

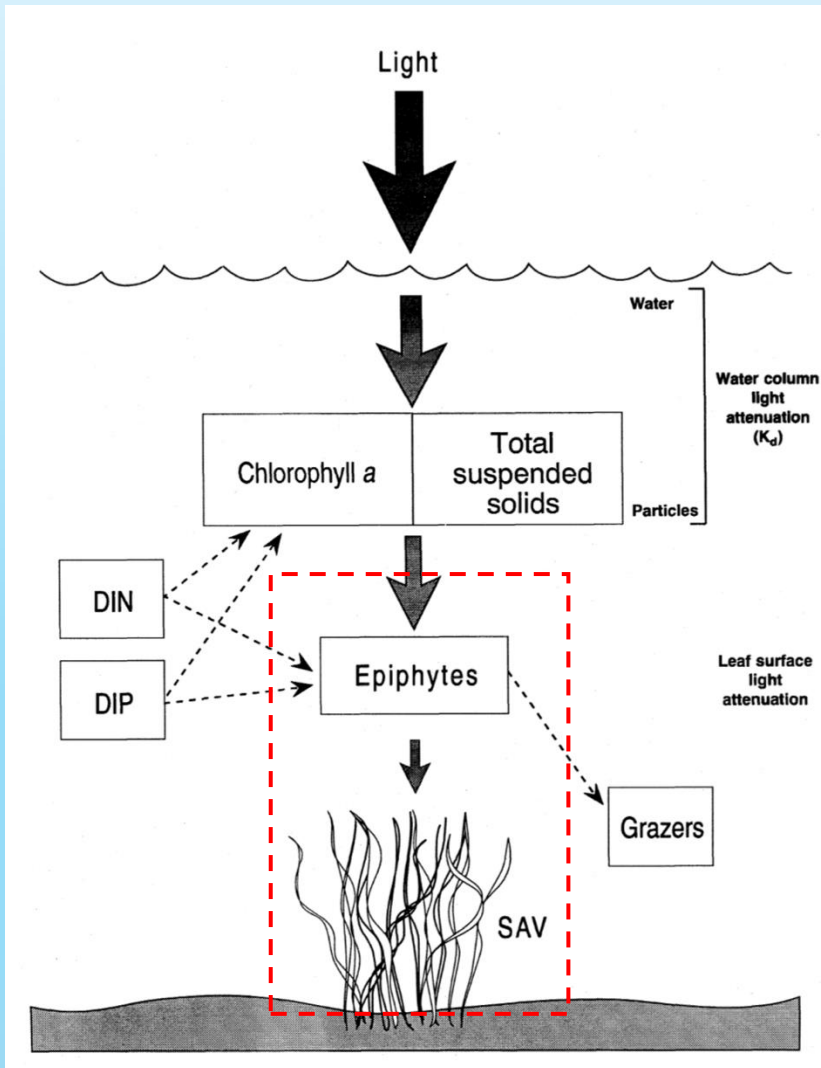
SAV Modeling Approach

Quarterly Review (7/14-15/2026)

