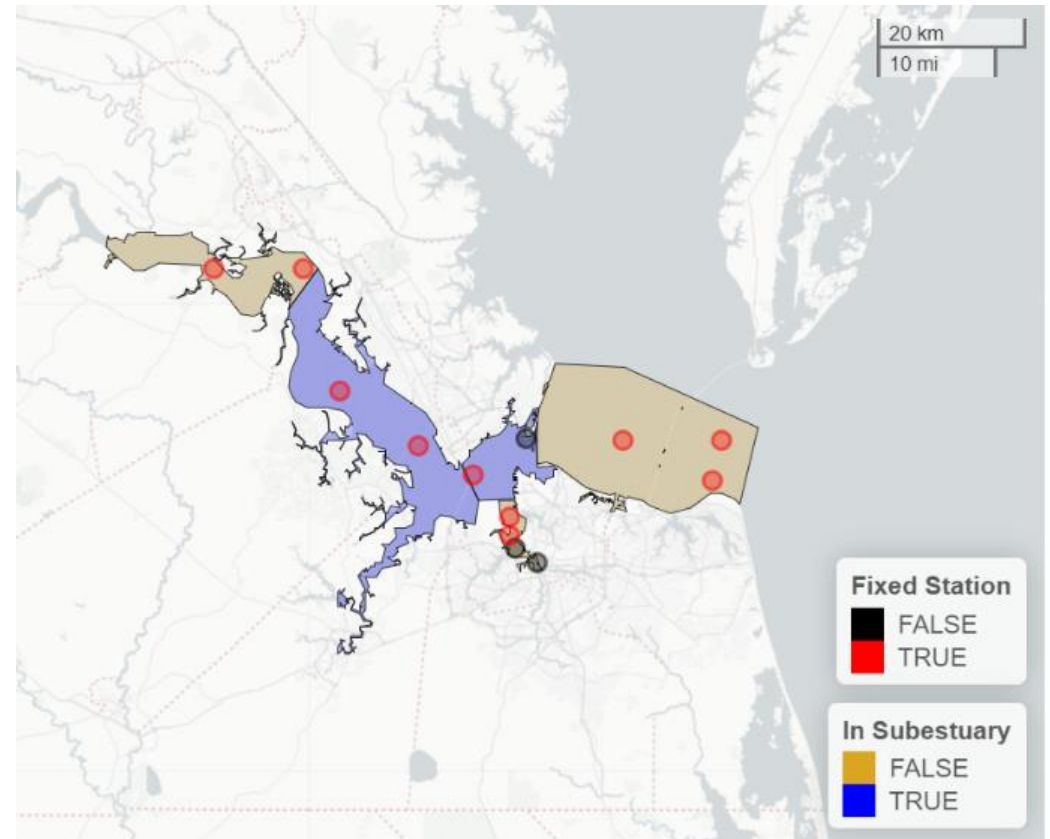
With boundary: JMSPH



### lowJMS

JMSPH, JMSMH

With boundary: CB8PH, ELIPH, JMSOH



# Details

- The regions are identified in an input spreadsheet to the 4-D tool, and adjustments are possible as we continue to test.
- Adjustments in the future will be possible too, especially if there are changes to the monitoring program.
- To get to this selected set of Segment Interpolation Regions, multiple options were evaluated.



	A	B	C	D	E	F	G	O
1	objectid	cbseg_92	cbpseg_st	state	cb_303d_segment	seg_int_reg	bndy_data	seg_int_reg_ID
2	11	CB2OH	CB2OH_MD	MD	Upper Chesapeake Bay	CB1_upE	TRUE	1
3	10	CB1TF	CB1TF_MD	MD	Northern Chesapeake Bay	CB1_upE	FALSE	1
4	31	ELKOH	ELKOH_MD	MD	Elk River	CB1_upE	FALSE	1
5	9	C&DOH_MD	C&DOH_MD	MD	C&D Canal, MD	CB1_upE	FALSE	1
6	8	C&DOH_DE	C&DOH_DE	DE	C&D Canal, DE	CB1_upE	FALSE	1
7	6	BOHOH	BOHOH_MD	MD	Bohemia River	CB1_upE	FALSE	1
8	52	NORTF	NORTF_MD	MD	Northeast River	CB1_upE	FALSE	1
9	10	CB1TF	CB1TF_MD	MD	Northern Chesapeake Bay	CB2_tribs	TRUE	2
10	12	CB3MH	CB3MH_MD	MD	Upper Central Chesapeake Bay	CB2_tribs	TRUE	2
11	75	SASOH	SASOH_MD	MD	Sassafras River	CB2_tribs	FALSE	2
12	11	CB2OH	CB2OH_MD	MD	Upper Chesapeake Bay	CB2_tribs	FALSE	2
13	4	BACOH	BACOH_MD	MD	Back River	CB2_tribs	FALSE	2
14	44	MIDOH	MIDOH_MD	MD	Middle River	CB2_tribs	FALSE	2
15	33	GUNOH	GUNOH_MD	MD	Gunpowder River	CB2_tribs	FALSE	2
16	7	BSHOH	BSHOH_MD	MD	Bush River	CB2_tribs	FALSE	2
17	11	CB2OH	CB2OH_MD	MD	Upper Chesapeake Bay	CB3	TRUE	3
18	13	CB4MH	CB4MH_MD	MD	Middle Central Chesapeake Bay	CB3	TRUE	3
19	24	CHSMH	CHSMH_MD	MD	Lower Chester River	CB3	TRUE	3
20	53	PATMH	PATMH_MD	MD	Patapsco River	CB3	TRUE	3
21	41	MAGMH	MAGMH_MD	MD	Magothy River	CB3	TRUE	3
22	12	CB3MH	CB3MH_MD	MD	Upper Central Chesapeake Bay	CB3	FALSE	3
23	53	PATMH	PATMH_MD	MD	Patapsco River	PAT_MAG	FALSE	4
24	41	MAGMH	MAGMH_MD	MD	Magothy River	PAT_MAG	FALSE	4
25	12	CB3MH	CB3MH_MD	MD	Upper Central Chesapeake Bay	PAT_MAG	TRUE	4



