



## Assessment of fish habitat in the Chesapeake Bay watershed

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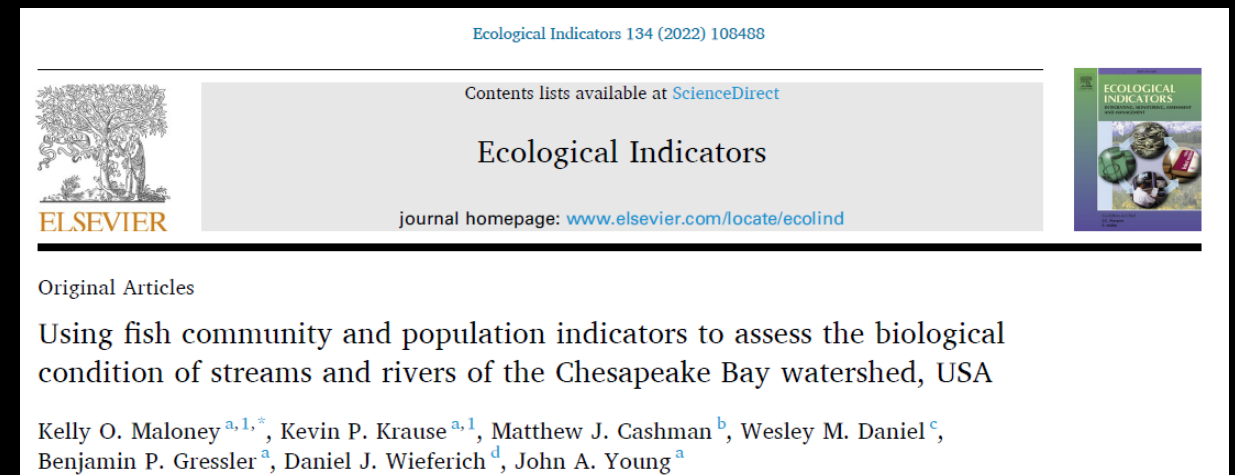
CBP FHAT 06 April 2022



# Objectives

Report on assessment of fish habitat condition for all non-tidal reaches in the Chesapeake Bay watershed

Get feedback and thoughts on web interface tool to present results



<https://doi.org/10.1016/j.ecolind.2021.108488>

- Recognized national efforts (NFHP) were limited
- More relevant information is available for the watershed
- USGS/NOAA inventoried available data and developed concepts
- 2018 Workshop presented ideas and gather feedback



STAC Publication 18-006

- Eventual assessment a fine spatial scale (1:24,000 map scale)
- Data gathering (2018-2019)
- Outreach and Training (2018-present)
- Assessment metrics (2021)
- Pilot assessment (Patuxent 2022-2023)
- Research (communicate research needs)

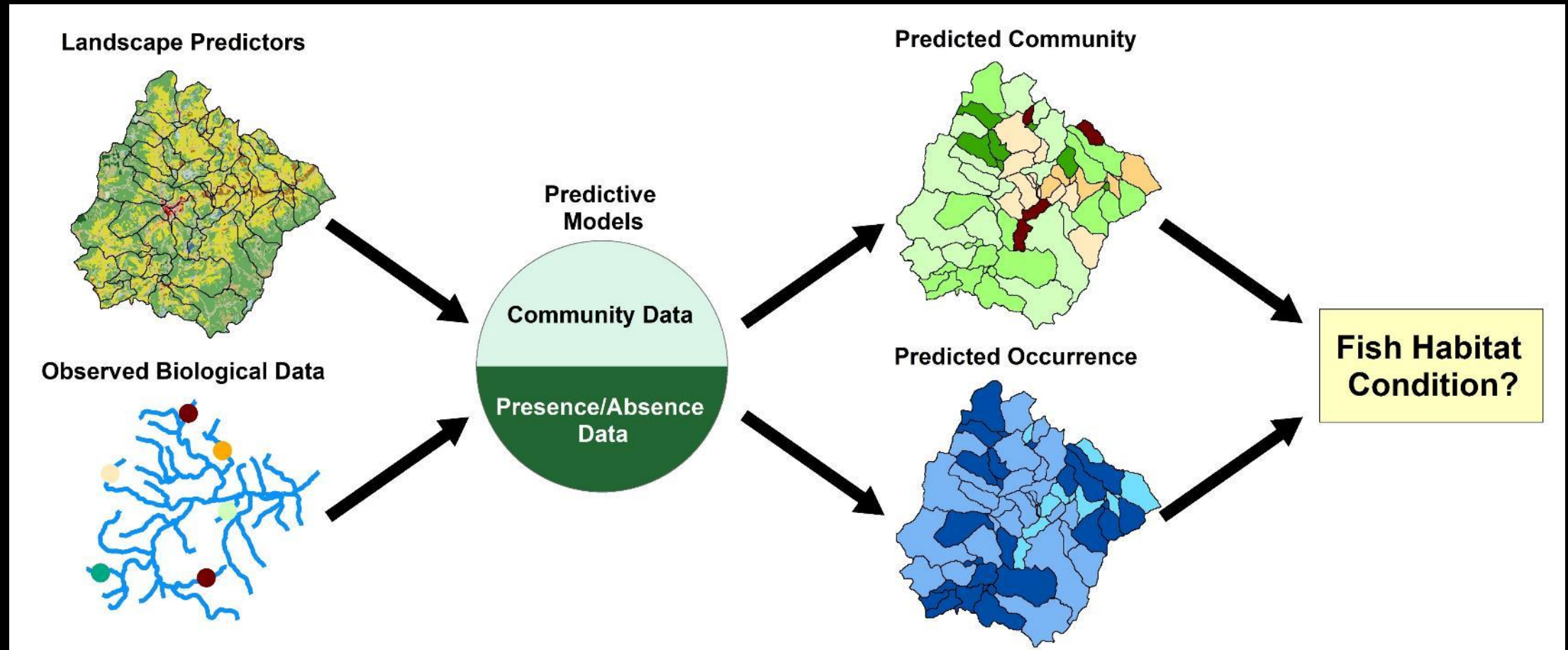
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STAC Publication 18-006