Jian Shen, Joseph Zheng, Zhengqui Wang

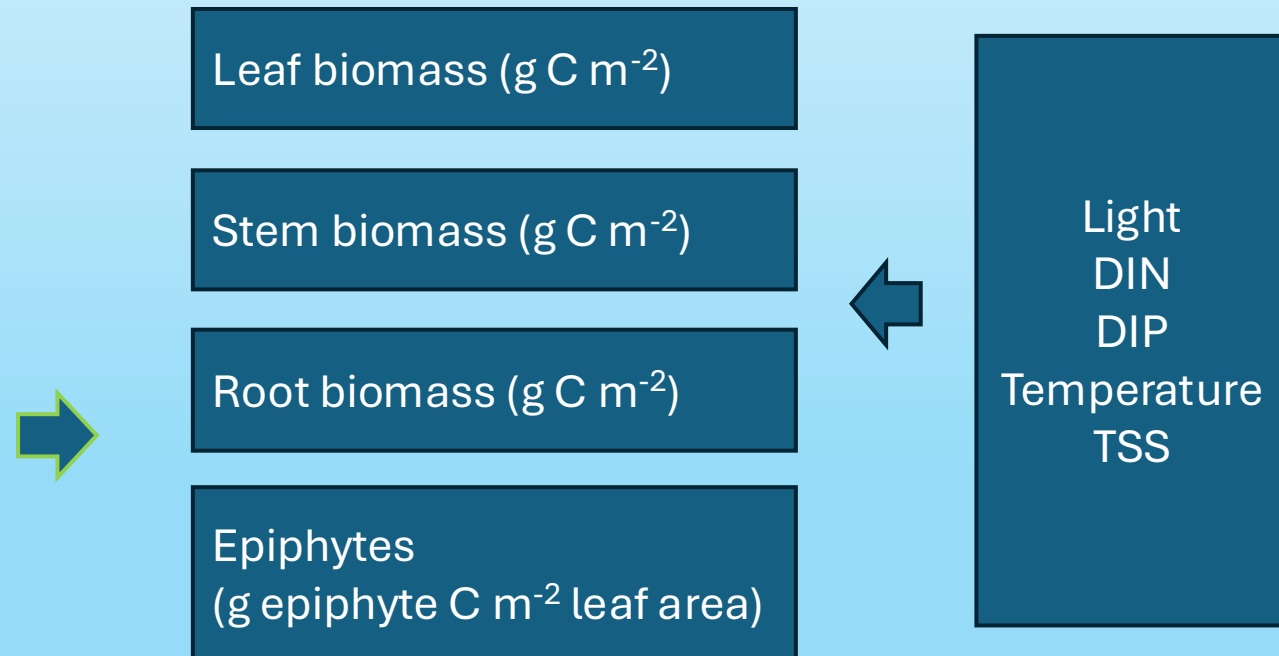
SAV Model

Conceptual Model



Dennison et al., 2008

SAV Numerical Model



Cerco et al. 2010

Essential Needs for Modeling SAV

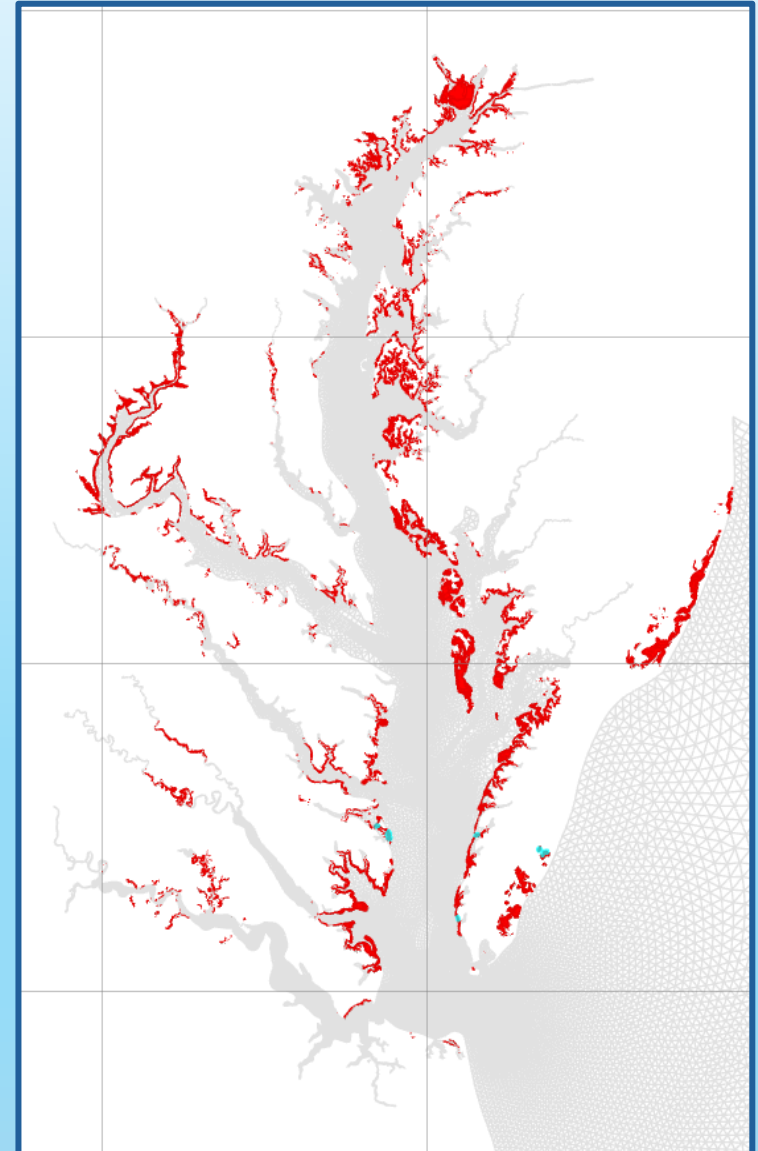
Estimate SAV biomass

Select SAV species

Convert biomass to area

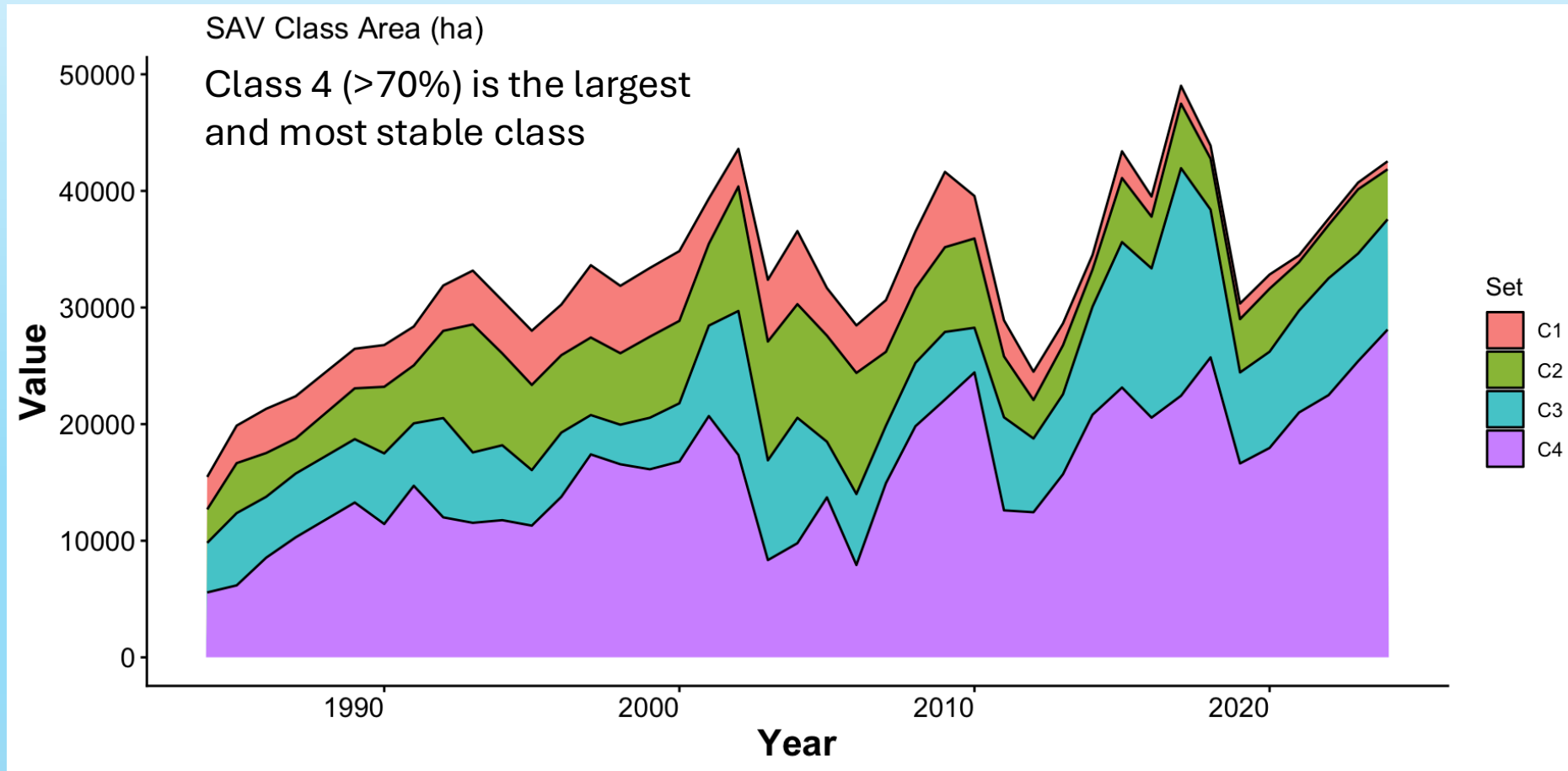
Set initial conditions for
the model

Model verification



Available Data

- Annual Distribution of SAV for each density class



Available Data

- Percentage areas for each class are reported by VIMS for each Bay segment

[Download Data](#) Year: 2024 Zone: Tidal Fresh

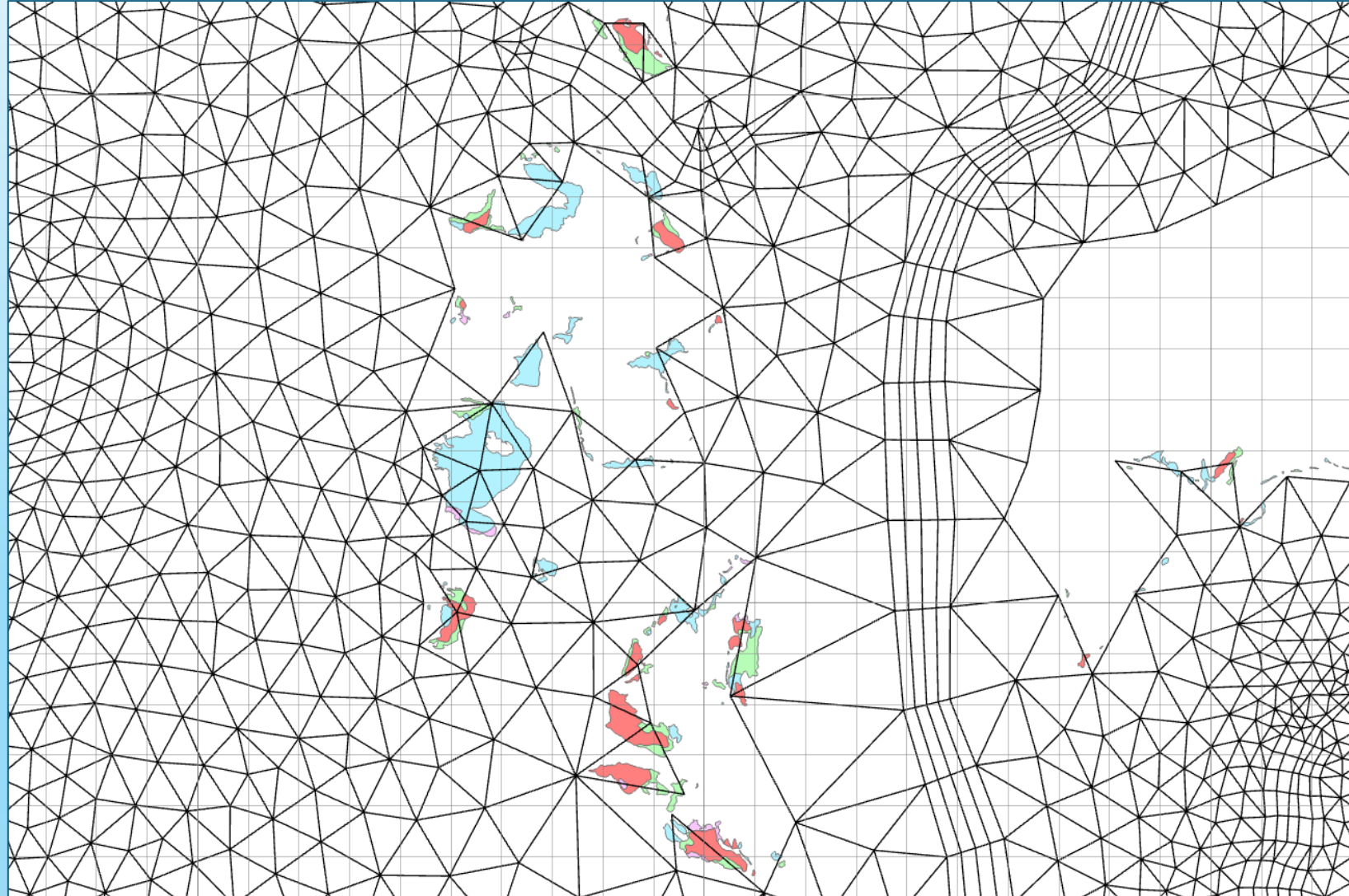
☒ Salinity Zones ☐ Geographic Zones

Segment	Class 1		Class 2		Class 3		Class 4		Total
CB1TF1	0.72	0 %	5.92	2 %	28.48	11 %	233.67	87 %	268.79
CB1TF2	31.69	1 %	267.97	6 %	110.74	3 %	4,010.94	91 %	4,421.34
NORTF	0.61	1 %	2.59	4 %	19.25	32 %	36.99	62 %	59.44
CHSTF	0.00	0 %	0.00	0 %	0.00	0 %	1.11	100 %	1.11
CHOTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
NANTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
POCTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
PAXTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
WBRTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
POTTF	0.00	0 %	15.26	1 %	89.83	5 %	1,536.56	94 %	1,641.65
MATTF	0.00	0 %	38.26	11 %	36.14	10 %	278.54	79 %	352.94
ANATF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00
PISTF	0.00	0 %	2.92	4 %	1.19	2 %	61.16	94 %	65.27
RPPTF	0.37	0 %	19.82	2 %	73.96	9 %	699.71	88 %	793.86
MPNTF	0.00	0 %	0.00	0 %	16.92	92 %	1.52	8 %	18.44
PMKTF	6.58	2 %	15.17	4 %	65.95	17 %	302.33	78 %	390.03
JMSTF1	0.87	1 %	24.36	14 %	60.20	36 %	83.88	50 %	169.30
JMSTF2	0.00	0 %	0.98	100 %	0.00	0 %	0.00	0 %	0.98
APPTF	0.00	0 %	0.00	0 %	0.00	0 %	0.00	0 %	0.00

