

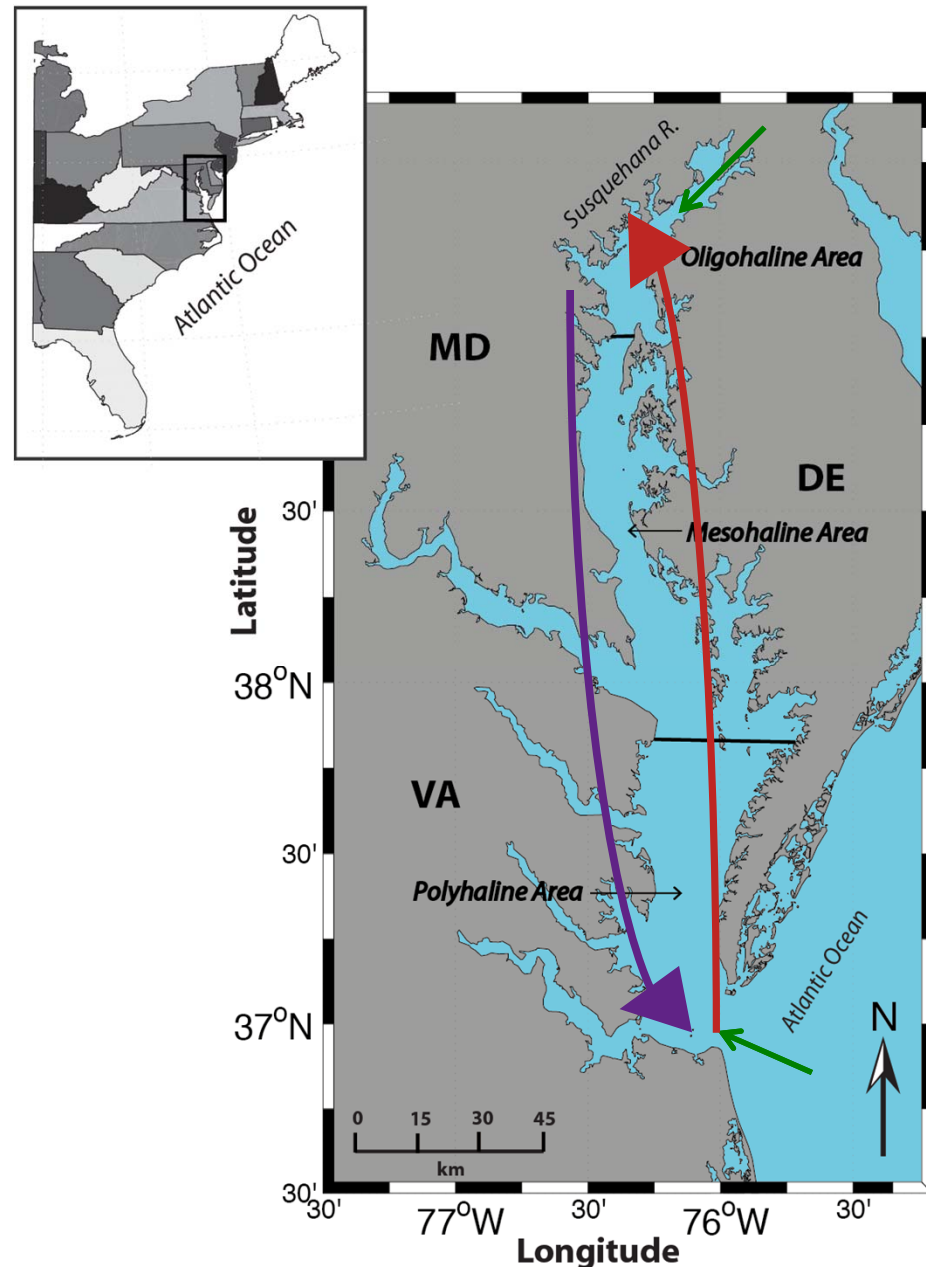
Remote Sensing of Environmental Vibrio in the Chesapeake Bay

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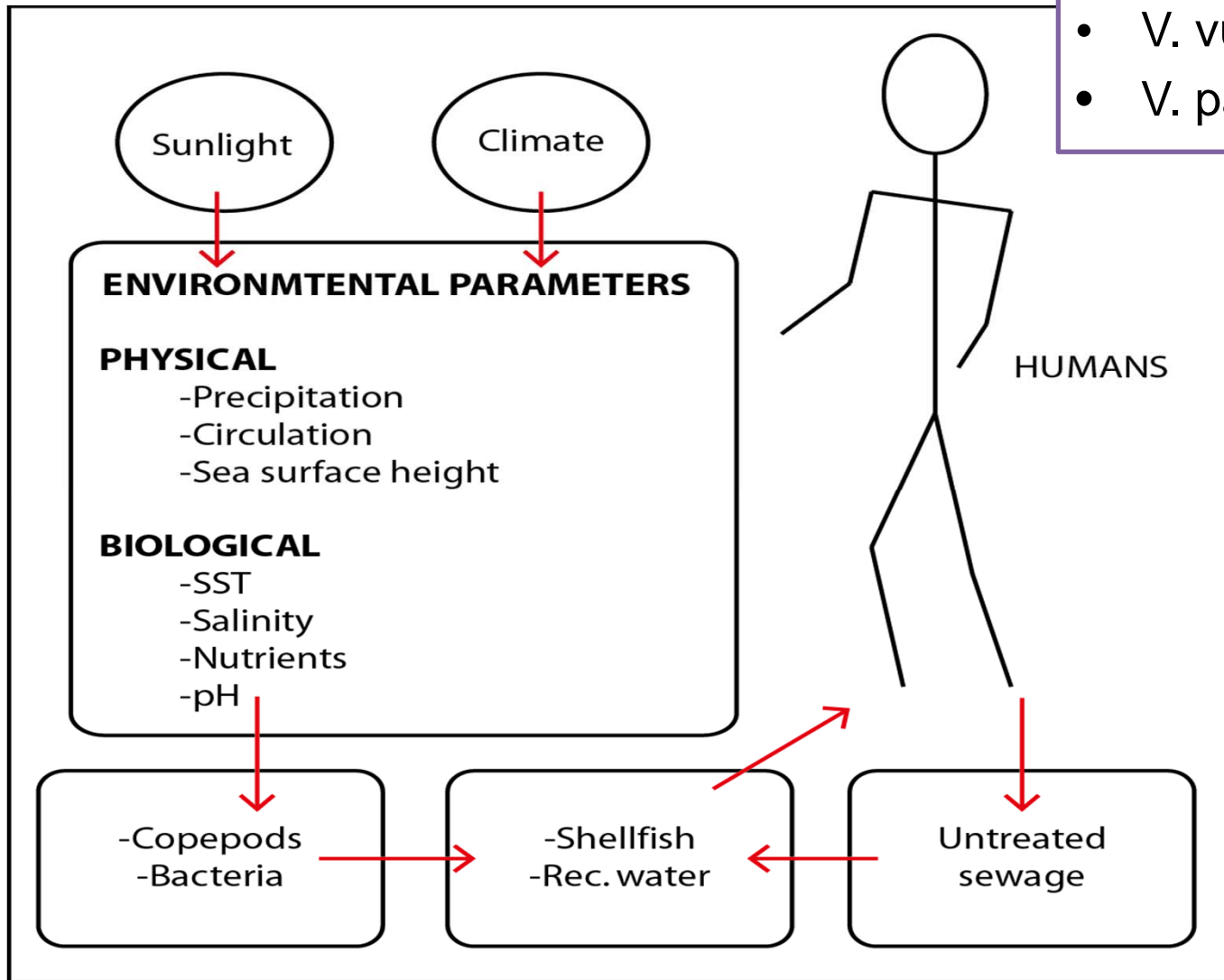
Study Area (The Chesapeake Bay)