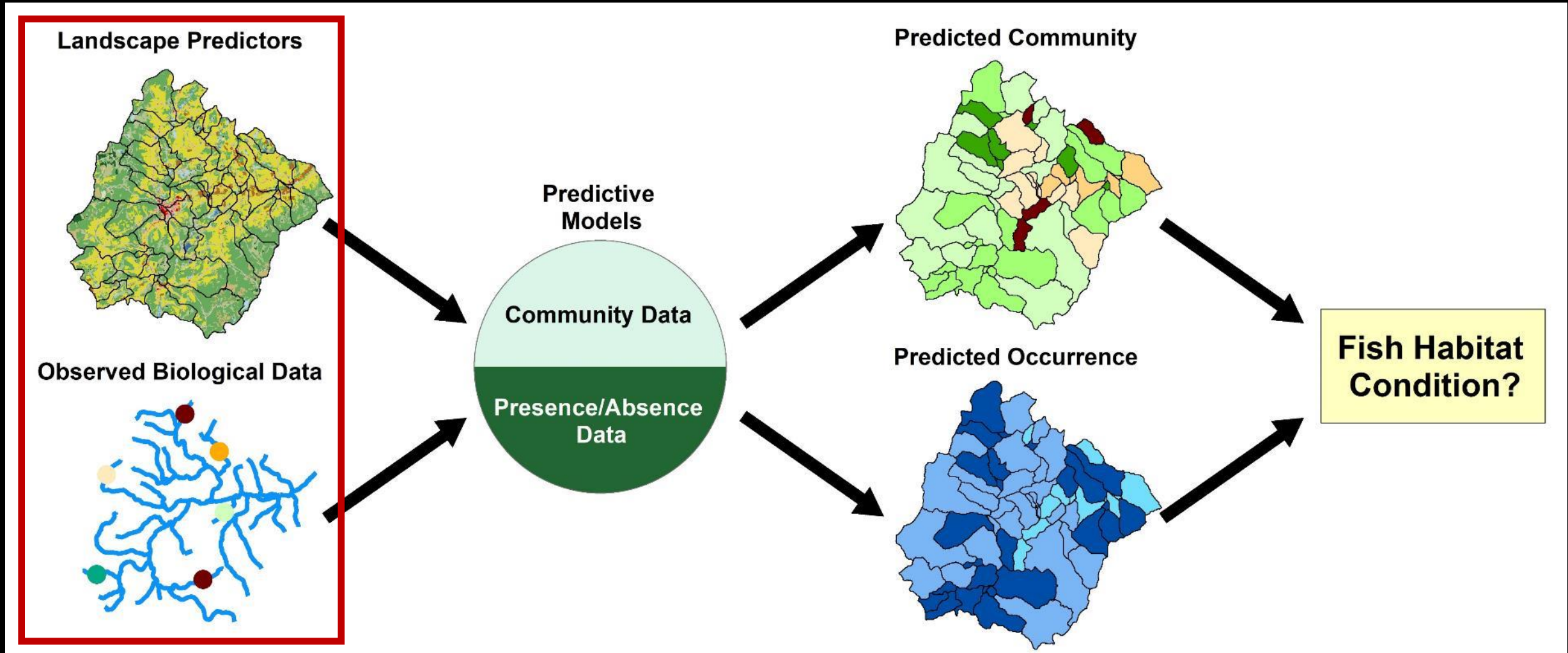
# Non-tidal Assessment: Two main approaches

	Community		Species
Measure	Metrics often rolled into multimetric indices		Key species or set of species
Breadth	Represents community - richness, diversity, abundance and functional aspects		Single or few species
Strength	Useful for overall condition		Detailed insight into key species
Weakness	Limited insight into individual species		Limited evaluation of overall condition