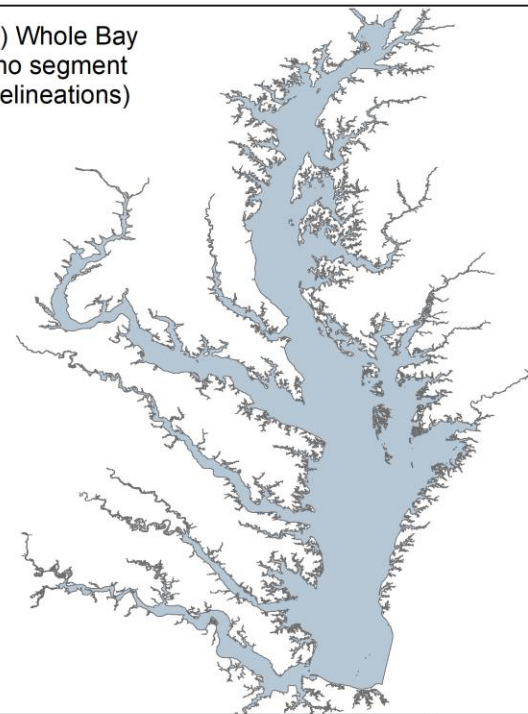
# Evaluation

## Some alternatives considered:

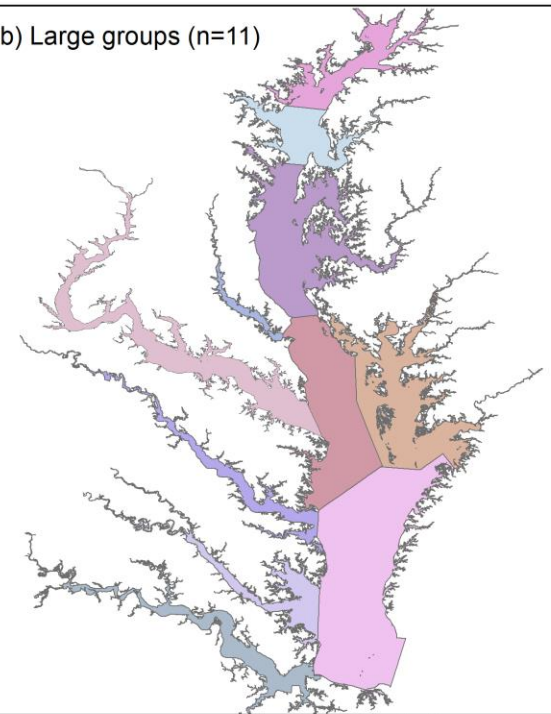
- A. Whole bay together, no segment delineations.
- B. 11 relatively large segment groups.
- C. 79 individual or small segment groups (*failures indicated with black*).
- D. 31 selected Segment Interpolation Regions–small, but balancing need to have enough data.

*Note: in all options except A, adjacent segments are used as boundary.*

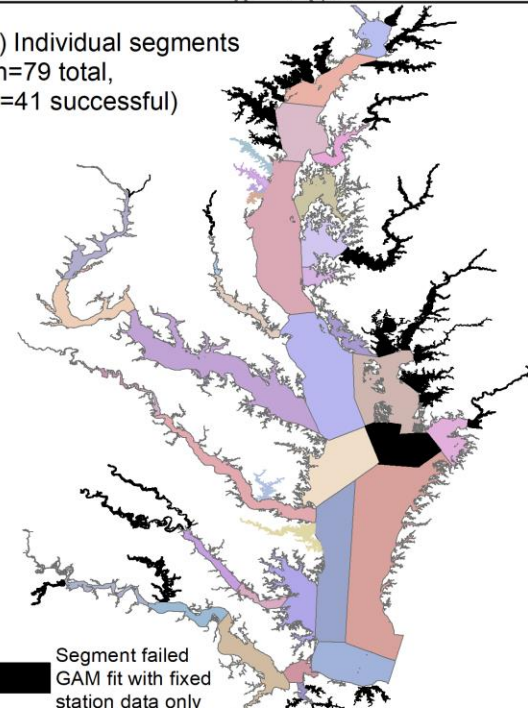
a) Whole Bay  
(no segment  
delineations)



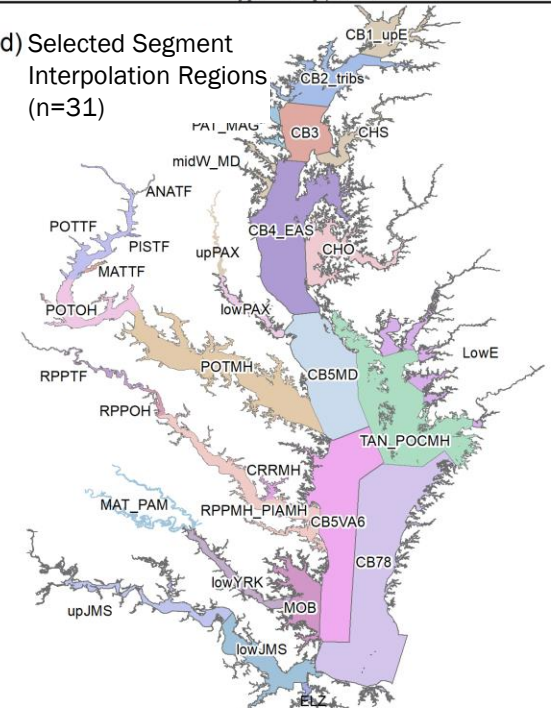
b) Large groups (n=11)



c) Individual segments  
(n=79 total,  
n=41 successful)