- **Salinity gradient**
 - Oligohalic (0-6 ppt)
 - Mesohalic (6-18 ppt)
 - Polyhalic (18-30+ ppt)
- **Sea surface temperature**
 - -0.5°C to 31°C
- **Major inputs**
 - Atlantic Ocean
 - Susquehanna River
- **2-Layer gravitational circulation scheme**



Vibrio in the Chesapeake Bay

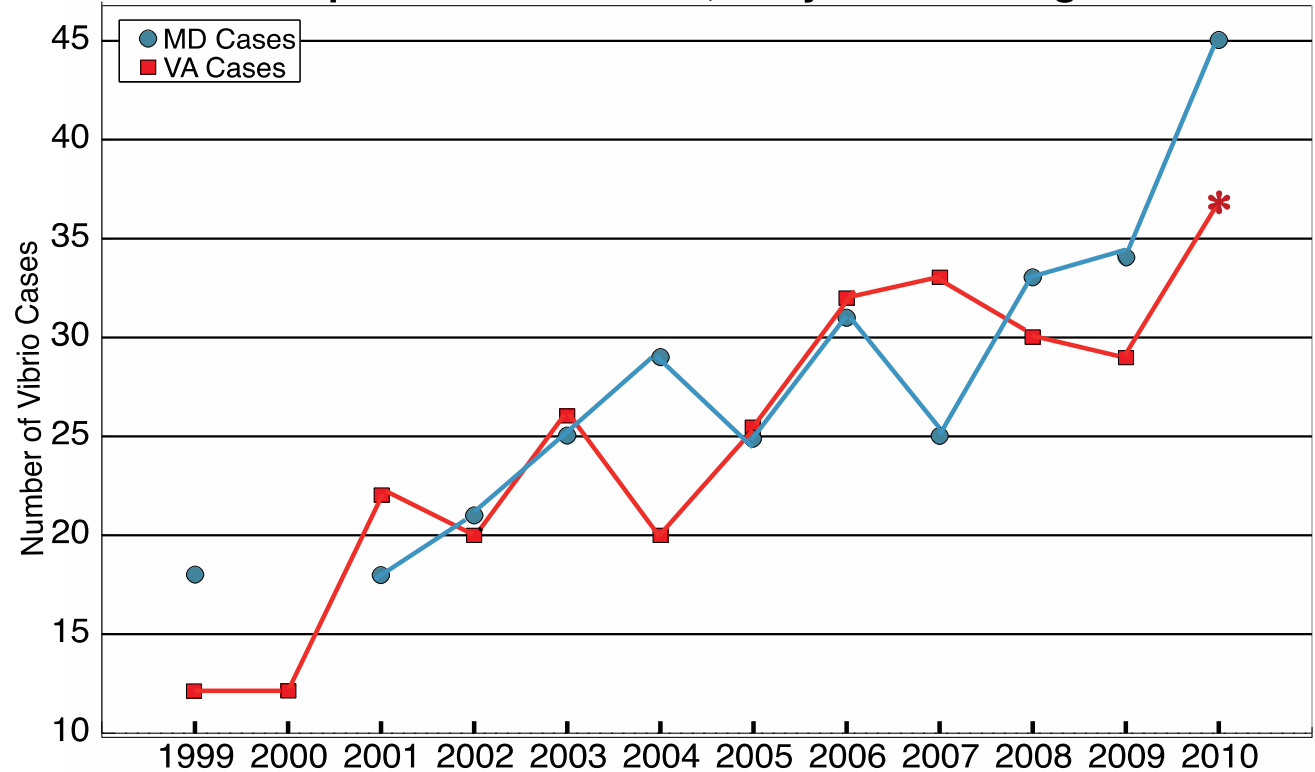
- *V. cholerae*
- *V. vulnificus*
- *V. parahaemolyticus*





Vibrio in the Chesapeake Bay

Reported Vibrio Cases, Maryland and Virginia



Maryland Dept. of Health and Mental Hygiene
Virginia Department of Health.

* Estimated 2010 VA Cases

Pre-existing Empirical Vibrio Models

$$z(V.v) = -7.867 + (0.316 * \text{Temp}) + (-0.342 * (|\text{Saln} - 11.5|))$$

(Jacobs et al., 2010)

$$z(V.c) = -1.1939 + (0.1233 * \text{Temp}) - (0.1997 * \text{Saln}) - (0.0324 * (\text{Temp} * \text{Saln}))$$