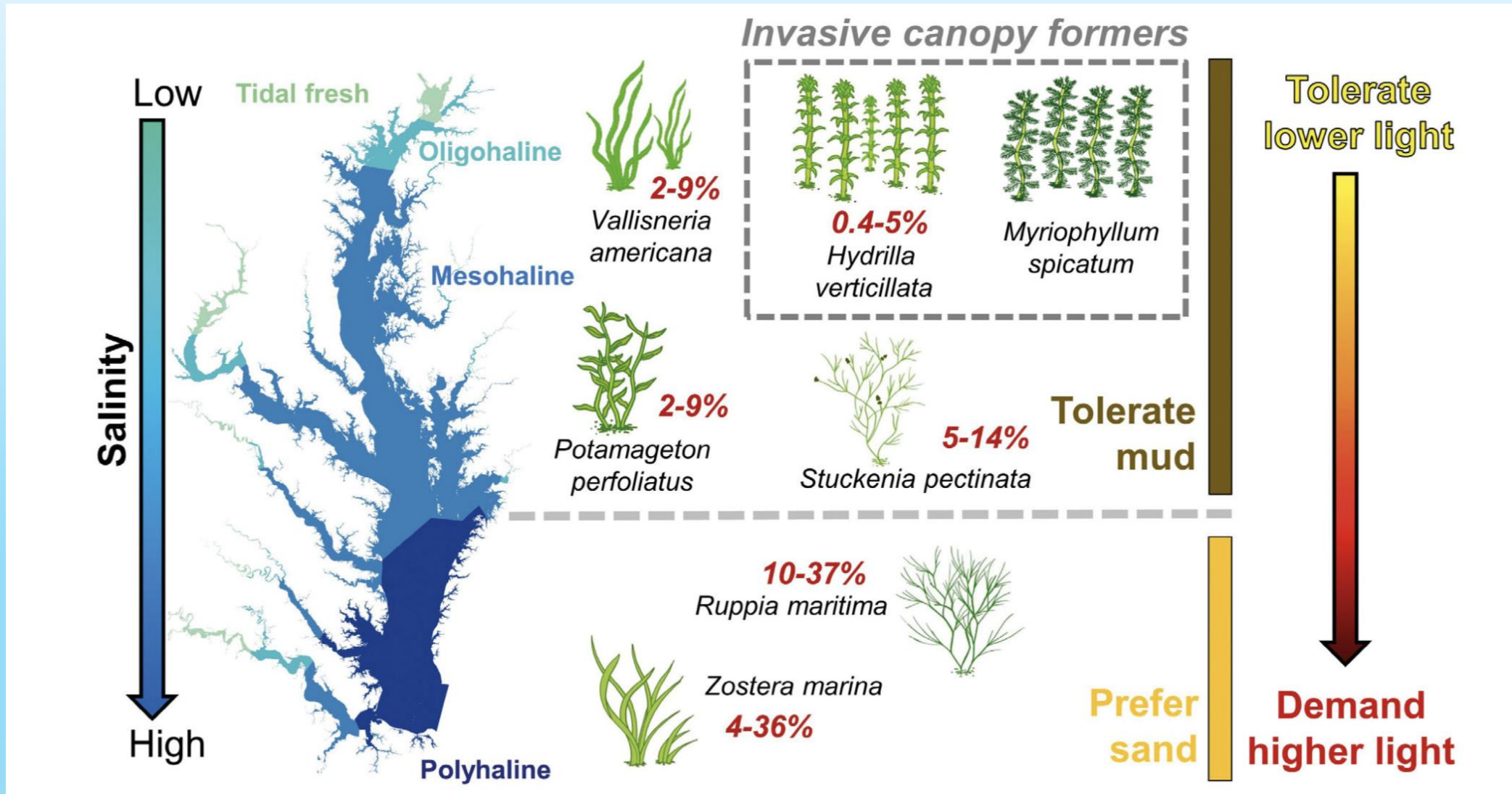
Areas in Hectares. "nd" indicates that the area was not mapped. "pd" indicates that the area was not fully mapped.

Biomass and Density Consideration

- SAV Density is divided into 5 categories
 - 1=<10%
 - 2=10-40%,
 - 3=40-70%,
 - 4=>70%,
 - -1=Unclassified
- Our grid size is larger than SAV area, it is difficult to model each density class.
- We can simulate averaged density



Species Distribution across Salinity Region



Biomass Distribution

- Moore (2000) provided typical biomass distribution of SAV in different salinity regions

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K. A. Moore et al.