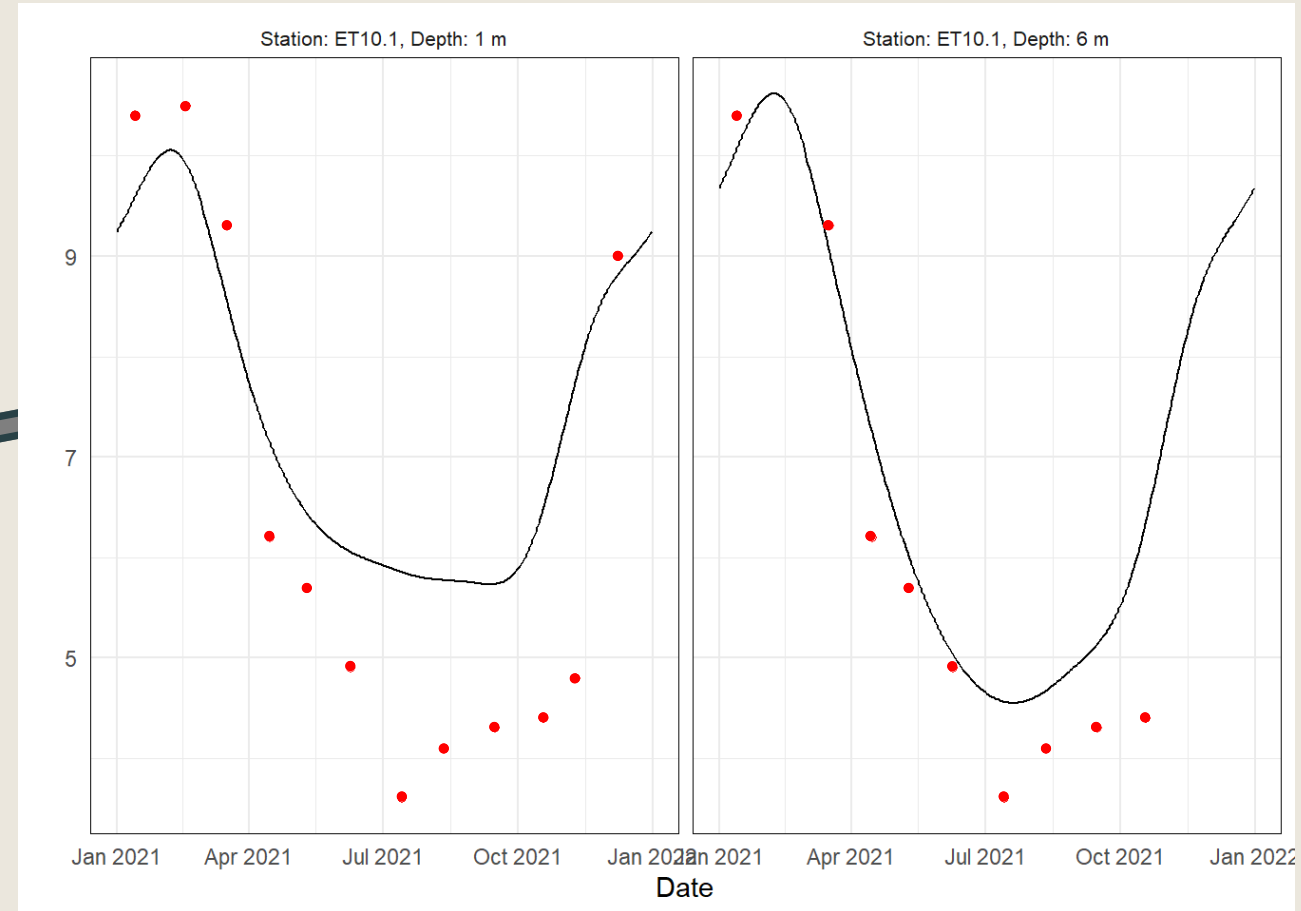
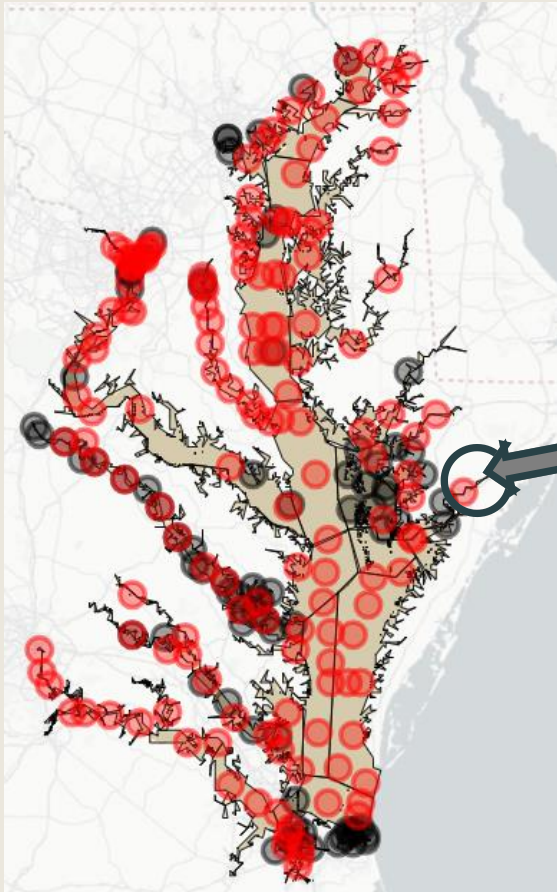


d) Selected Segment  
Interpolation Regions  
(n=31)