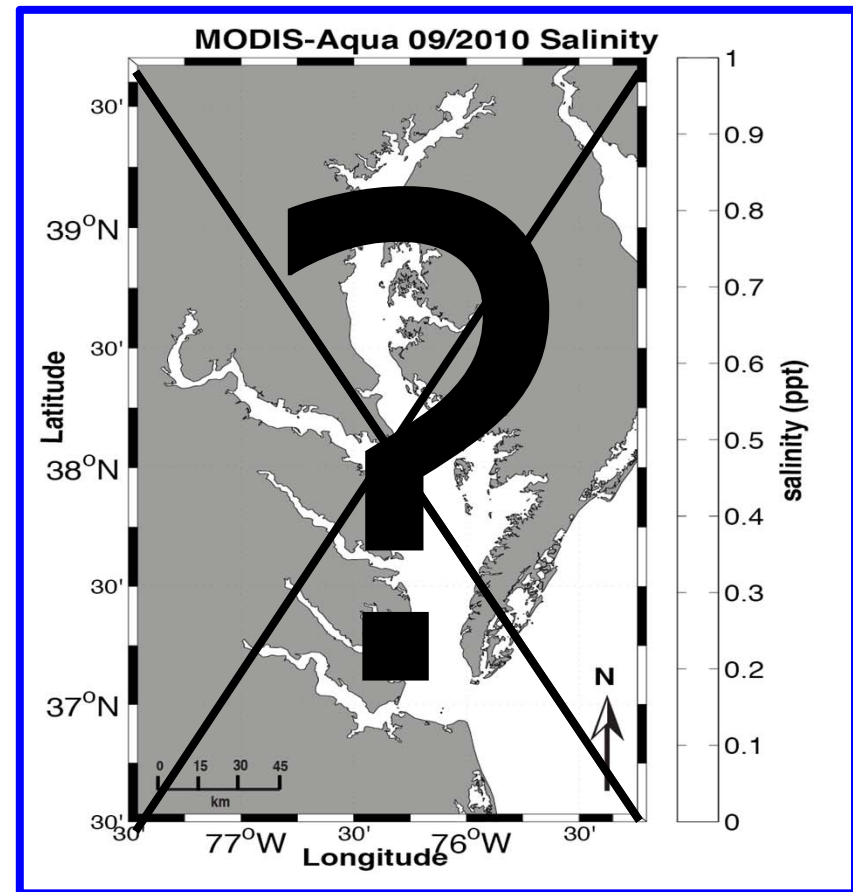
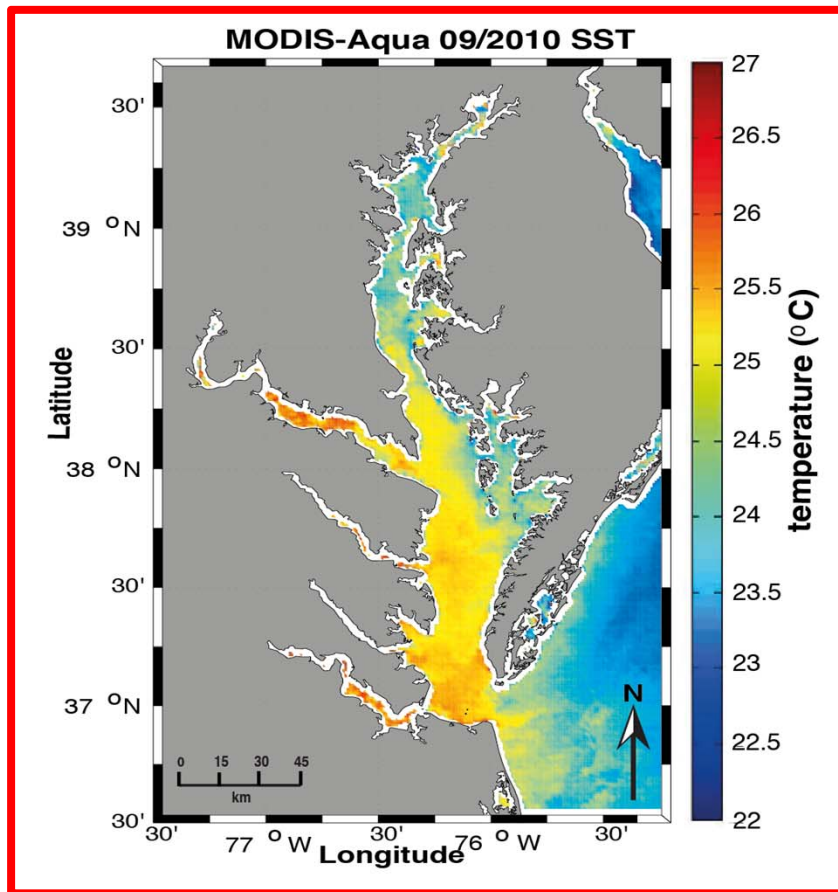
(Louis et al., 2003)

$$f(z) = e^z / (1 + e^z)$$

- **In situ** and **modeled** temperature and salinity inputs
- Probability of occurrence Vibrio spp. models
- Historical cruise (V.c) and CBay Program (V.v) bacteria collection

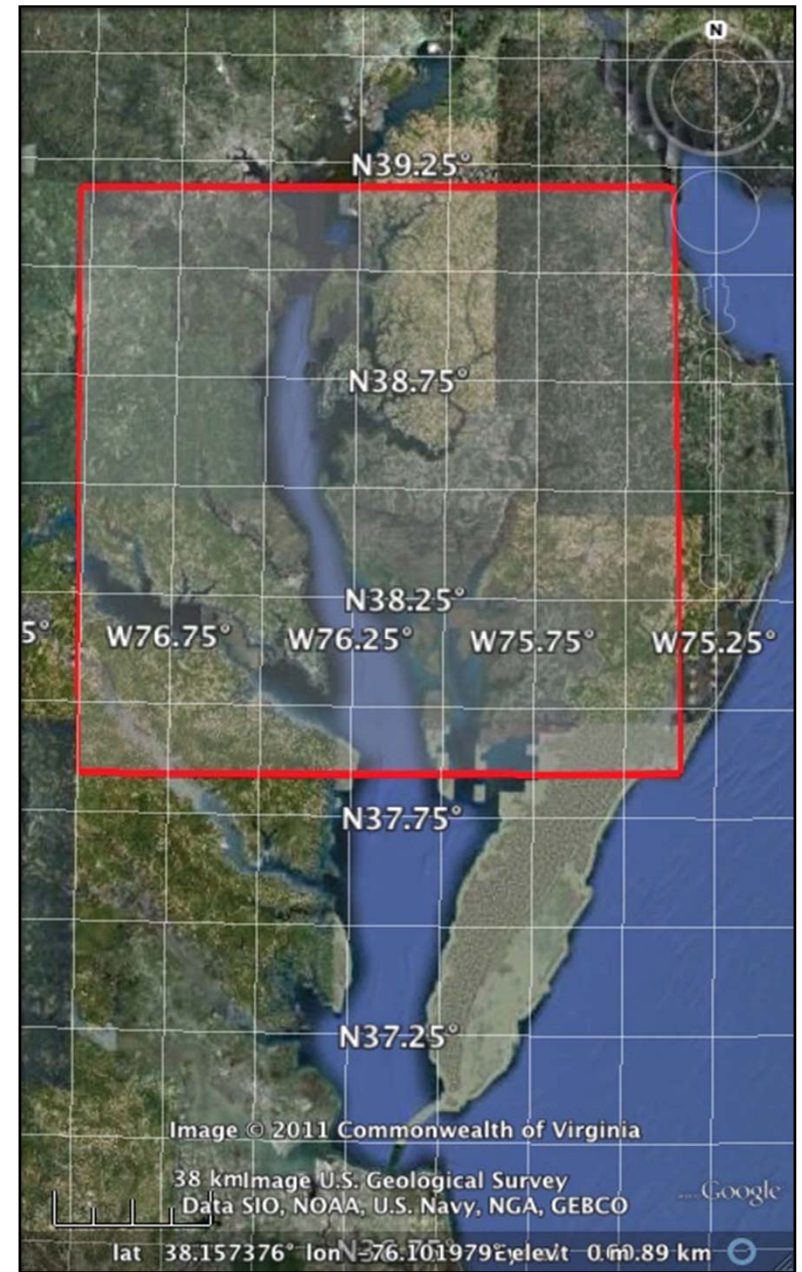
Remote Sensing of Vibrio in the Chesapeake Bay

$$z(\text{V.c.}) = -1.1939 + (0.1233 * \text{Temp}) - (0.1997 * \text{Saln}) - (0.0324 * (\text{Temp} * \text{Saln}))$$