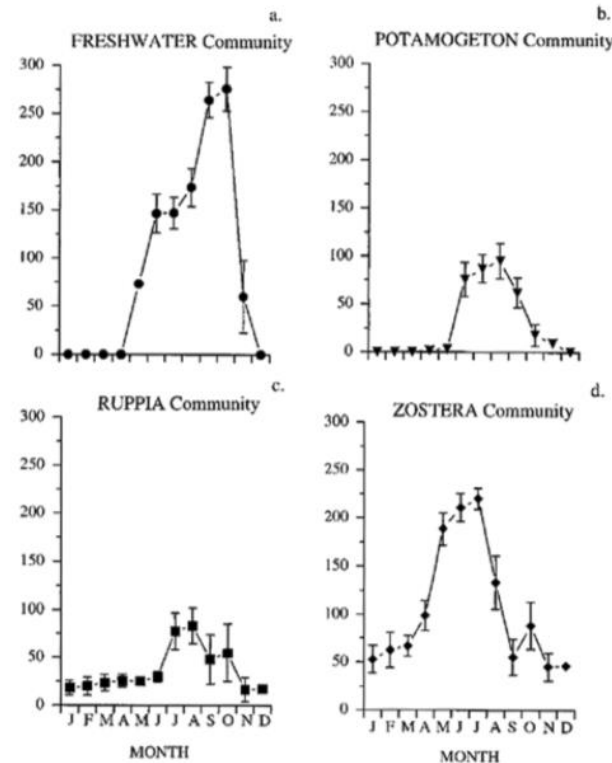


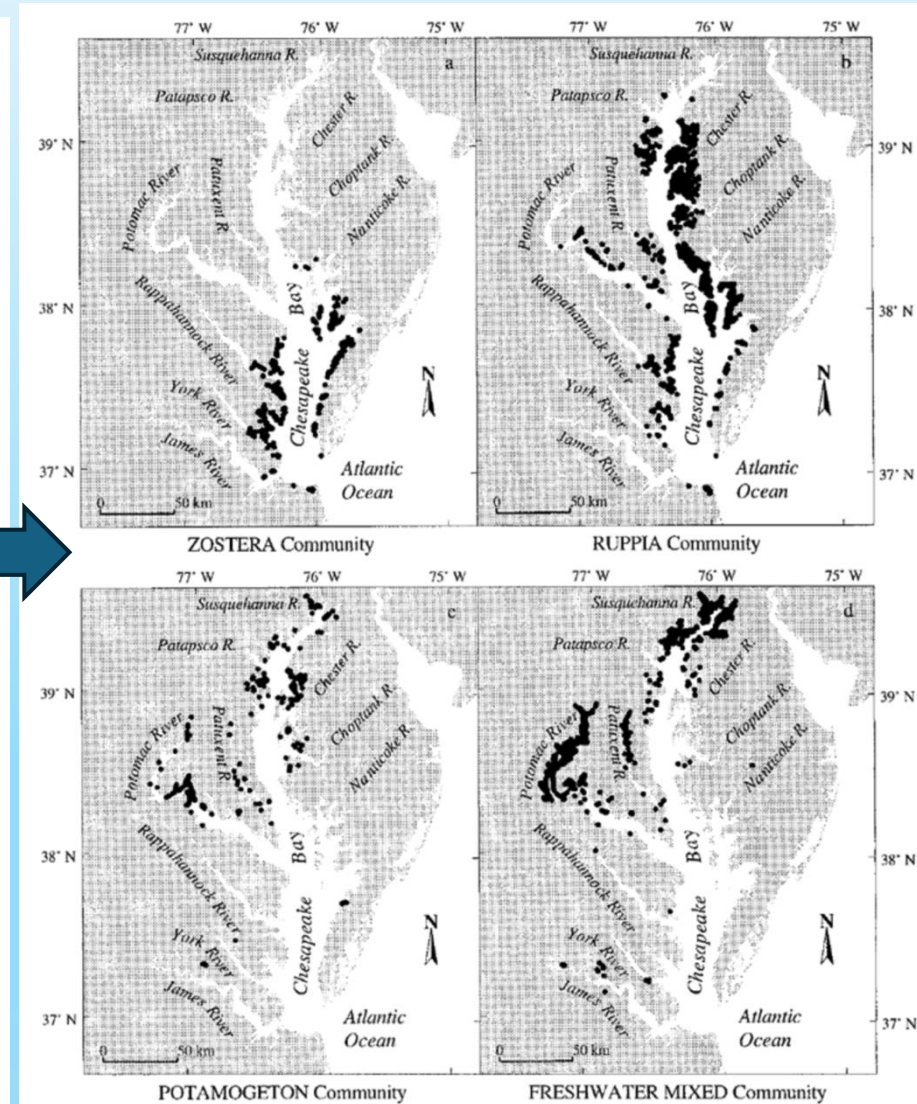
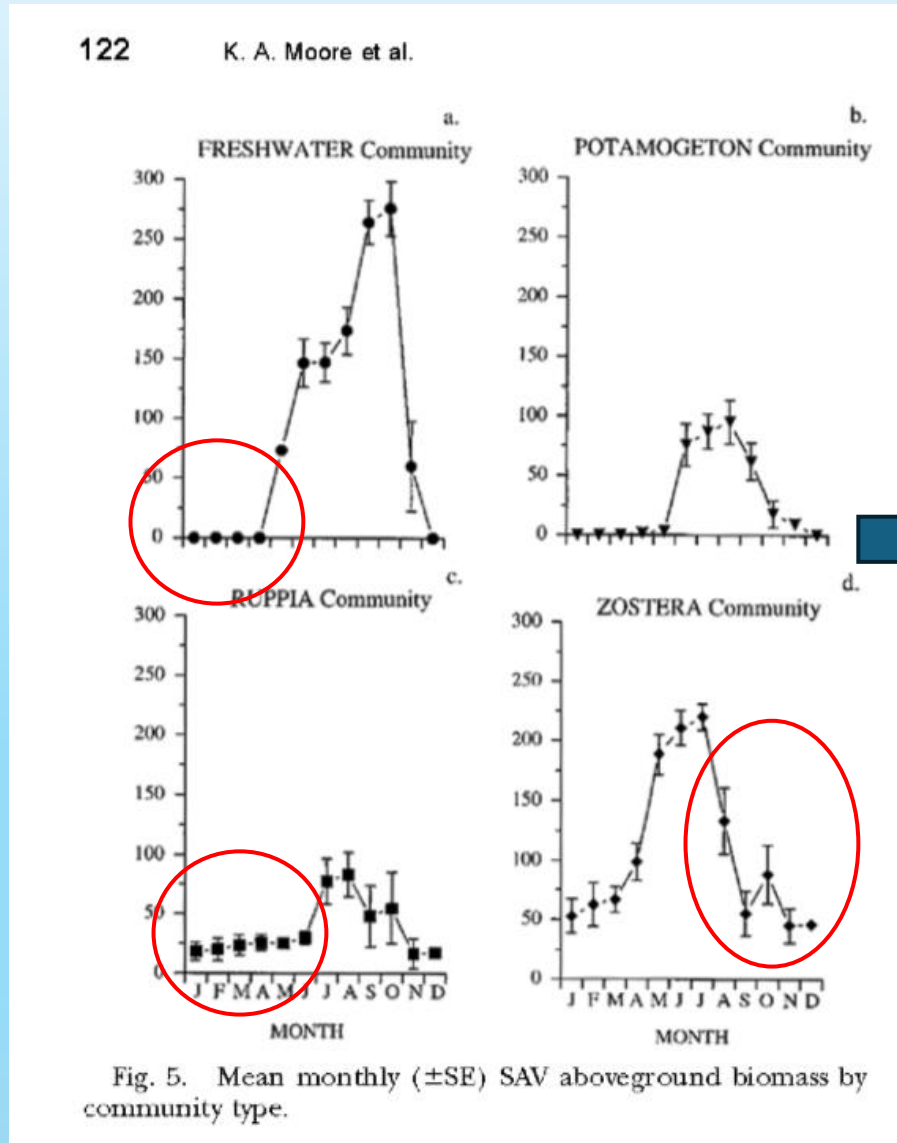
Fig. 5. Mean monthly ($\pm$ SE) SAV aboveground biomass by community type.

POTAMOGETON community reached a peak standing crop of 100 gdm m⁻² or more by August with complete loss by December. In contrast, shoot biomass of the FRESHWATER MIXED community increased throughout the summer and early fall, and reached an average maximum of nearly 300

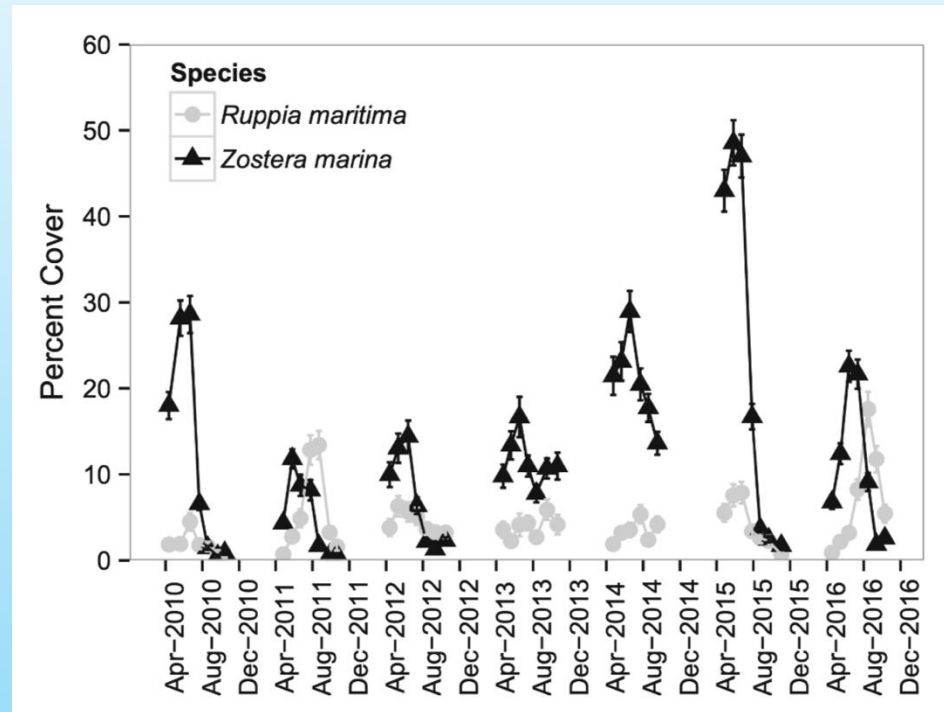
PERCENT COVER			DENSITY CLASS
5%			Very Sparse <10%
15%			
25%			
35%			Sparse 10 - 40 %
45%			
55%			
65%			Moderate 40 - 70%
75%			
85%			
95%			Dense 70 - 100%

Species Distribution

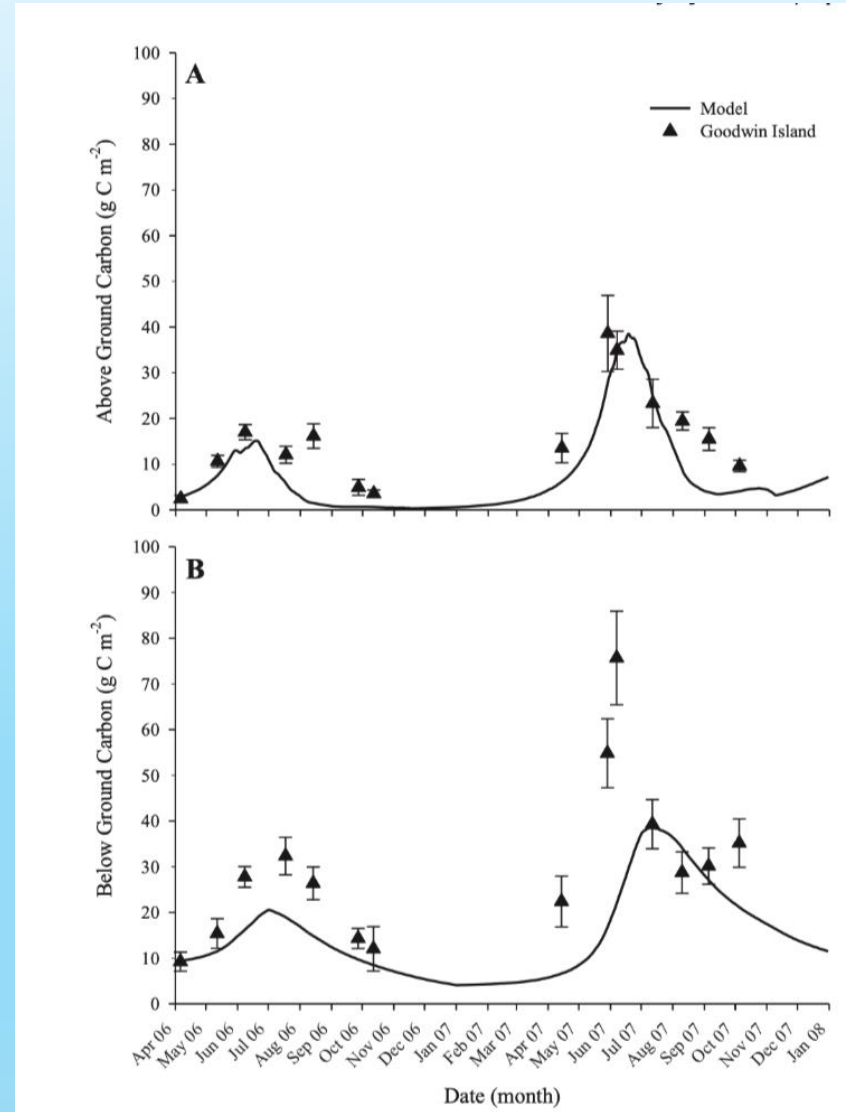
- Select typical 4 species for 4 salinity regions
- Key differences
 - Magnitude of peak biomass
 - Duration