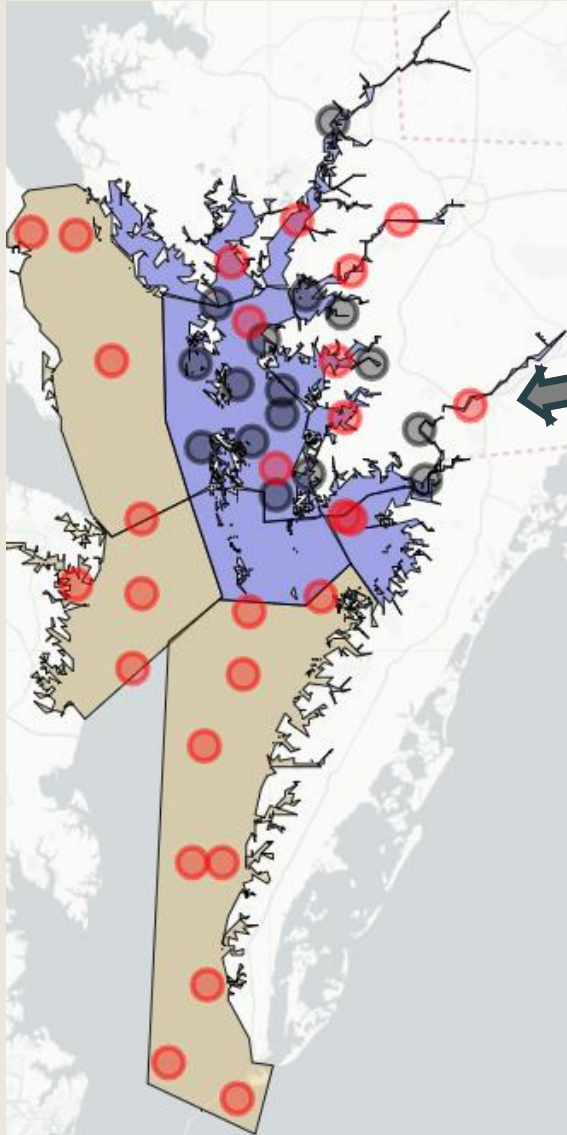
# Example: impact at one station of region size