# Non-tidal Assessment: Integrated approach

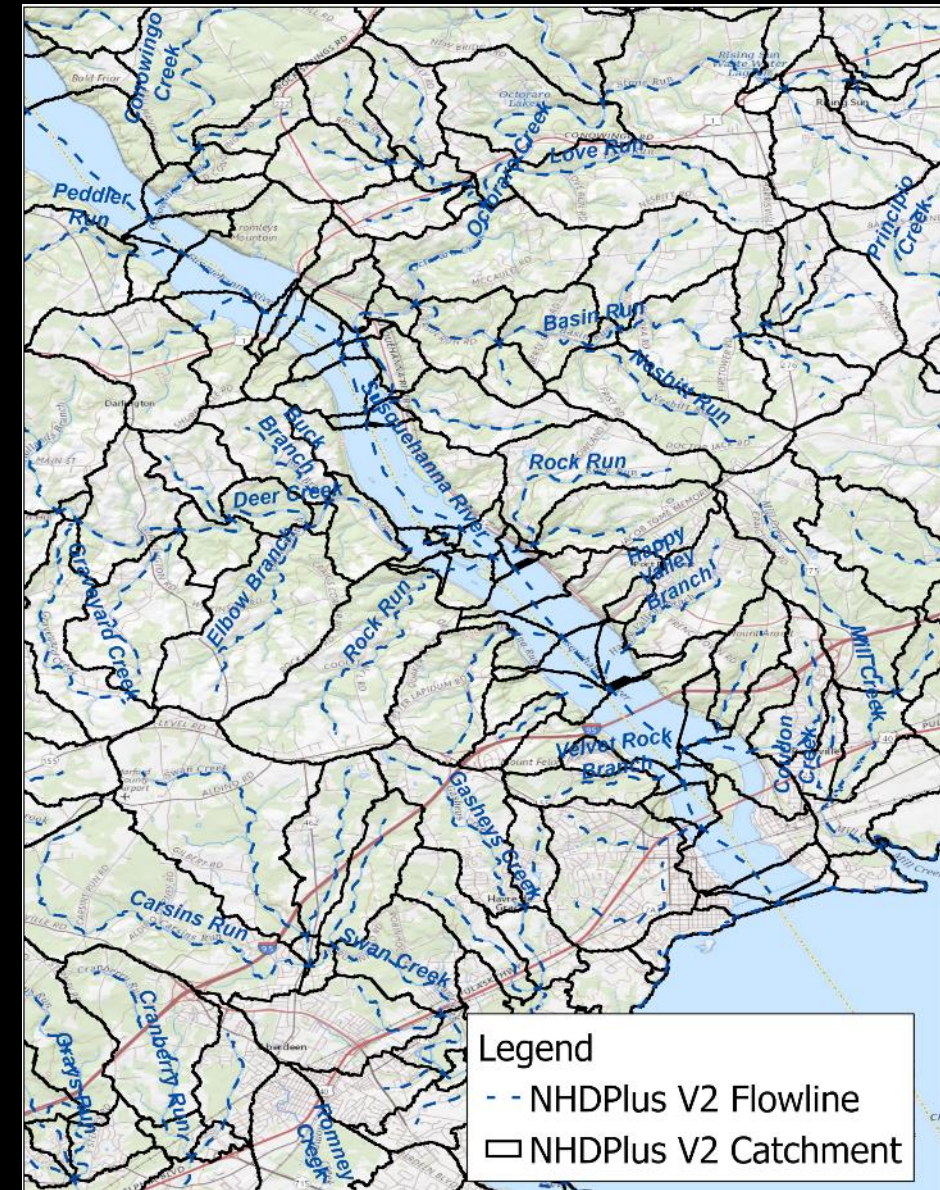






## Data

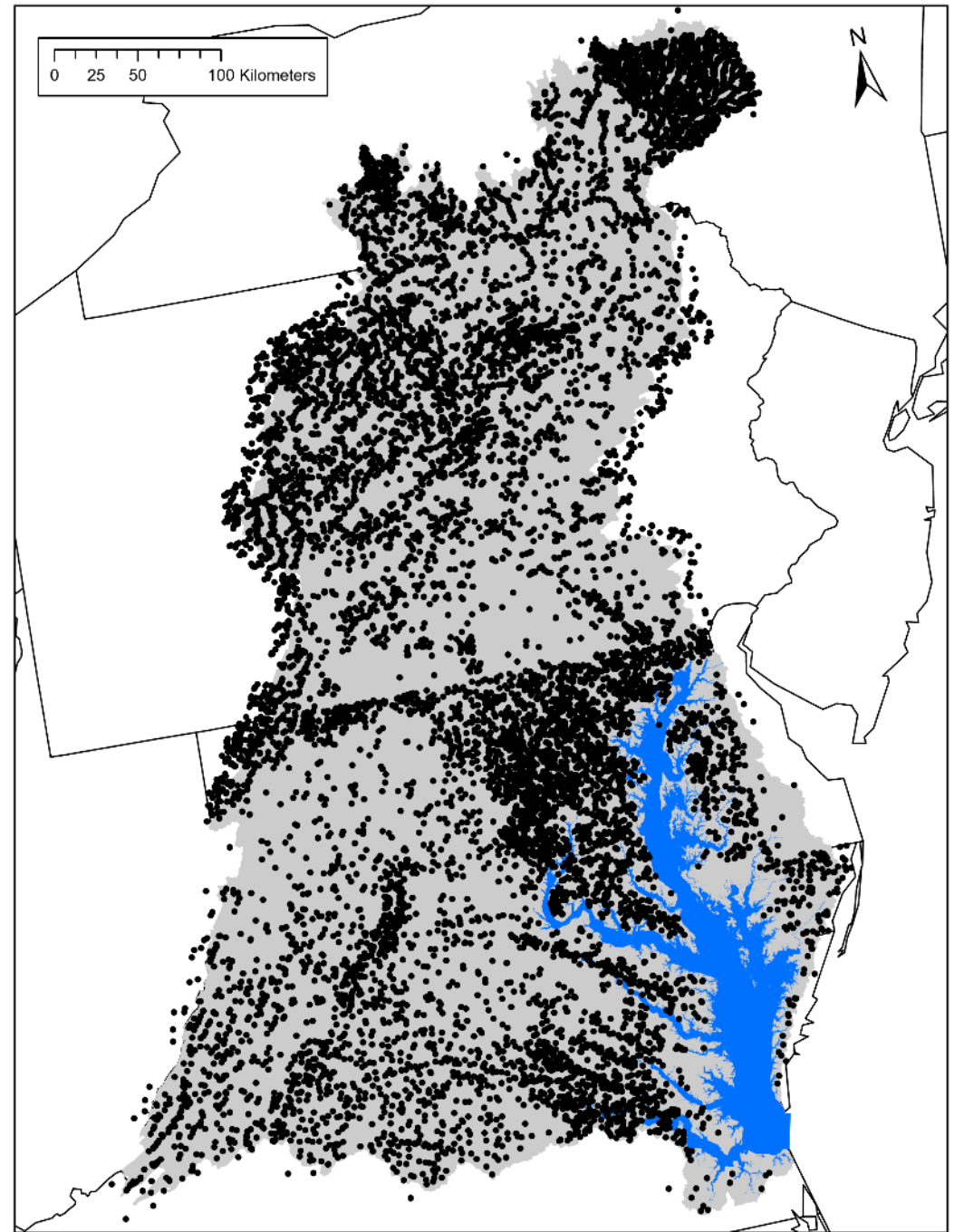
- 56 uncorrelated landscape predictors (upstream accumulated)
  - LULC, climate, topography, geology, soils, human impact
  - From US EPA StreamCat (Hill et al. 2016) and USGS NAWQA (Wieczorek et al. 2018)
- 1:100,000 map scale (NHDPlus v2.1)
  - Next move to 1:24,000 as data become available