Remote Sensing of Sea Surface Salinity

- **NASA Aquarius Salinity Product**
 - 150 km spatial resolution
 - Monthly composites
- **Neural Network Salinity Product**
 - Geiger et al. (2011): UDEL
 - Statistically Derived from MODIS-Aqua Ocean Color
 - Additional RS input products
 - Trained on Mid-Atlantic region
 - Historical cruise data



Satellite Based Salinity Algorithms

- MODIS-Aqua Ocean Color Standard Products

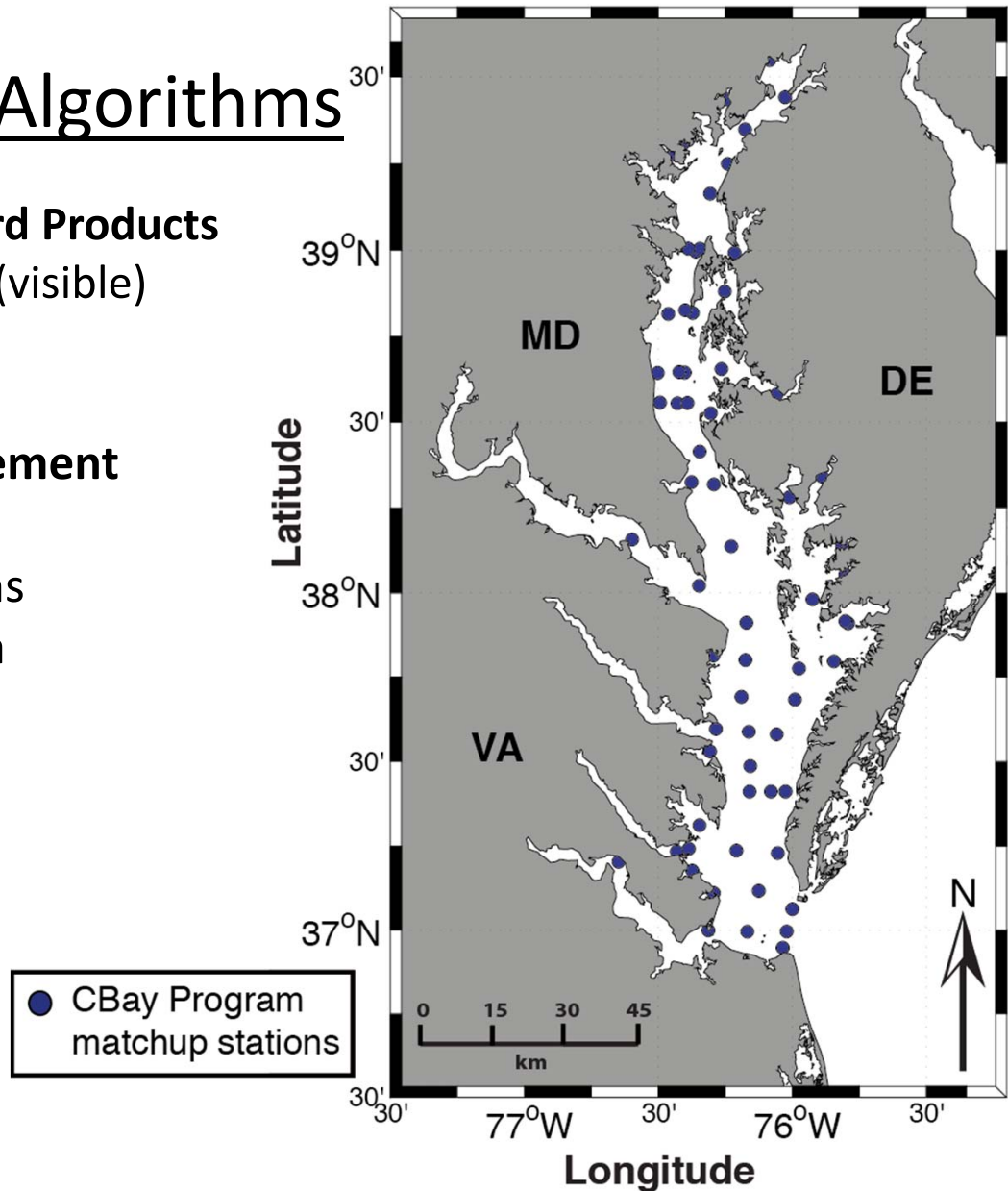
- 10 Remote sensing reflectances (visible)
- 2003-2010

- In situ – remote sensed measurement matchups

- 68 CBay Program in situ stations
- Single pass RS ocean color data
- 1km radius RS averaging
- 2003-2010

- Salinity Prediction Models

- | | |
|---------|--------|
| • GLM | • BART |
| • GAM | • RF |
| • CART | • ANN |
| • BCART | • MARS |



Satellite Based Salinity Algorithms

- Generalized Linear Model (GLM)
- Generalized Additive Model (GAM)
- Artificial Neural Network (ANN)
- Multivariate Adaptive Regression Spline (MARS)

Tree-Based Data Mining

- Categorical and Regression Tree (CART)
- Bagged Categorical and Regression Tree (BCART)
- Bayesian Additive Regression Tree (BART)
- Random Forest (RF)