A. Whole bay together: results in estimates at ET10.1 clearly mis-matched to the data

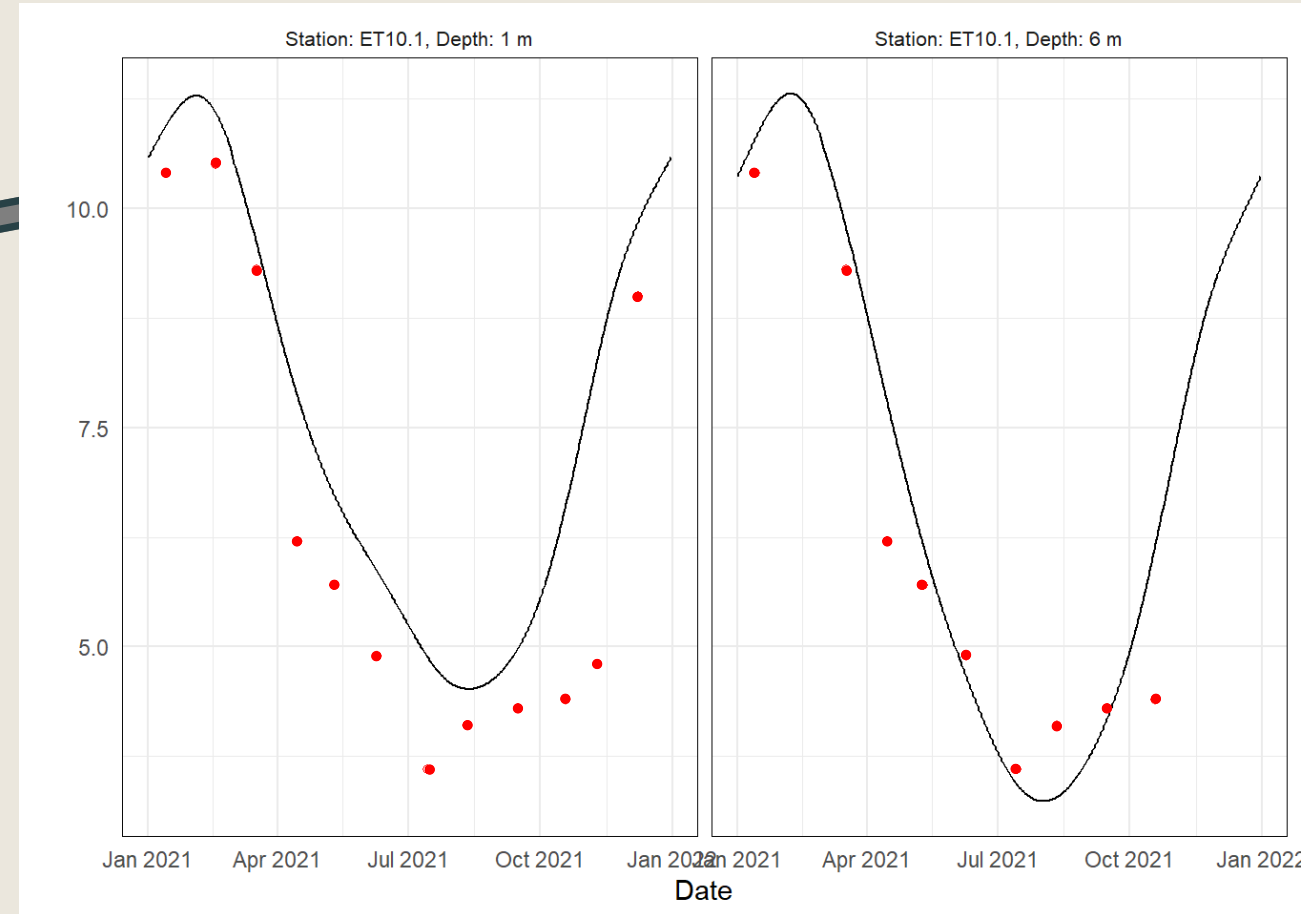




# Example: impact at one station of region size

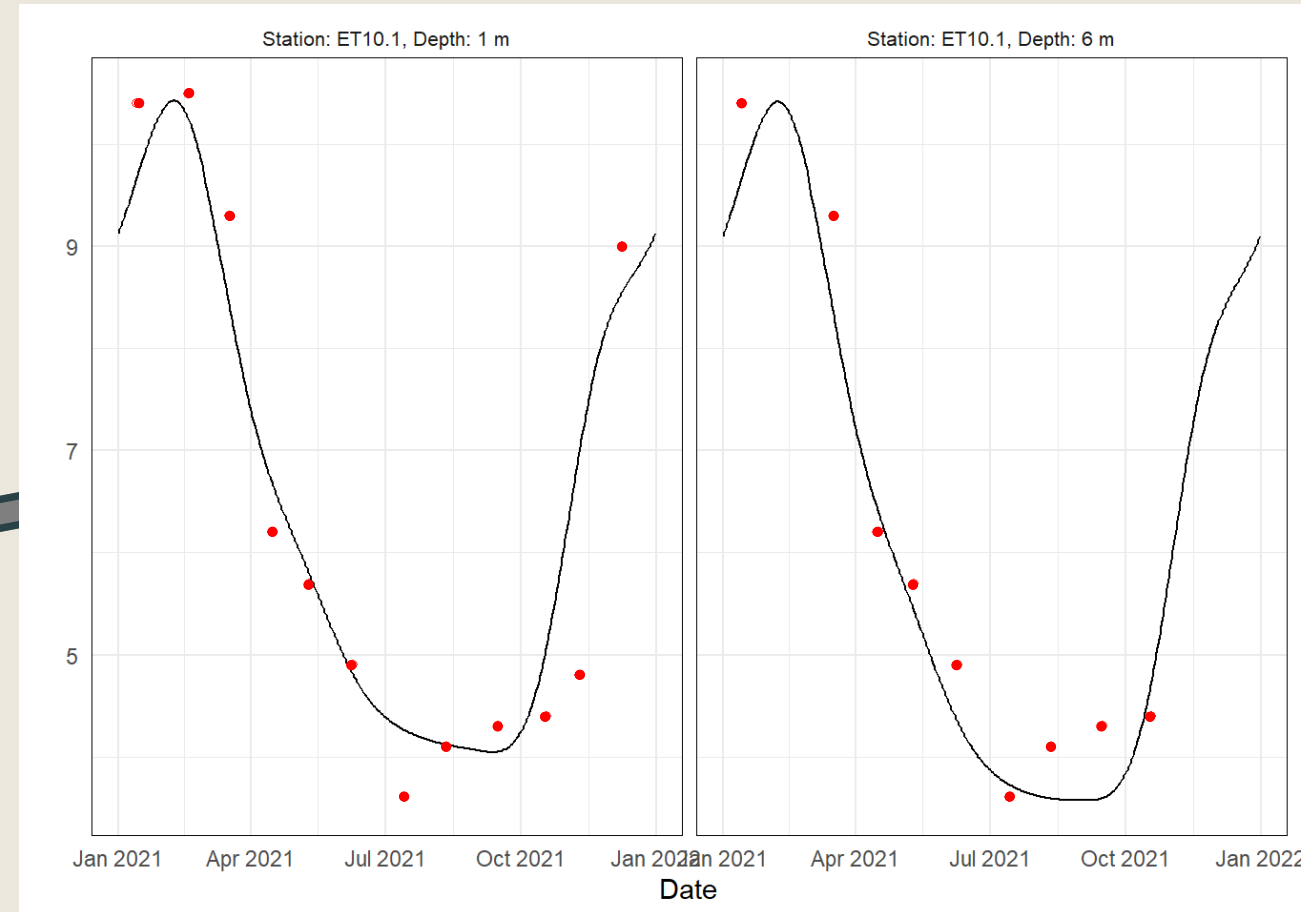
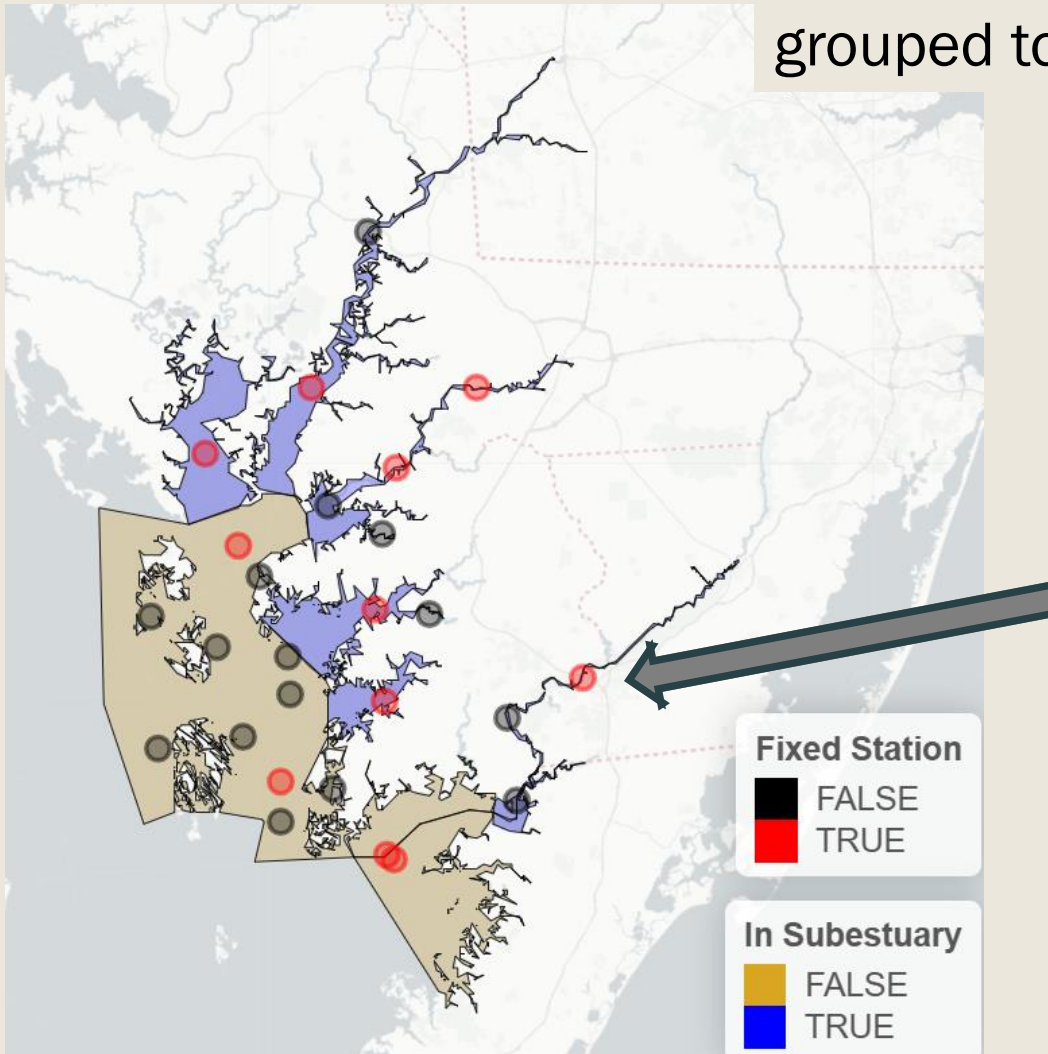