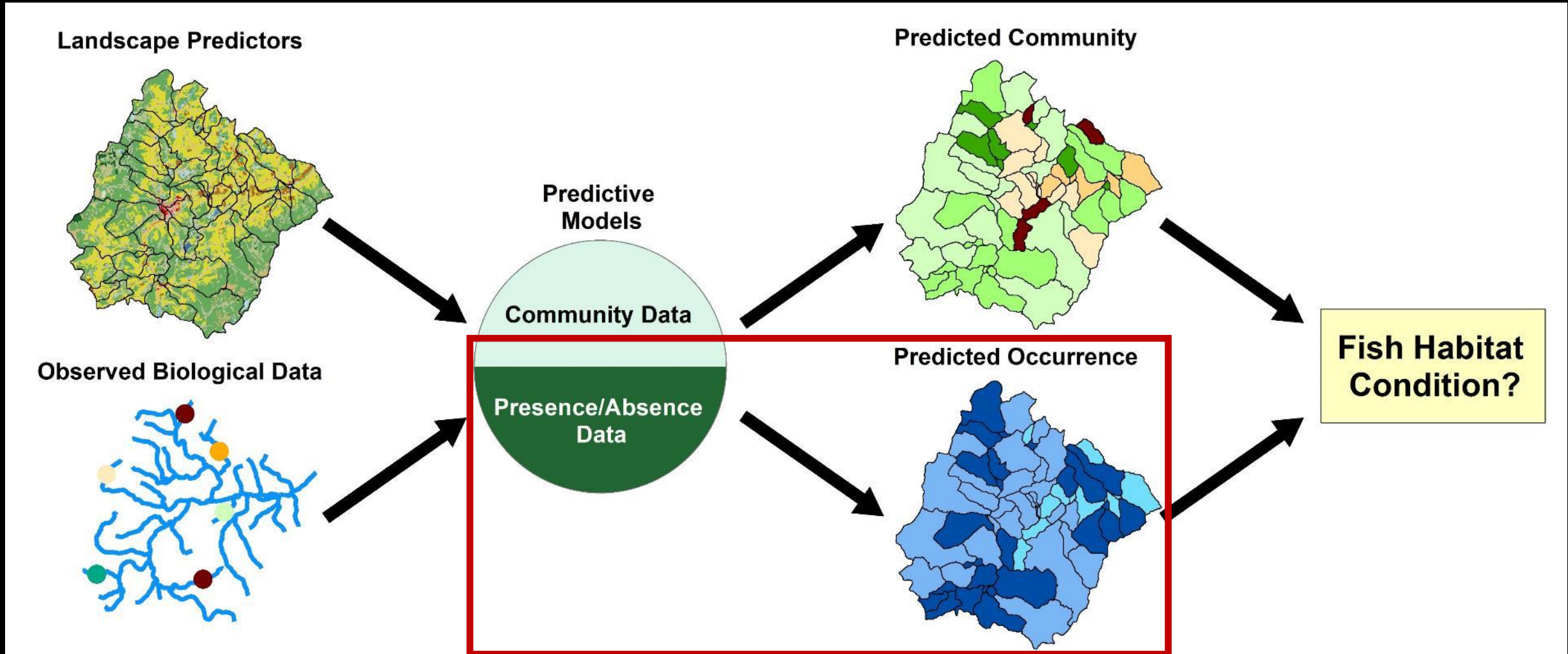




## Non-Tidal Fish Database

- Over 20 separate data sources/sampling programs
- Over 30,000 sampling events
- Summary metrics used by NRSA (EPA, 2020), NFHP (2016), and others of interest calculated for Community samples (Krause and Maloney 2021)
- Spatially linked samples to NHDPlus V2.1 where appropriate (Krause et al. 2021)