Examples of Seasonal Variation



Shields et al., 2019

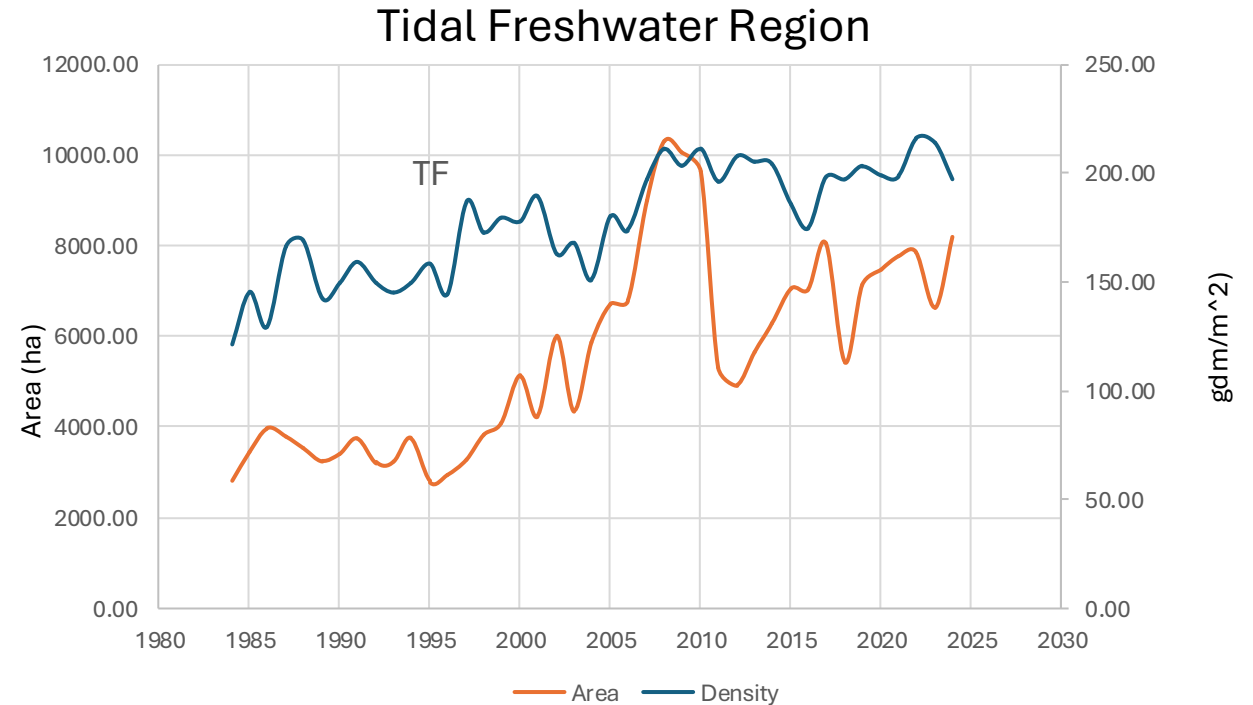
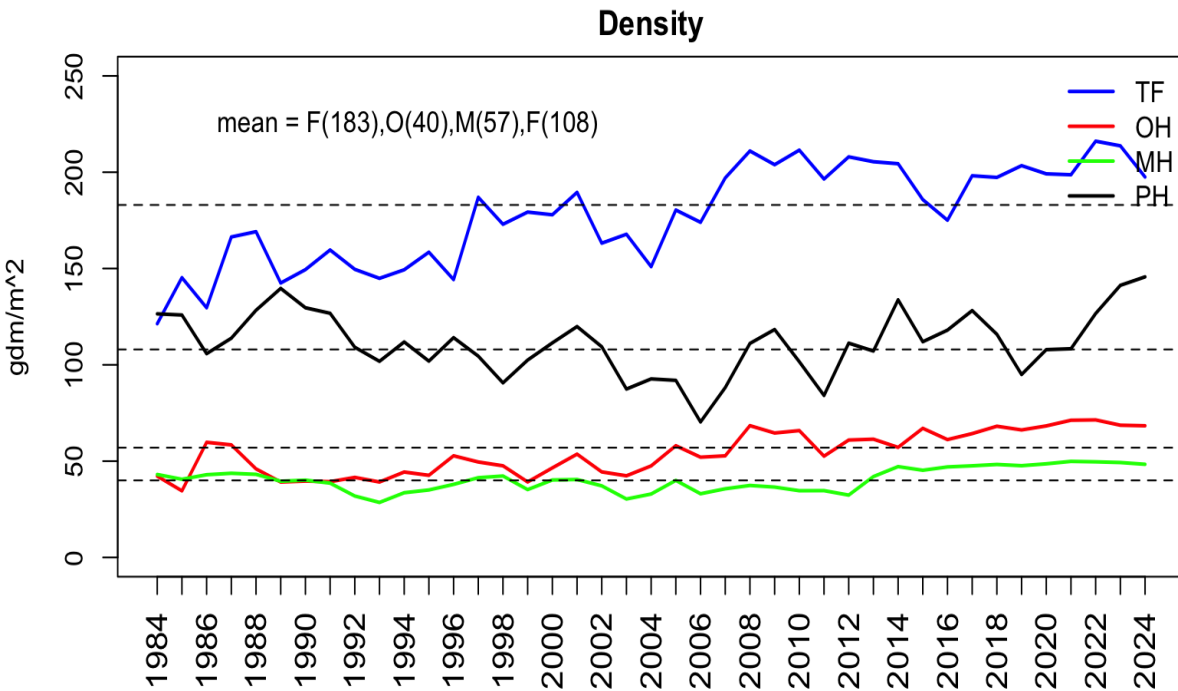


Jessie et al., 2014

Estimate of Typical Mean Biomass

- Use maximum biomass as the baseline biomass for each salinity regions (TF ~ 275 (gdm/m²), OH ~ 100, MH ~ 80, PH ~ 220)
- Biomass density ρ x % of each SAV class to estimate biomass for each SAV class
- Mean density is estimated as area weighted mean:
Density (ρ) = Sum(Density (i)x area(i))/Sum of total area