B. Large groups – segment is with Tangier and other eastern shore tribs, but also has boundary in main bay



# Example: impact at one station of region size

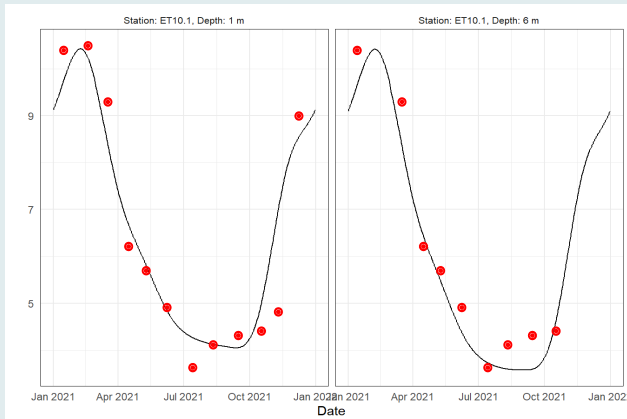
D. Selected Seg. Interp. Regions: Eastern shore tributaries grouped together, with only nearby boundary



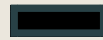
# Computing Root Mean Squared Error (RMSE) difference

ET10.1 in 2021

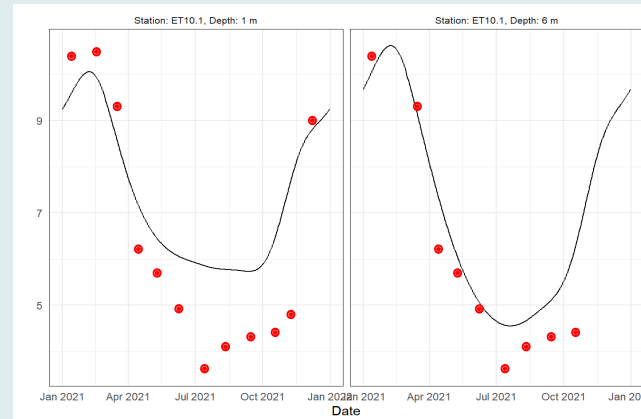
D. Selected Seg. Interp. Reg.  
RMSE 0.69



minus



A. Bay-wide GAM  
RMSE 1.4

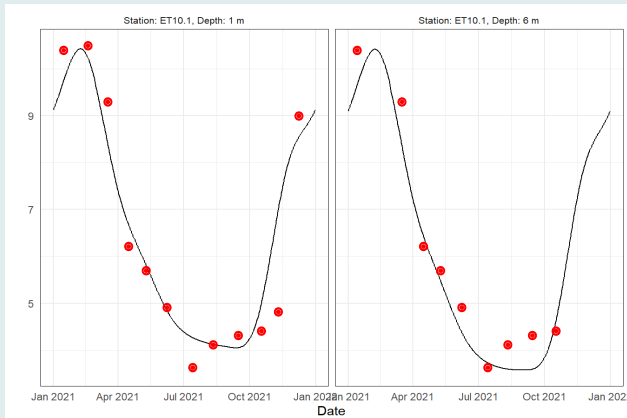


-0.71

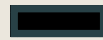
# Computing Root Mean Squared Error (RMSE) difference