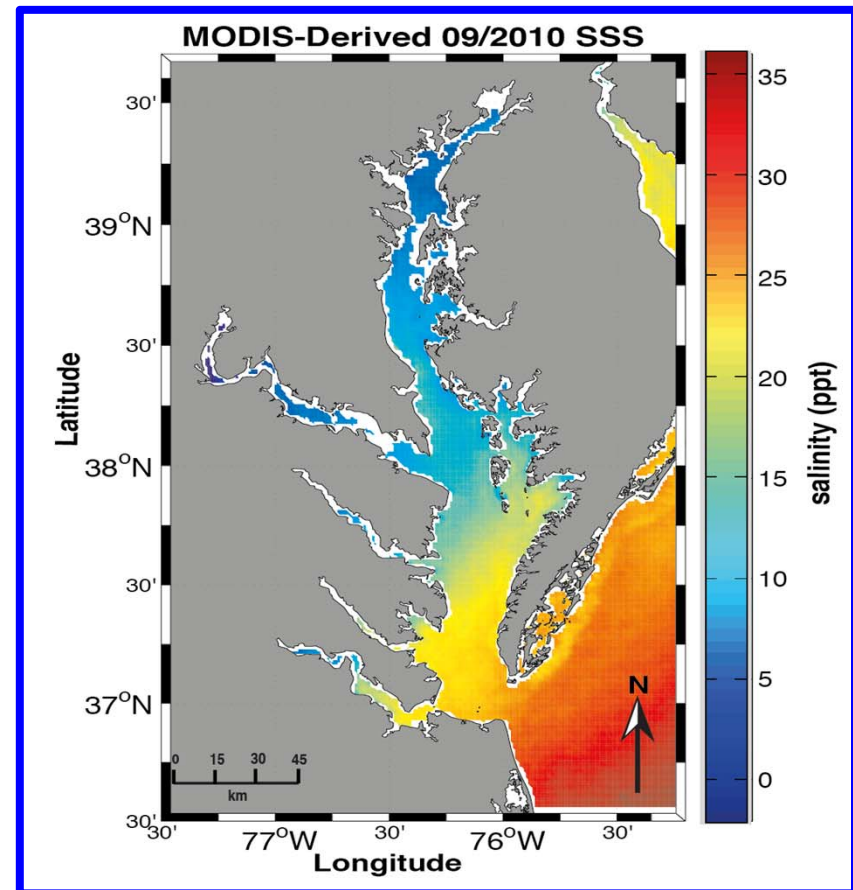
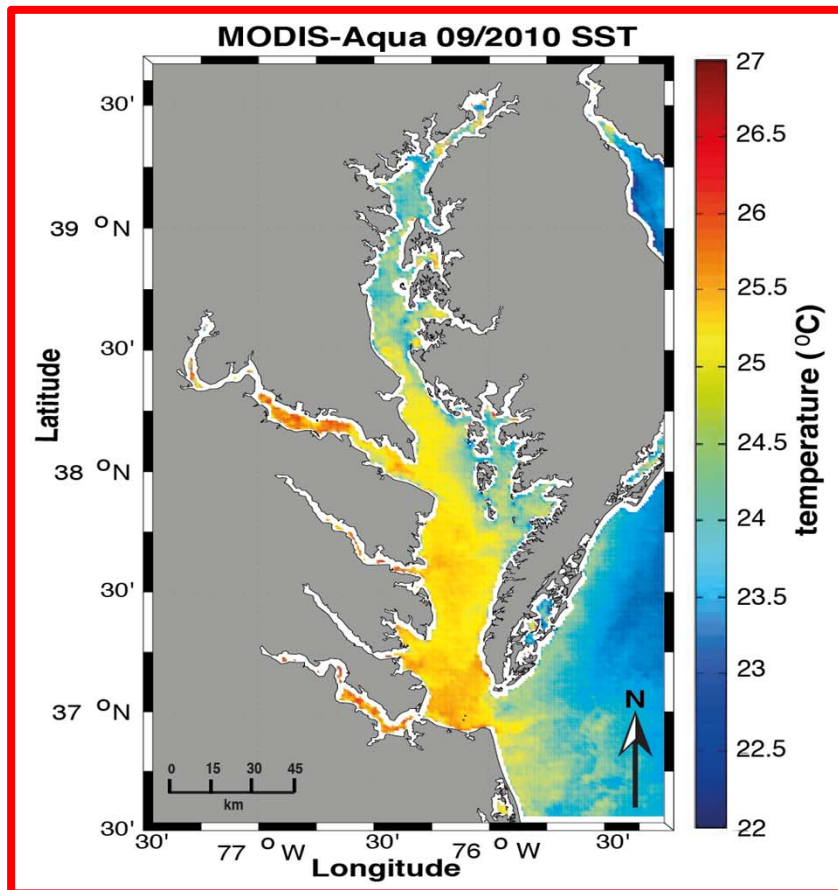
Satellite Based Salinity Algorithms

	GAM	ANN	GLM	CART	BCART	RF	MEAN	BART	MARS
MAE	1.82	1.85	1.93	2.39	2.38	2.06	3.72	2.04	1.98
RMSE	2.38	2.50	2.53	3.03	3.01	2.67	4.69	2.60	2.52
MSE	5.67	6.28	6.40	9.17	9.08	7.14	22.07	6.77	6.33

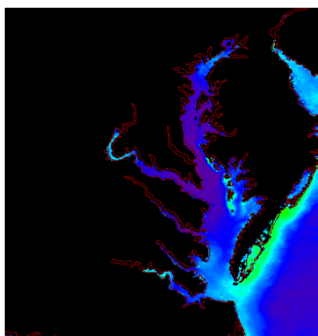
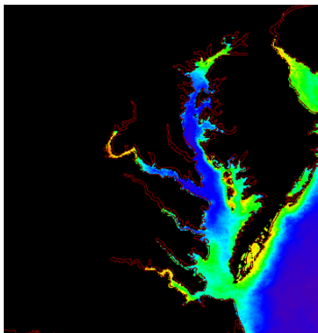
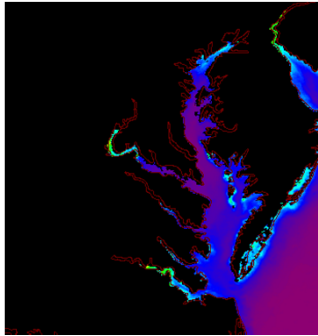
- Top performing prediction models: **GAM** and **ANN**
- GAM and ANN are not statistically different
- All models outperform the mean salinity prediction

Remote Sensing of Vibrio in the Chesapeake Bay

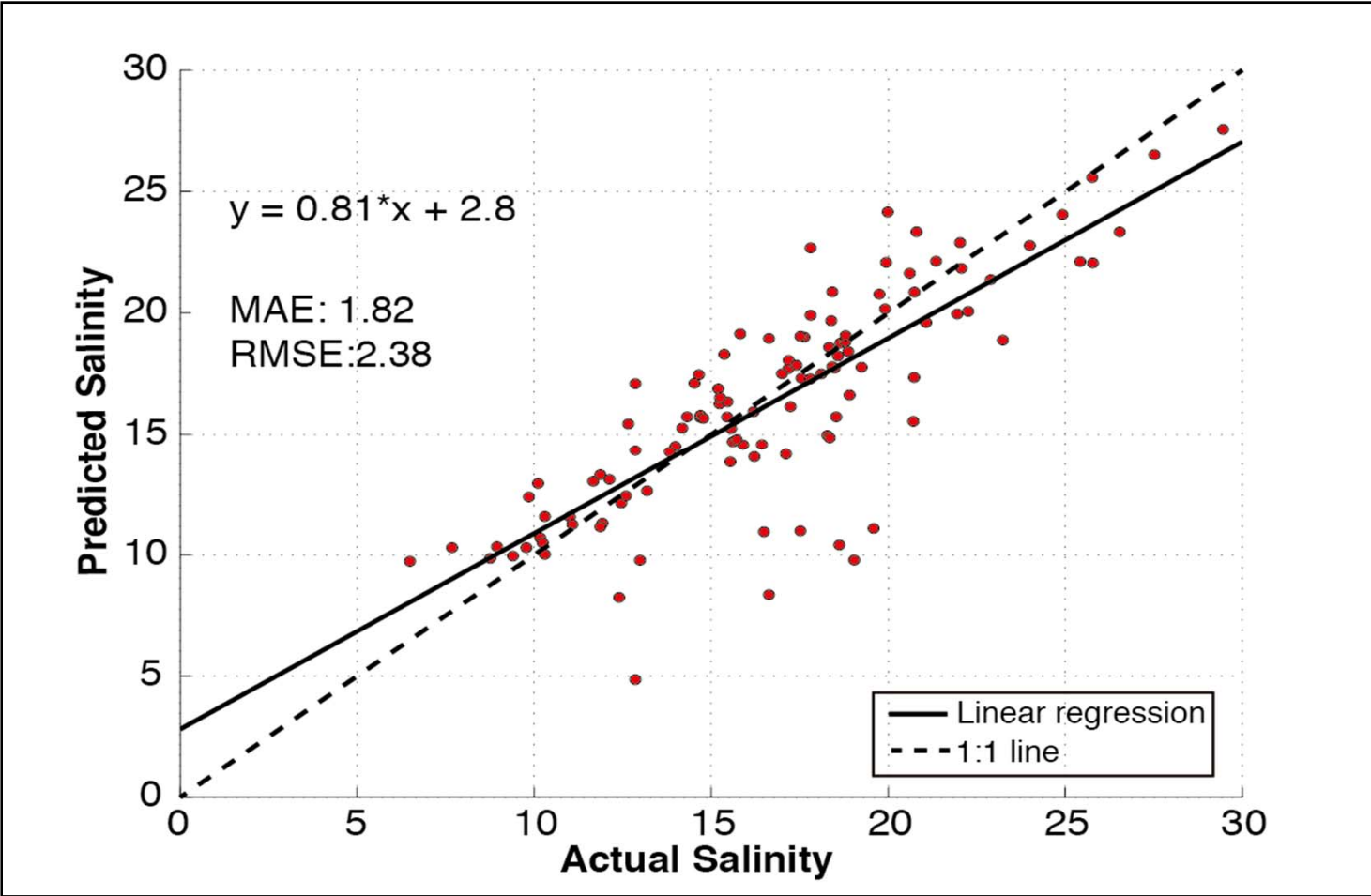
$$z(V.c) = -1.1939 + (0.1233 * \text{Temp}) - (0.1997 * \text{Saln}) - (0.0324 * (\text{Temp} * \text{Saln}))$$