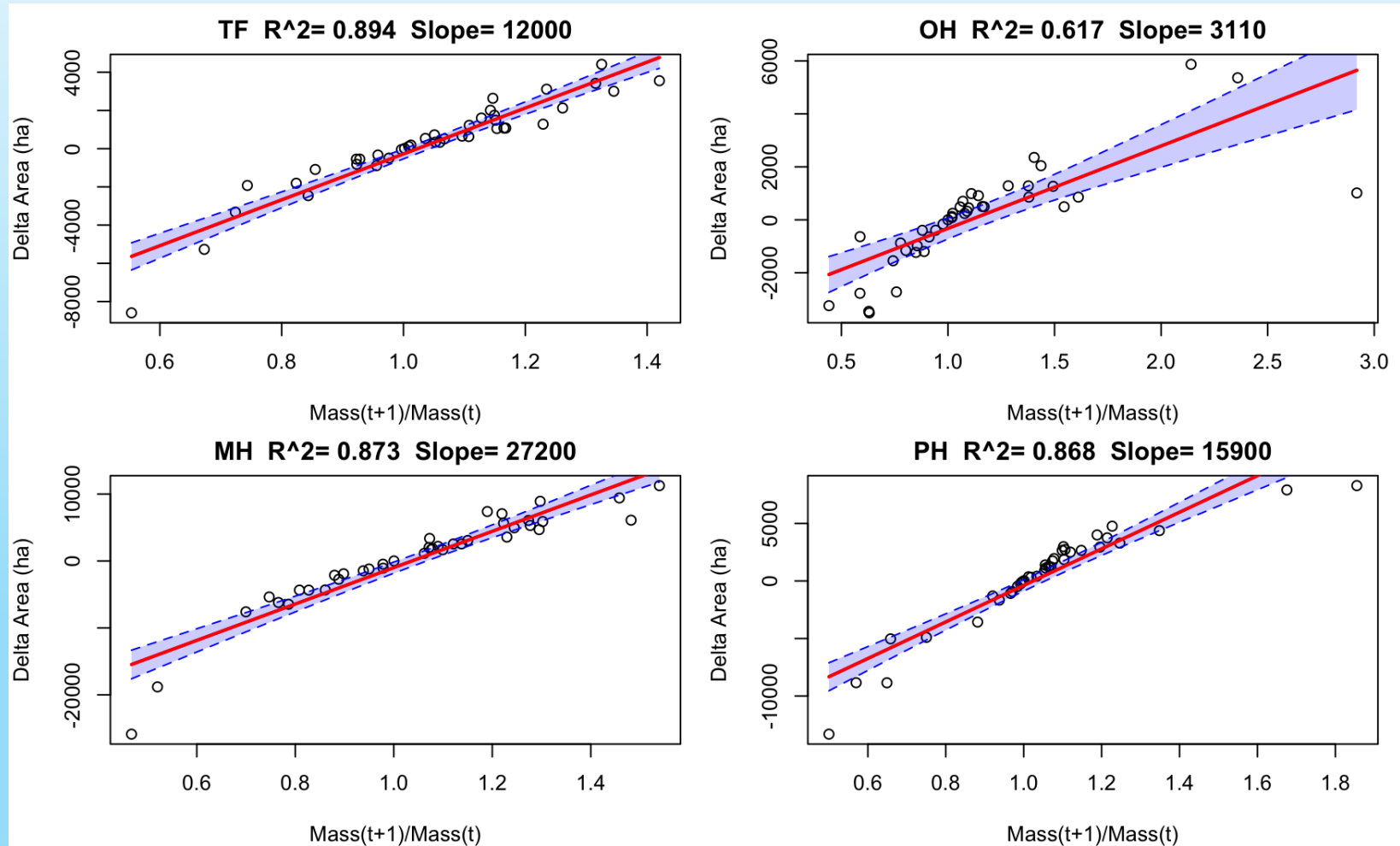
Distribution of Biomass



- Area has larger variation than density
- The difference can be partially attributed to the precondition

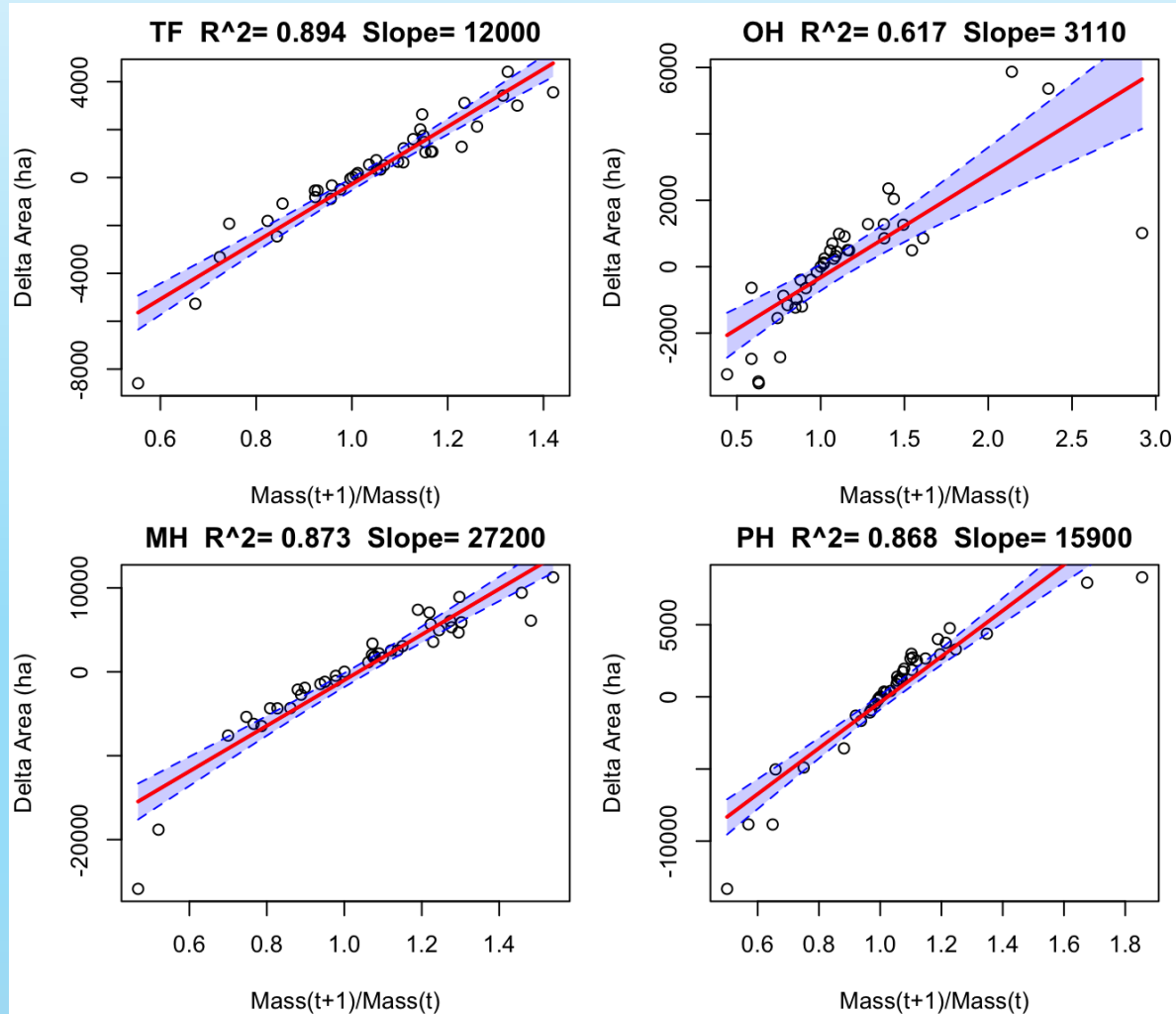
Estimated Biomass Ratio vs. Delta Area

- Assume average ρ does not change dramatically, ratio of biomass $r = \rho * A(t) / \rho * A(t-1)$ can quantify the changes in biomass
- $\rho(t)/\rho(t-1)$ ranges from 0.77-1.72
- On average, $r \sim (A(t+1)-A(t))$ has a good linear correlation
- It suggests that we can use this relationship to Convert biomass to area and match the mean condition in each salinity region