ET10.1 in 2021

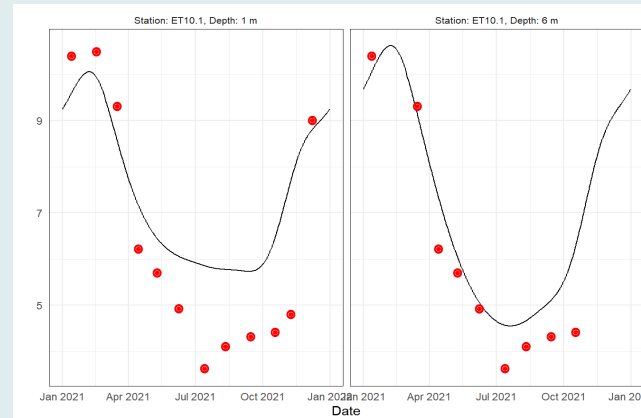
D. Selected Seg. Interp. Reg.  
RMSE 0.69



minus







A. Bay-wide GAM  
RMSE 1.4



-0.71



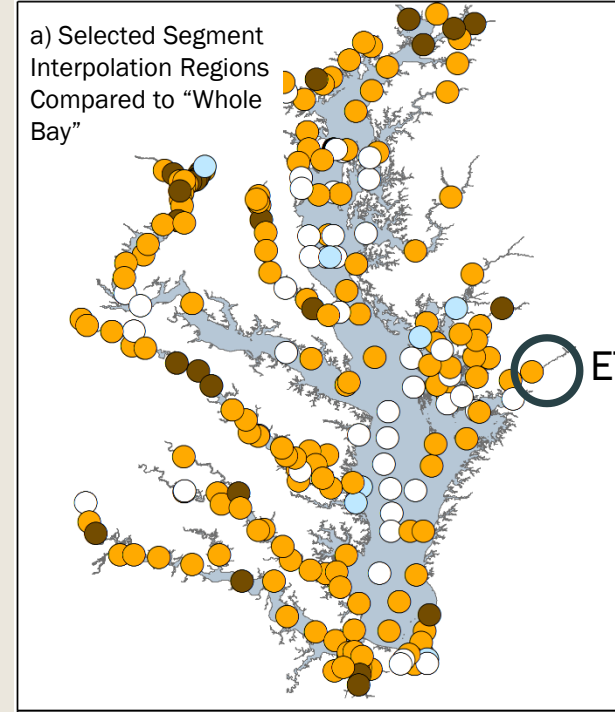
Each station's RMSE difference indicated with a symbol:

-  Brown and orange = The selected 31 Seg. Interp. Regions are LOWER error
-  White = almost the same
-  Blue = Selected Regions are HIGHER error
-  Black = GAM unable to be fit

# All stations

(D) Selected  
31 Seg. Interp. Regions  
= (A) Whole  
bay together =

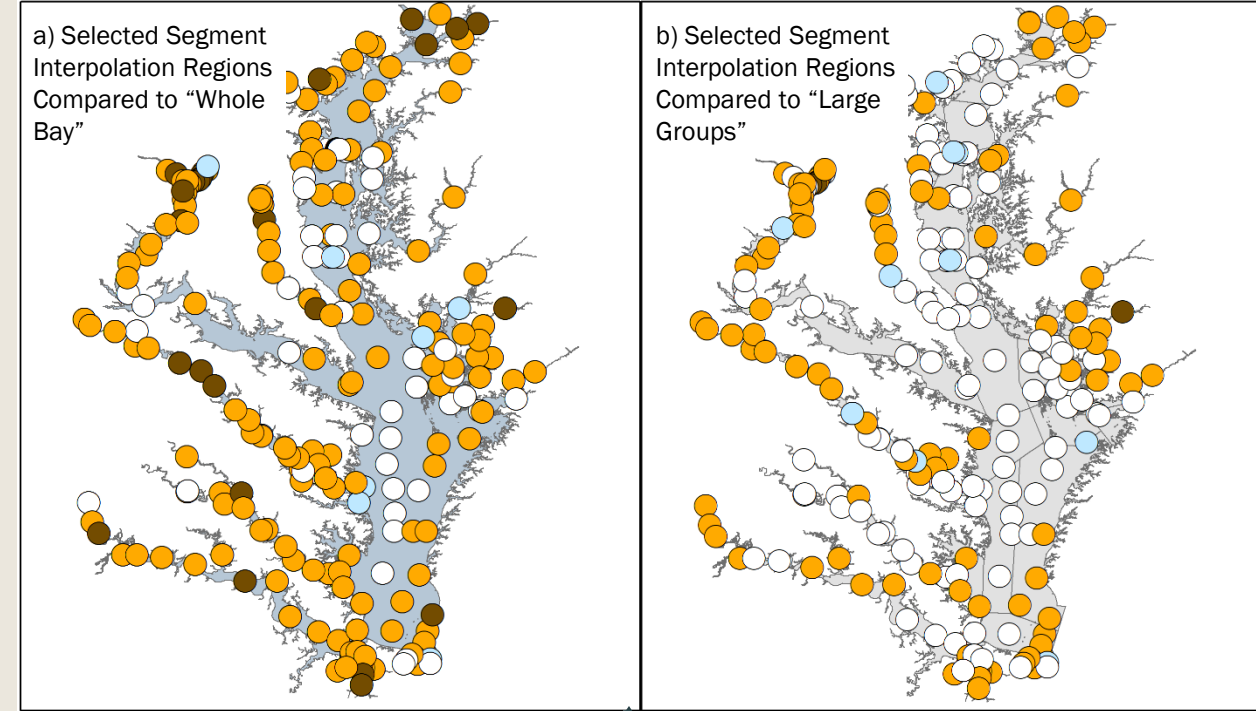
*Majority of stations have better interpolation results with smaller regions.*