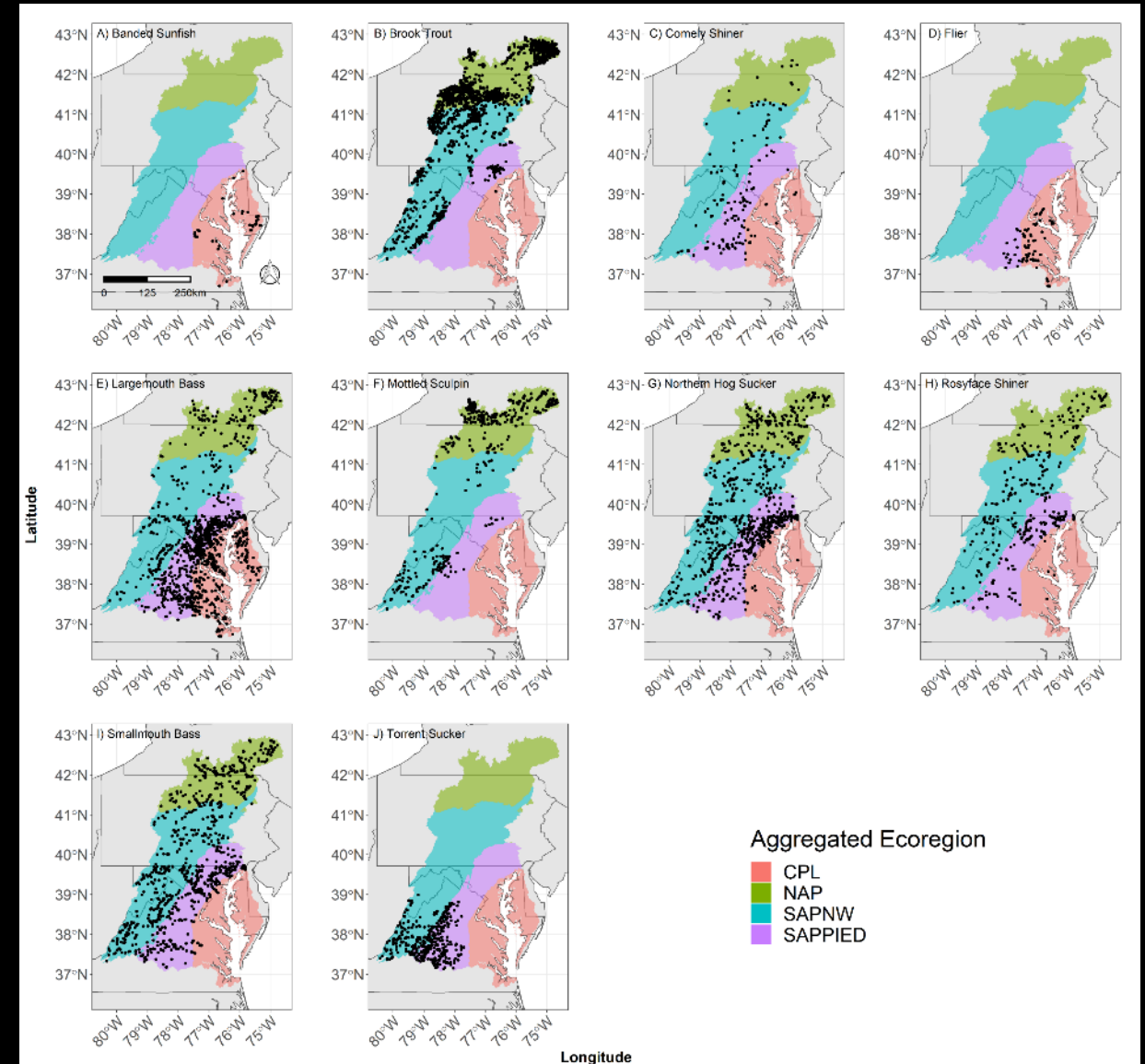






# Methods - Species

- Examined species if:
  - > 100 occurrences
  - USEPA identified as sensitive or were gamefish of interest
- Random forest models built for each species
- Modeling extent set by native range and habitat size
- Models for species selected if:
  - Kappa and TSS > 0.40 (training and test)
  - Specificity and sensitivity > 0.70 (training and test)



# Results – 4 Species met all criteria

**Brook Trout**  
(*Salvelinus fontinalis*)



Kevin P. Krause

Kappa	Sens.	Spec.	TSS
0.85	0.91	0.94	0.86

**Northern Hog Sucker**  
(*Hypentelium nigricans*)



Brian Gratwicke (commons.wikimedia.org) ©

Kappa	Sens.	Spec.	TSS
0.58	0.80	0.81	0.61

**Smallmouth Bass**  
(*Micropterus dolomieu*)



U.S. Fish and Wildlife Service

Kappa	Sens.	Spec.	TSS
0.71	0.74	0.94	0.68

**Torrent Sucker**  
(*Thoburnia rathbuni*)



U.S. Fish and Wildlife Service

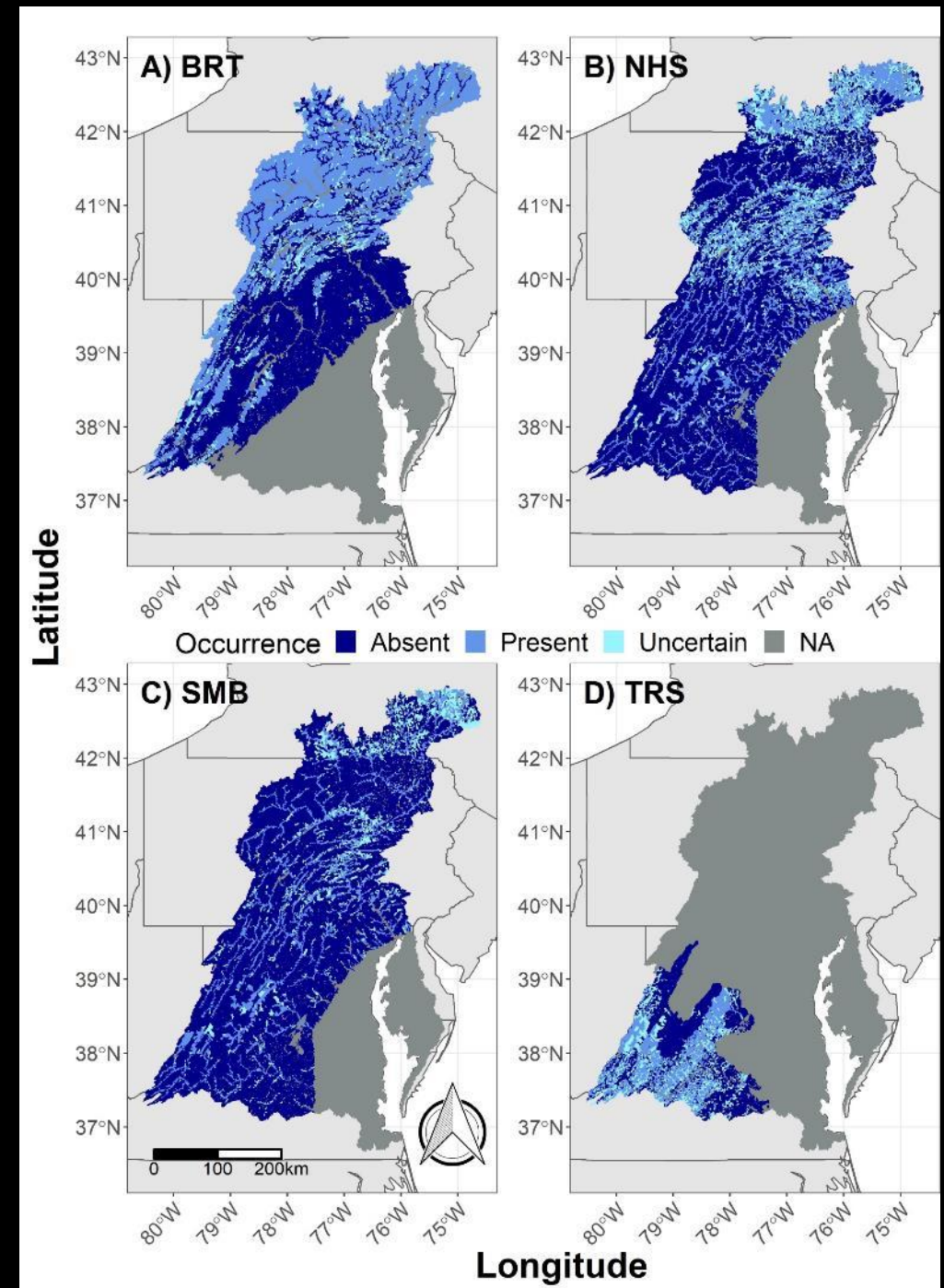
Kappa	Sens.	Spec.	TSS
0.54	0.82	0.72	0.54