General Additive Model (GAM) for Predicting Salinity

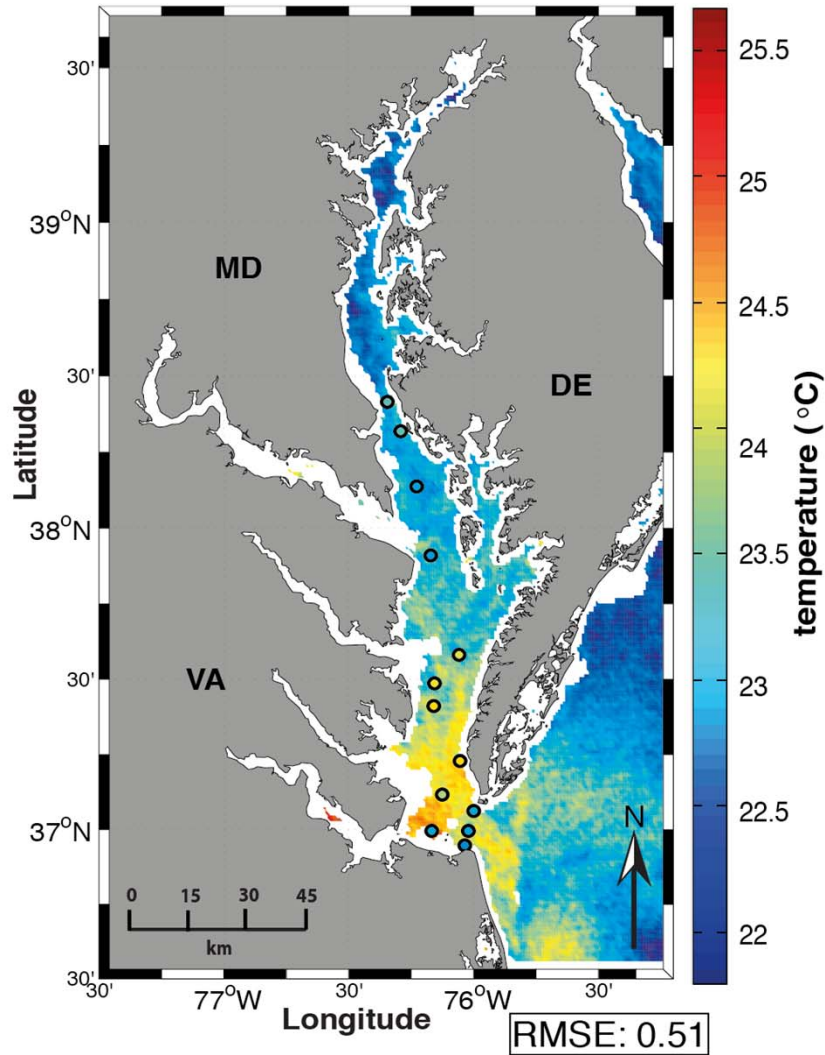


```
GAM.MODEL<-gam(Salinity~s(Rrs_678)+s(Rrs_547)+s(Rrs_488.....))
```