- Brown and orange = The selected 31 Seg. Interp. Regions are LOWER error
- White = almost the same
- Blue = Selected Regions are HIGHER error
- Black = GAM unable to be fit

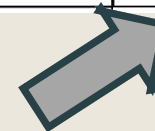


# All stations



(D) Selected  
31 Seg. Interp.  
Regions

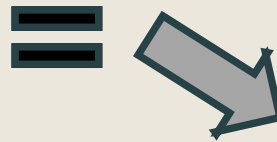
(B) 11 Large  
groups






- 
- Brown and orange = The selected 31 Seg. Interp. Regions are LOWER error
  - White = almost the same
  - Blue = Selected Regions are HIGHER error
  - Black = GAM unable to be fit

# All stations

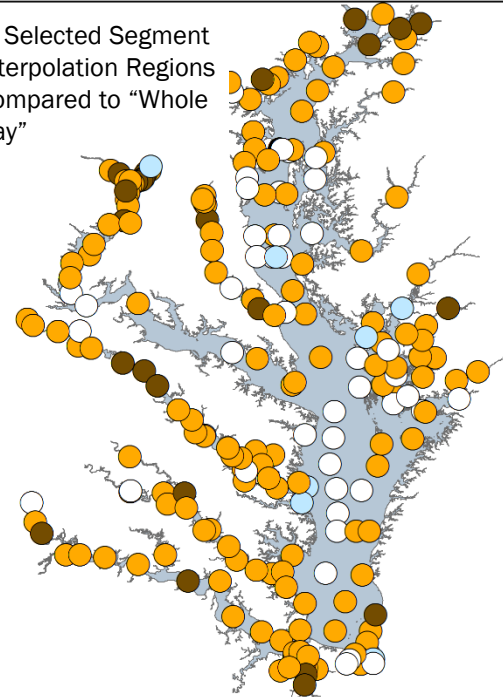
D) Selected  
31 Seg. Interp. Regions  
= Individual  
Segments

(C) 79

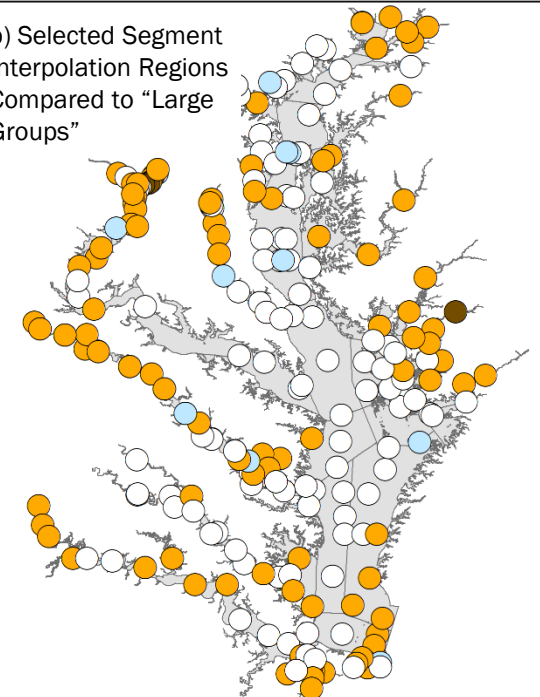


-  Brown and orange = The selected 31 Seg. Interp. Regions are LOWER error
-  Orange
-  White = almost the same
-  Blue = Selected Regions are HIGHER error
-  Black = GAM unable to be fit

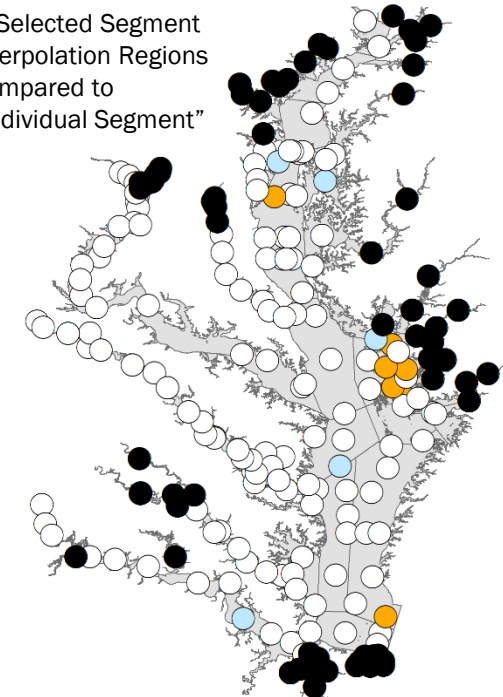
a) Selected Segment Interpolation Regions Compared to "Whole Bay"



b) Selected Segment Interpolation Regions Compared to "Large Groups"



c) Selected Segment Interpolation Regions Compared to "Individual Segment"



*Conclusion: Small regions are best, but they need to be large enough to fit the interpolation equation.*

# Summary/discussion

- Evaluation of multiple options resulted in current set of 31 Segment Interpolation Regions. Regions contain on average 3 segments grouped together (range =1 to 11), plus boundary.
- Possible discussion
  - *Impact of needing data across state lines.*
  - *The tool will identify if an individual segment does not have data in a year, since in that case it should not be assessed.*