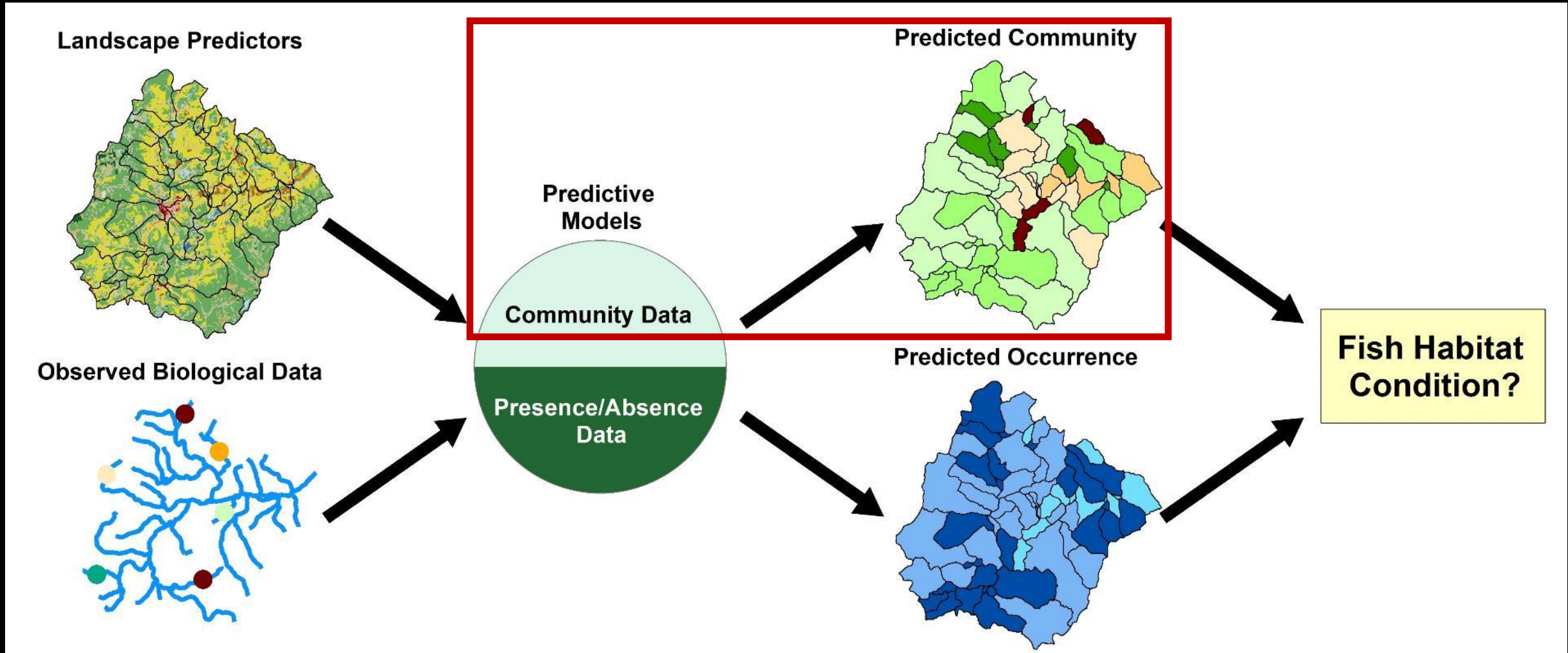
Predicted probability of occurrence (present, absent, uncertain)  
to unsurveyed reaches (2001, 2006, 2011, 2016)



## Results – Species (2016 only)

- Brook Trout (BRT) predicted to smaller systems in the NAP and SAPNW
- Northern Hog Sucker (NHS) predicted for most stream sizes in CBW except CPL
- Smallmouth Bass (SMB) - predicted for small to large rivers in CBW except CPL
- Torrent Sucker (TRS) – predicted for smaller habitats in HUC8 in the southwest portion