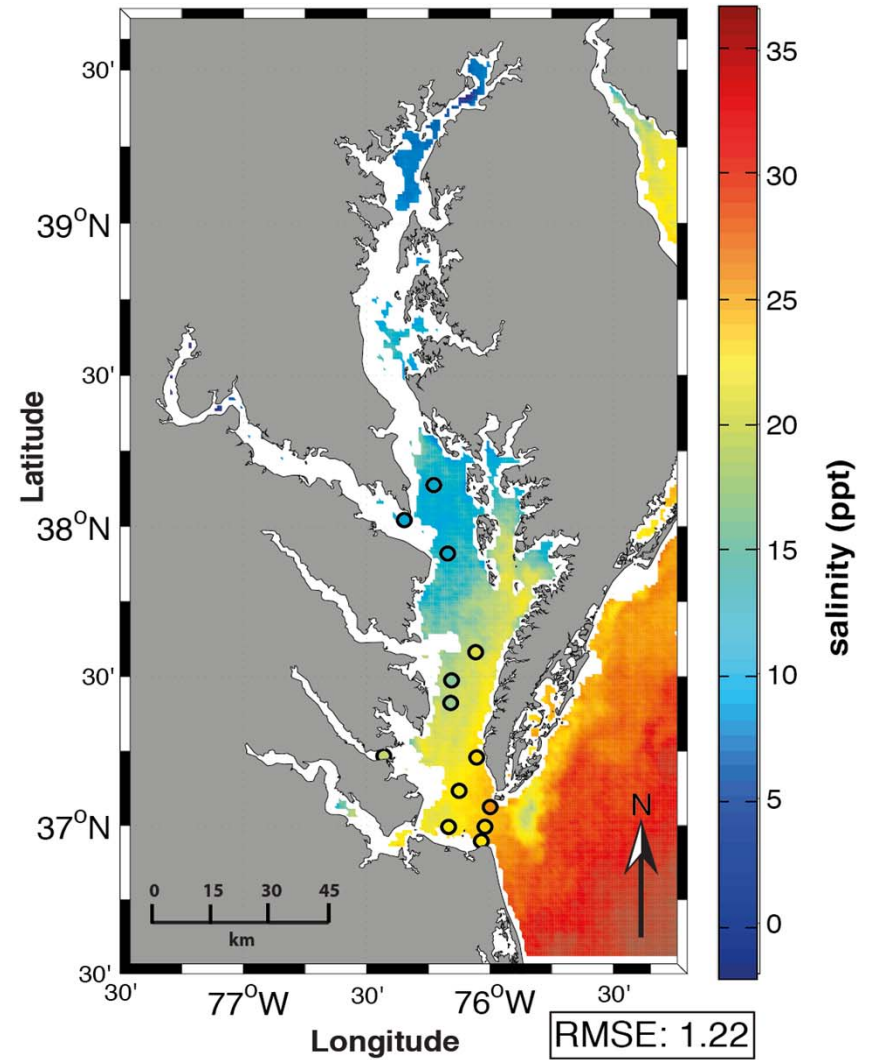


Daily Remote Sensing of SST and Saln

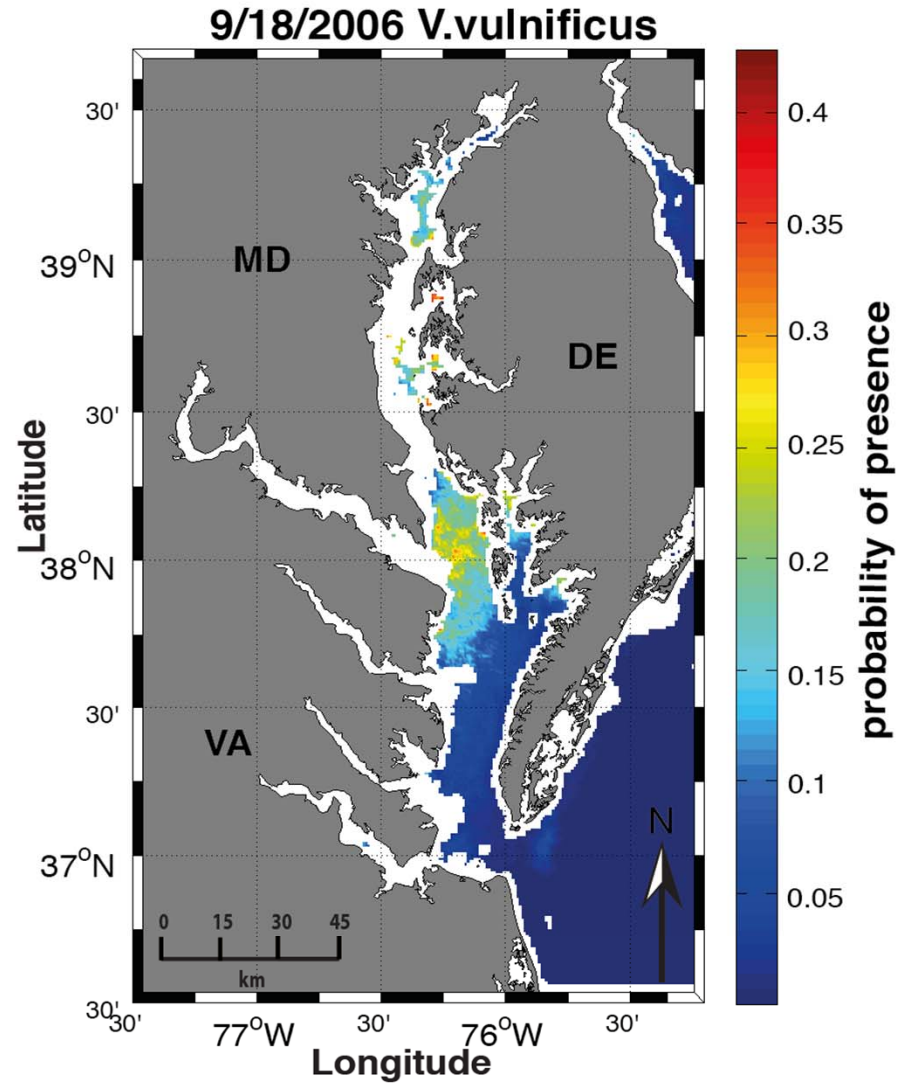
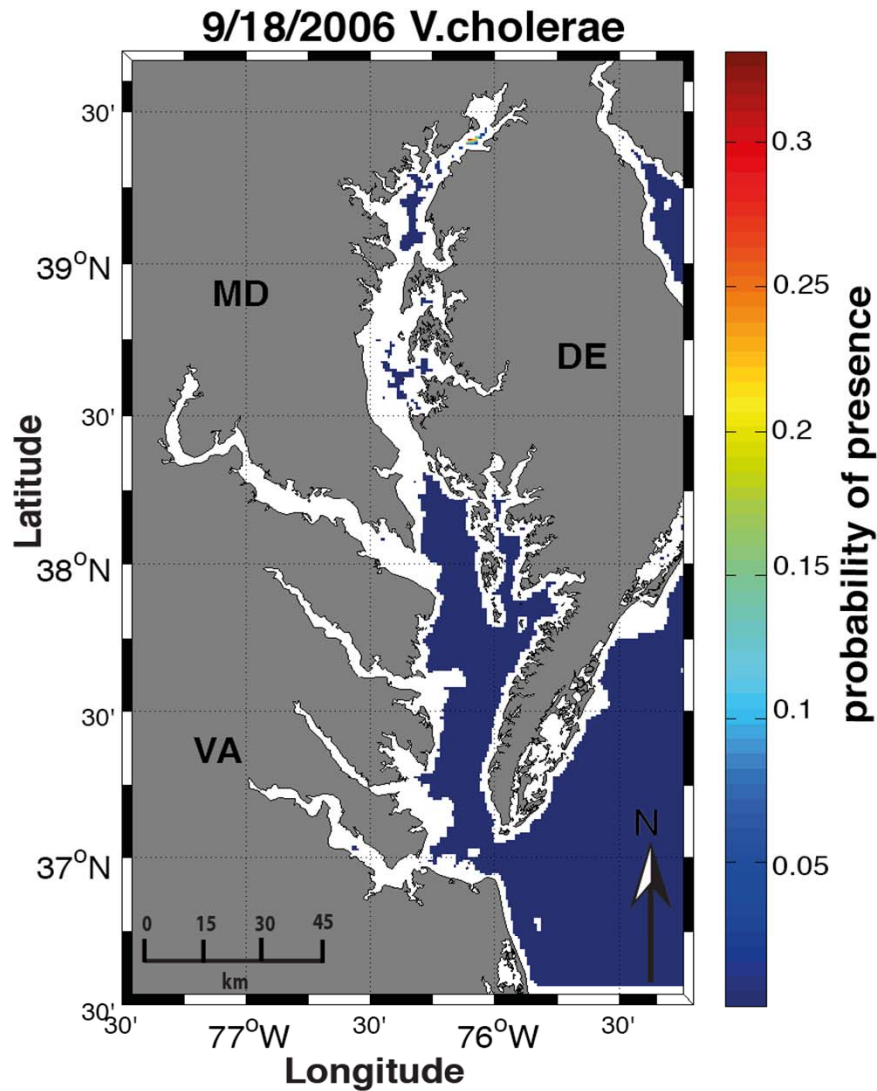
MODIS- Aqua SST 9/18/2006