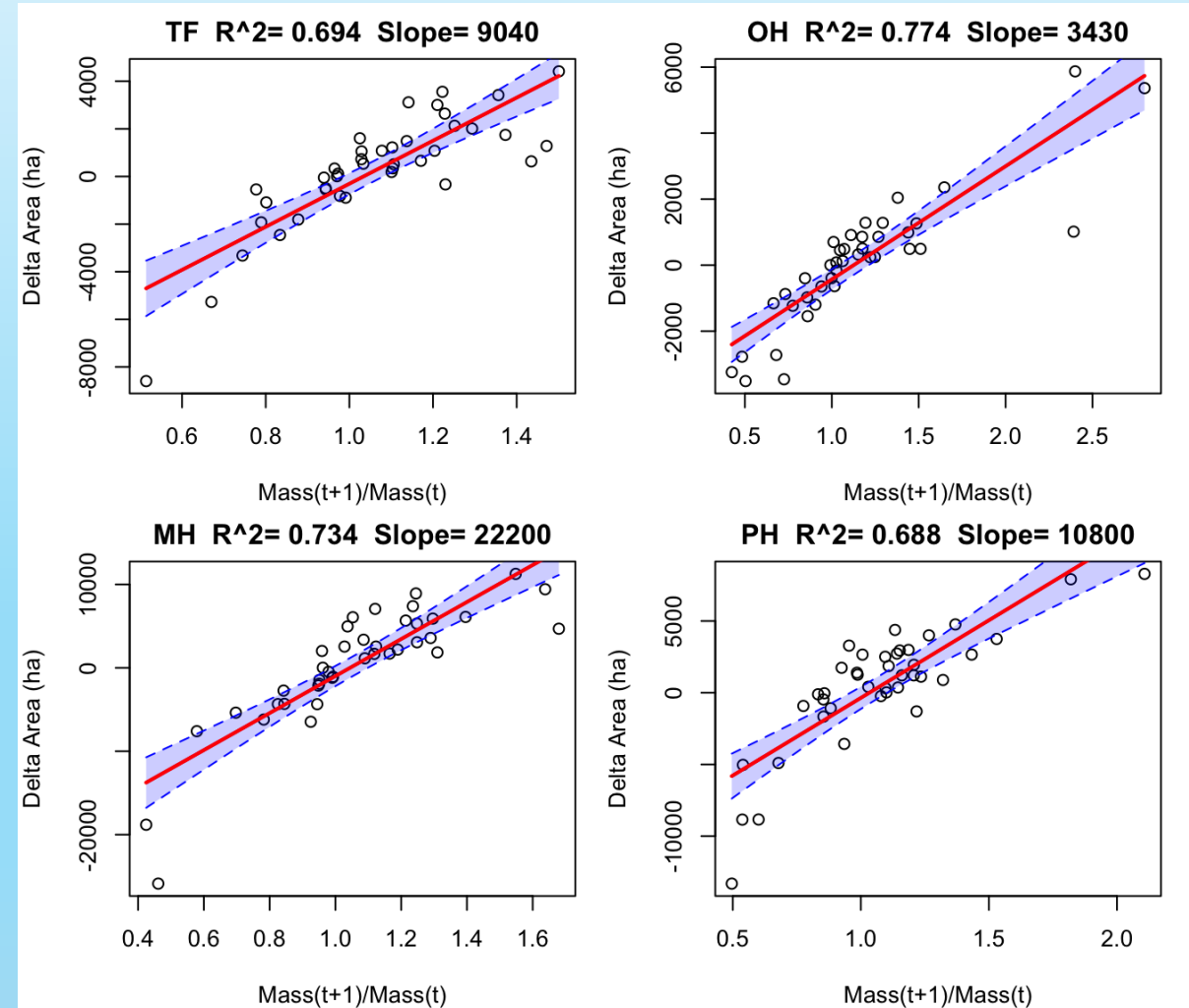


Compare using Constant and Varying Density

Constant density

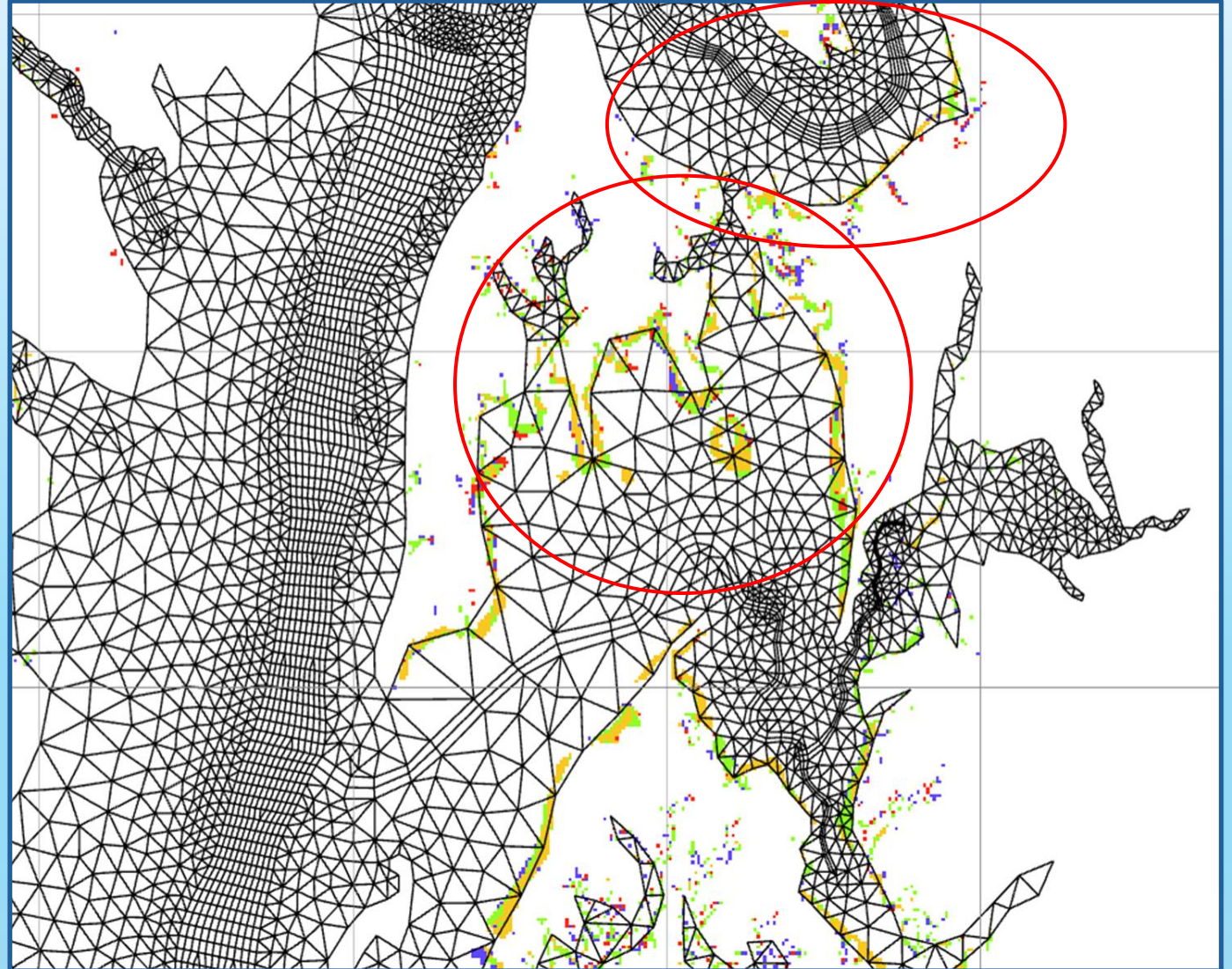


Varying density

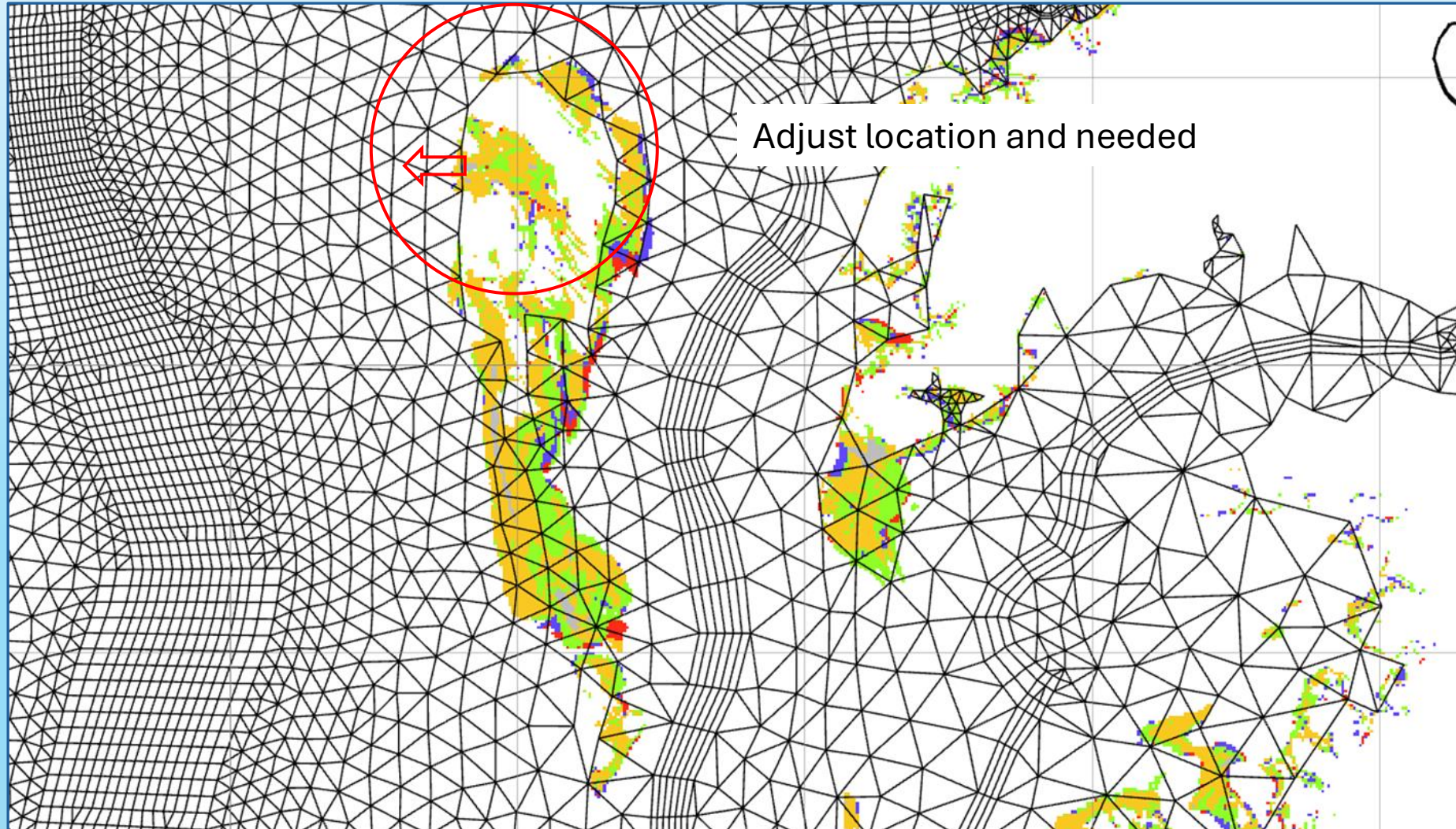


Initial Condition of SAV

- Convert observed SAV area to 100 x100 m grid
- Set criteria to include all the SAV outside of the model grid to the nearest model grid.
- Estimate total area in each grid (count 100x100 grids)
- Use typical mean density to estimate the initial condition