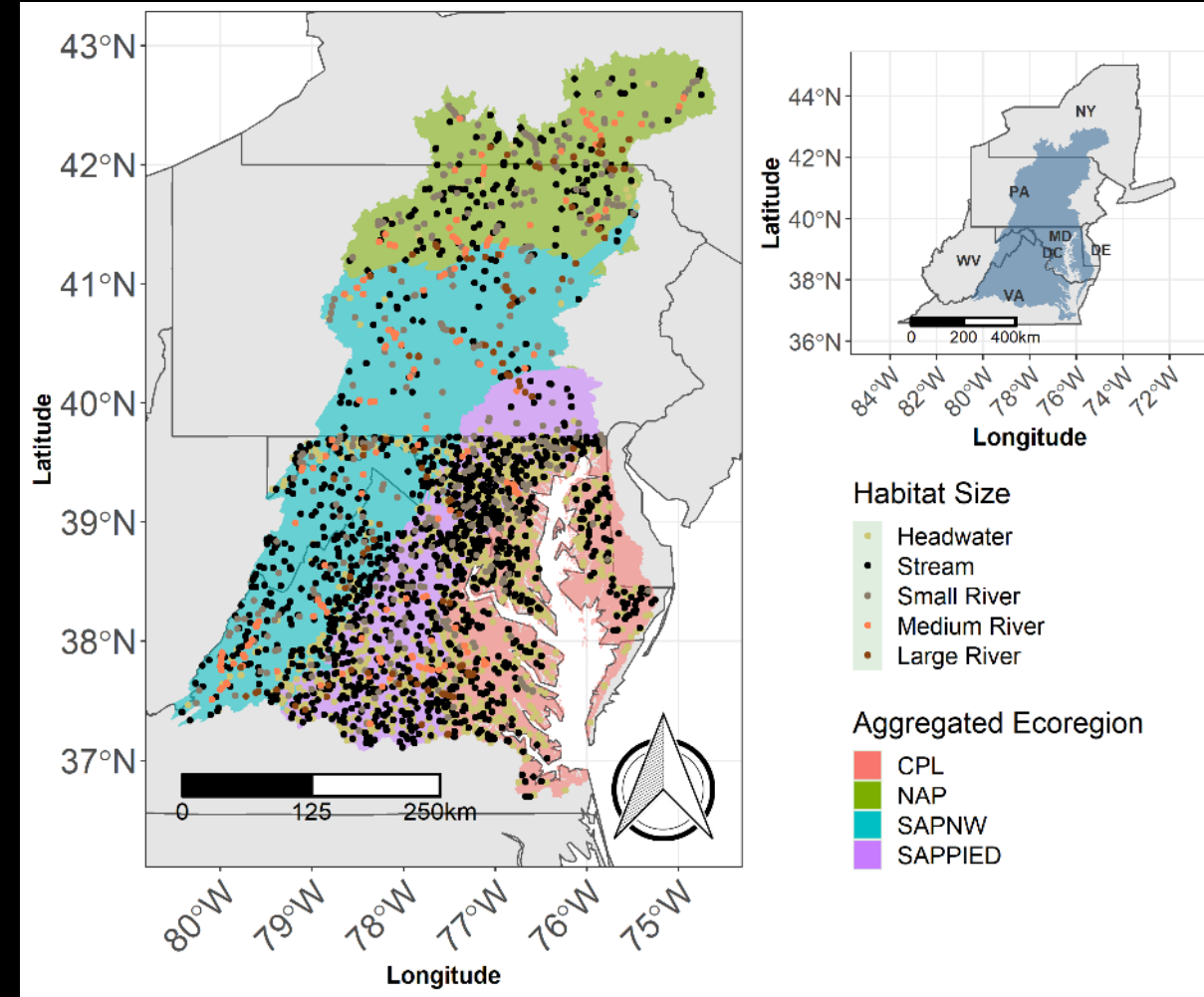






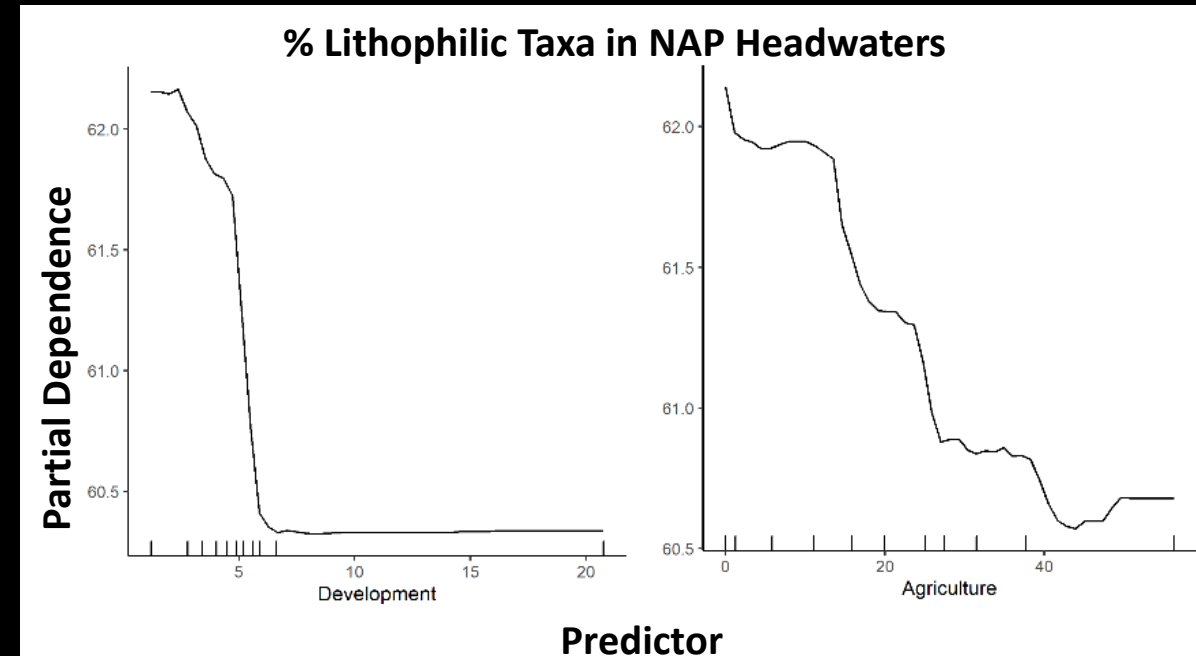
# Methods - Community

- Separate analyses by aggregated ecoregion
- Evaluated > 200 metrics
- Unable to develop MMI
- Random forest models built for metrics with sufficient range and variability (screening criteria used by USEPA 2020)



# Methods - Community

- Separate analyses by aggregated ecoregion
- Evaluated > 200 metrics
- Unable to develop MMI
- Random forest models built for metrics with sufficient range and variability (screening criteria used by USEPA 2020)
- Selected metrics for ecoregion if:
  - Metric was uncorrelated
  - Model had  $R^2 > 0.40$  (training and test)
  - Relationships with development and agriculture were “uni-directional”
- Predicted metrics to unsurveyed reaches (based on 2001, 2006, 2011, 2016 LULC)