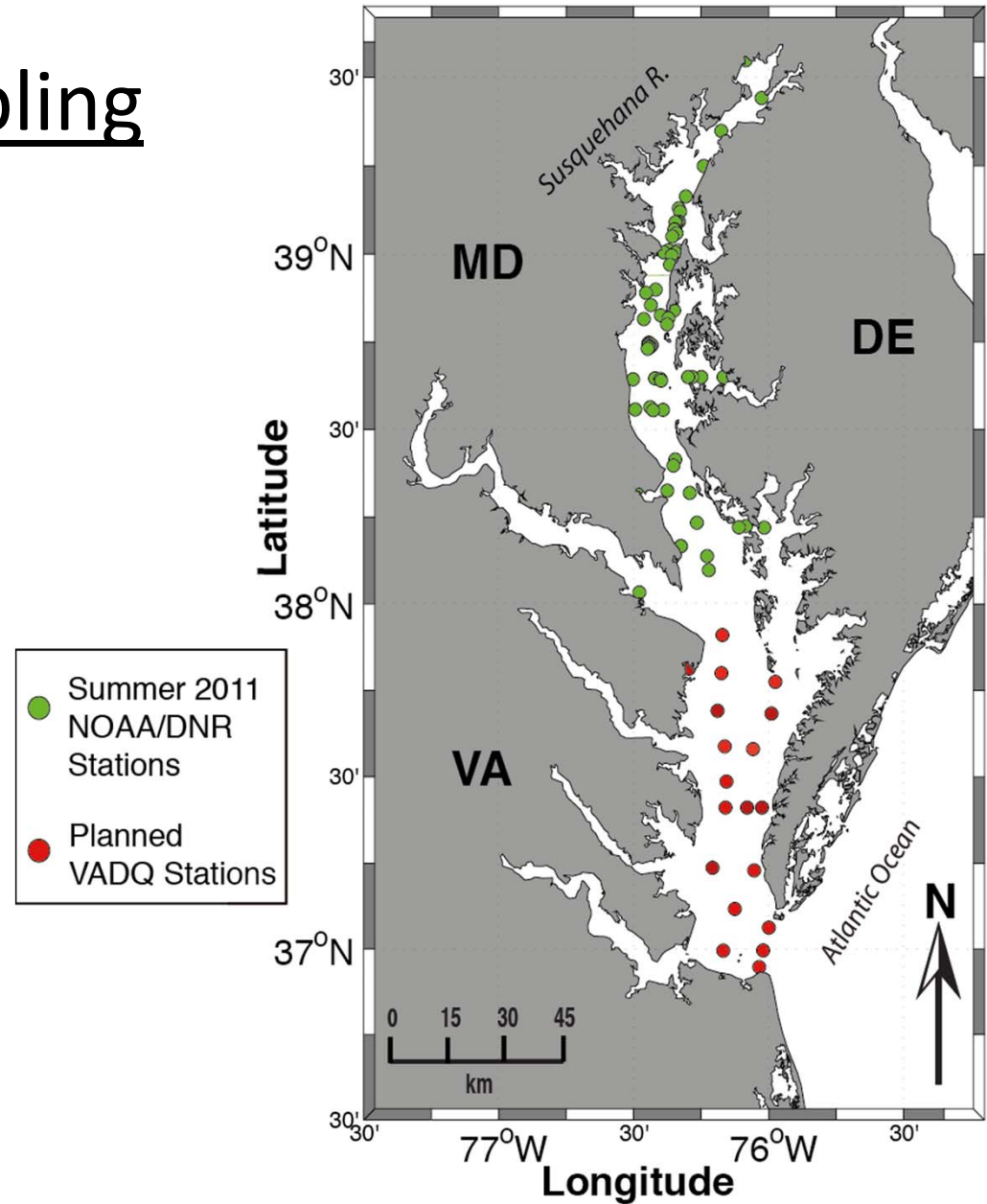
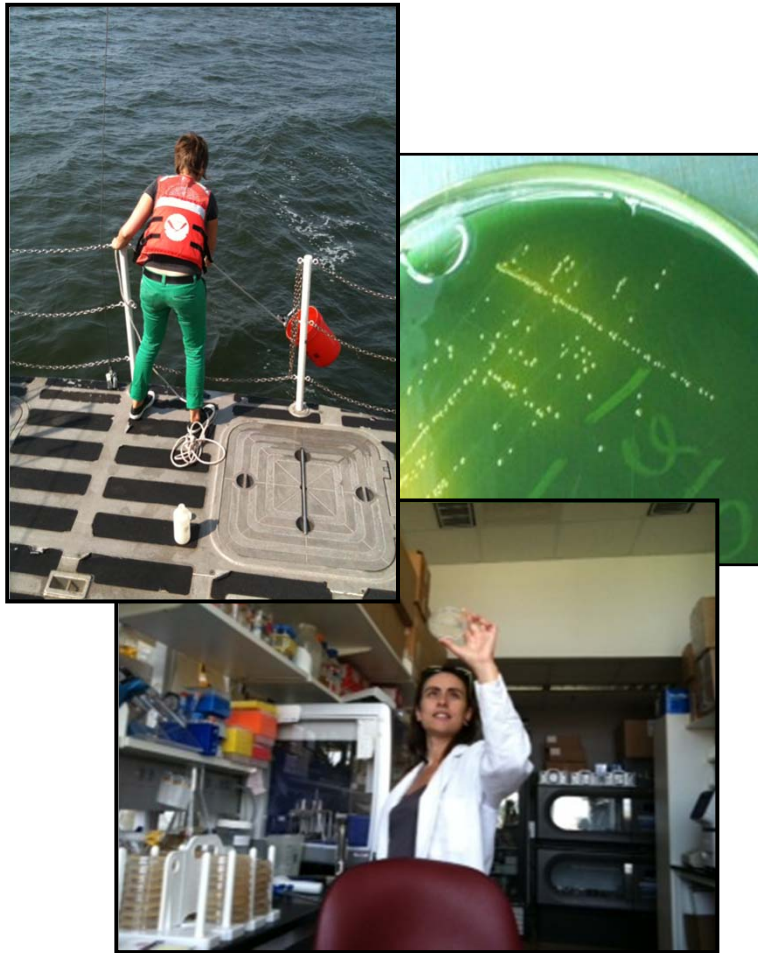
MODIS- Derived SSS 9/18/2006



Remote Sensing of Vibrio in the Chesapeake Bay



In situ Vibrio Sampling



Future Research Directions

- Salinity algorithm applications
- Applications within the Chesapeake Bay
- Applications beyond the Bay

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