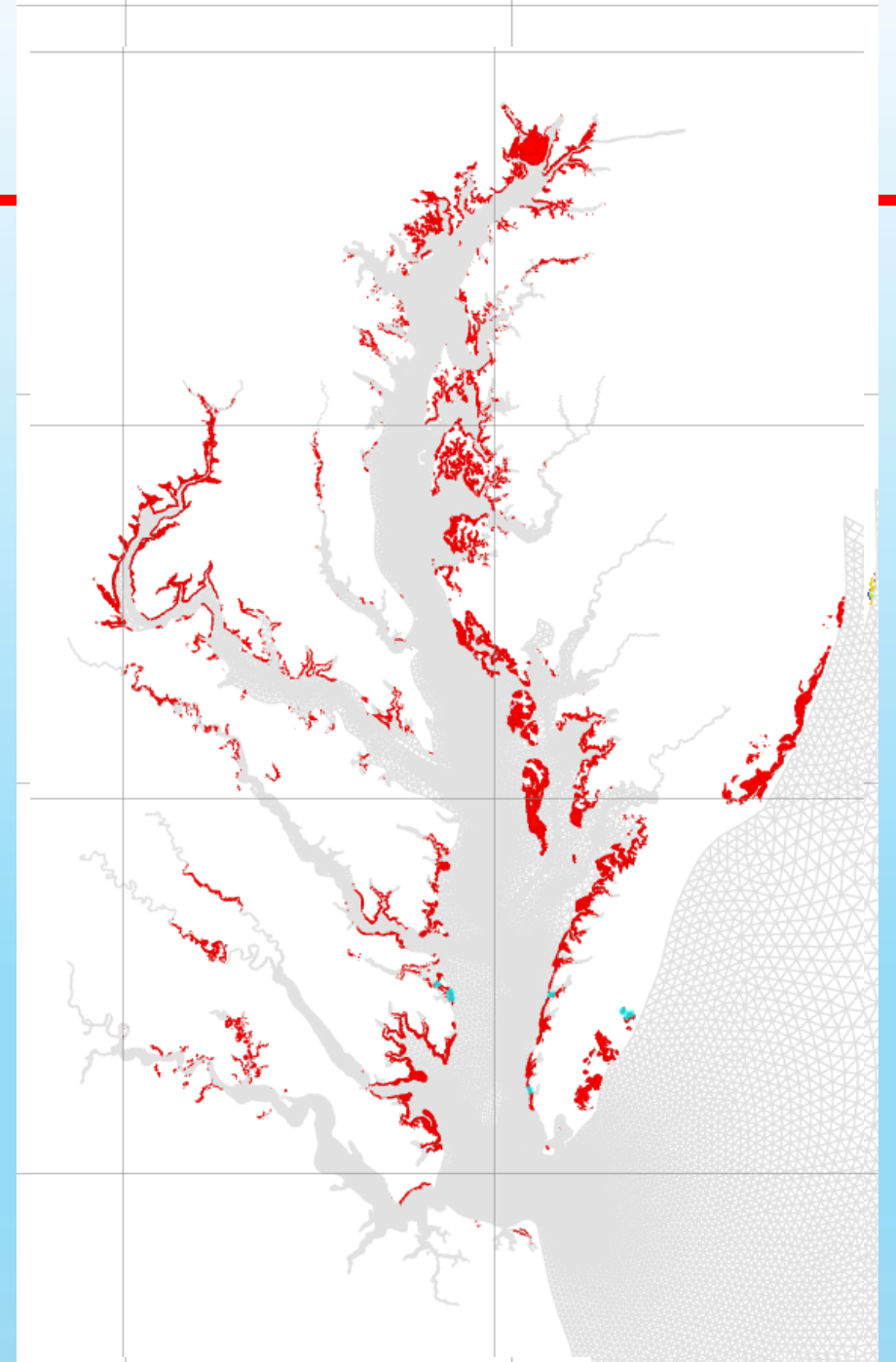


Add SAV to Model Grid

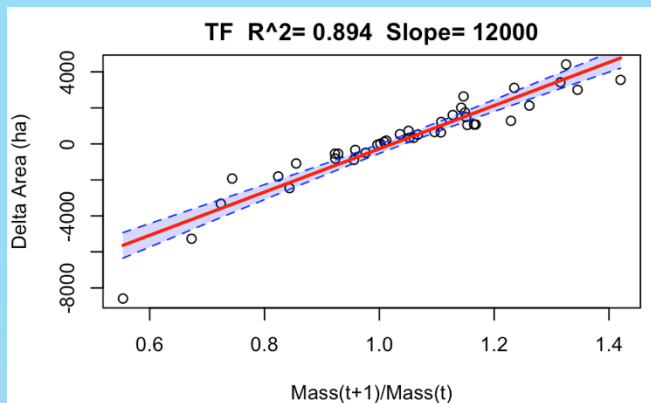
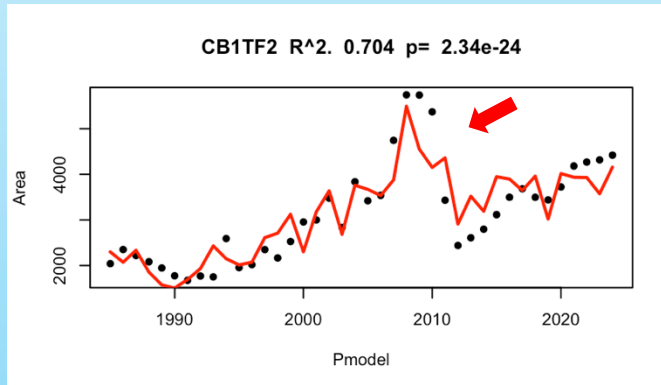
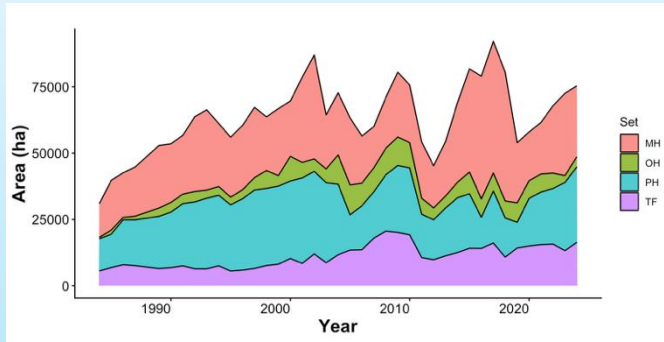


Designate Area for SAV

- Not all shallow water have SAV
- We plan to include the largest area (1990-2020) to set SAV and corresponding biomass
- We will do sensitivity test to see if SAV will survive in the region that currently has no SAV
- Because model does not consider the bed condition, SAV will likely to grow if salinity, temperature, and light are favorable
 - Under which bed condition preventing SAV growth?



Model Calibration and Verification



Set initial SAV
in different
salinity region
with typical
species

Select a
typical year
(e.g., average
temperature
condition)

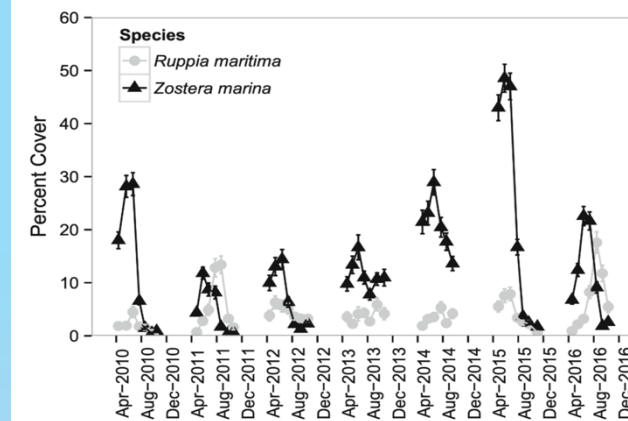
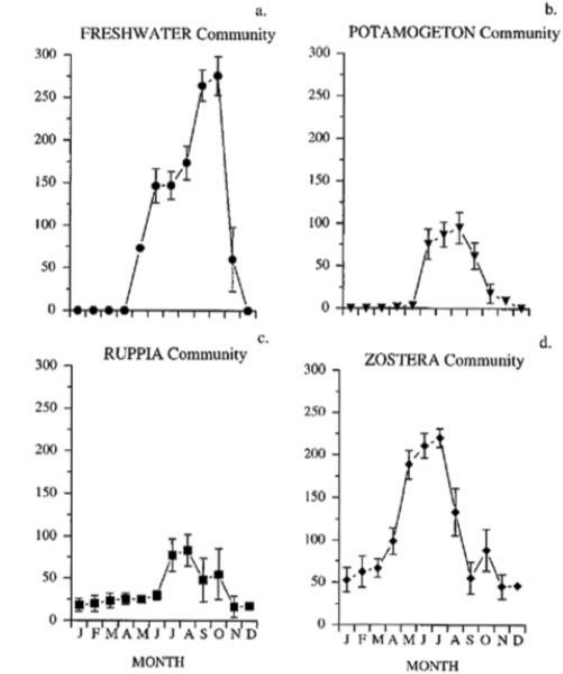
Compare
observations.
Focuses on
key events of
changes in
SAV area

Calibrate
model to
match typical
SAV in
different
salinity
regions

Use ratio ~
delta area to
convert
biomass to
area

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Summary Modeling Approach

- Select 4 species for 4 salinity regions (tidal fresh, oligohaline, mesohaline, polydalin)
- Use observations to obtain weighted biomass (gdm/m^2)
- Conver SAV area to 100m x 100m grid
- Use defined criteria to include SAV outside of the model grid into the model grid
- Count the area and biomass in each grid
- Calibrate SVA in four regions in average year (mean flow and temperature)
- Use varying parameters to account for species variations
- Convert model biomass to area (using typical linear relationship)
- Compare model results to SAV area for salinity region (or Bay segment)