# Results - Community

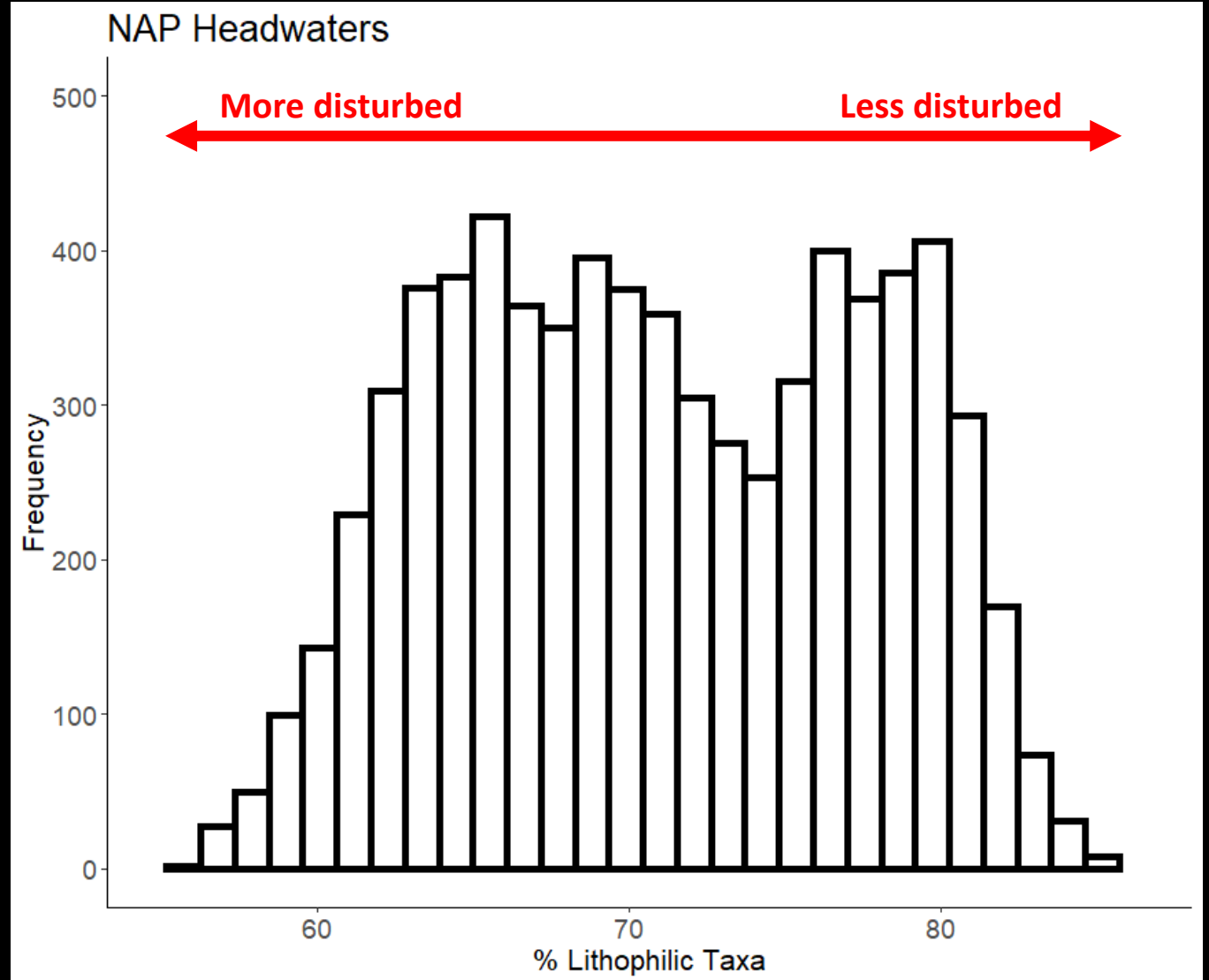
7 metrics in SAPNW, 3 in SAPPIED, 5 in CPL, and 14 in NAP

Region	Metric description	Num. Train n	ind. vars.	R2 (OOB)	Test n	R2 Test	Pattern with Development	Pattern with Agriculture
SAPNW	Number of intolerant lotic taxa	585	47	0.48	147	0.52	↓ to →	↓ to →
	Number of distinct intolerant taxa	585	47	0.48	147	0.52	↓ to →	↓ to →
	Number of native Cyprinidae taxa	585	47	0.49	147	0.53	↑ to →	↑
	% native taxa as intolerant lotic	585	47	0.41	147	0.42	↓ to →	↓ to →
	% Intolerant native taxa	585	47	0.41	147	0.42	↓ to →	↓ to →
	% native taxa as intolerant rheophilic	585	47	0.45	147	0.43	↓ to →	↓ to →
	Number of distinct Tolerant taxa	585	47	0.62	147	0.61	↑	↑
SAPPIED	% large river taxa	1108	51	0.52	278	0.54	rapid ↑ to → to ↓ at extremes	rapid ↑ then slow ↓
	Number of native rheophilic taxa	1108	51	0.52	278	0.53	↑	↑
	Number of distinct Tolerant native taxa	1108	51	0.49	278	0.49	rapid ↓ to ↑	↑
CPL	Number of lithophilic taxa	687	49	0.53	172	0.60	↑	↑
	Number of lotic taxa	687	49	0.46	172	0.56	↑	↑
	Number of native benthic invertivore taxa	687	49	0.43	172	0.41	↑	↑
	Number of native Cyprinidae taxa	687	49	0.53	172	0.63	↑	rapid ↓ then ↑
	Number of distinct Tolerant taxa	687	49	0.41	172	0.51	↑	↑

Region	Metric description	Num. Train n	ind. vars.	R2 (OOB)	Test n	R2 Test	Pattern with Development	Pattern with Agriculture
NAP	Number of Cyprinidae taxa	252	44	0.43	63	0.56	↑	↑
	% lithophilic taxa	252	44	0.46	63	0.53	↓	↓
	% large river taxa	252	44	0.73	63	0.74	↑	↑
	Number of native benthic invertivore taxa	252	44	0.51	63	0.58	↑	↑
	% native individuals as Centrarchidae	252	44	0.55	63	0.47	↑	↑
	% native individuals as coldwater	252	44	0.72	63	0.56	↓	↓ to → to ↑ at extremes
	% native individuals as intolerant rheophilic	252	44	0.47	63	0.43	↓	↓
	Number of native invertivores taxa	252	44	0.44	63	0.57	↑	↑
	Number of native lotic taxa	252	44	0.50	63	0.57	↑	↑
	Number of native non-tolerant benthic taxa	252	44	0.48	63	0.64	↑	↑
	% native individuals as rheophilic	252	44	0.45	63	0.56	↓	↓ to slight ↑ to ↓
	% native taxa as rheophilic	252	44	0.57	63	0.60	↓	↓
	% native taxa as salmonid	252	44	0.65	63	0.81	↓ to → to light ↑	↓ to → to ↑ at extremes
	Number of distinct Tolerant taxa	252	44	0.48	63	0.53	↑	↑

# Methods - Community

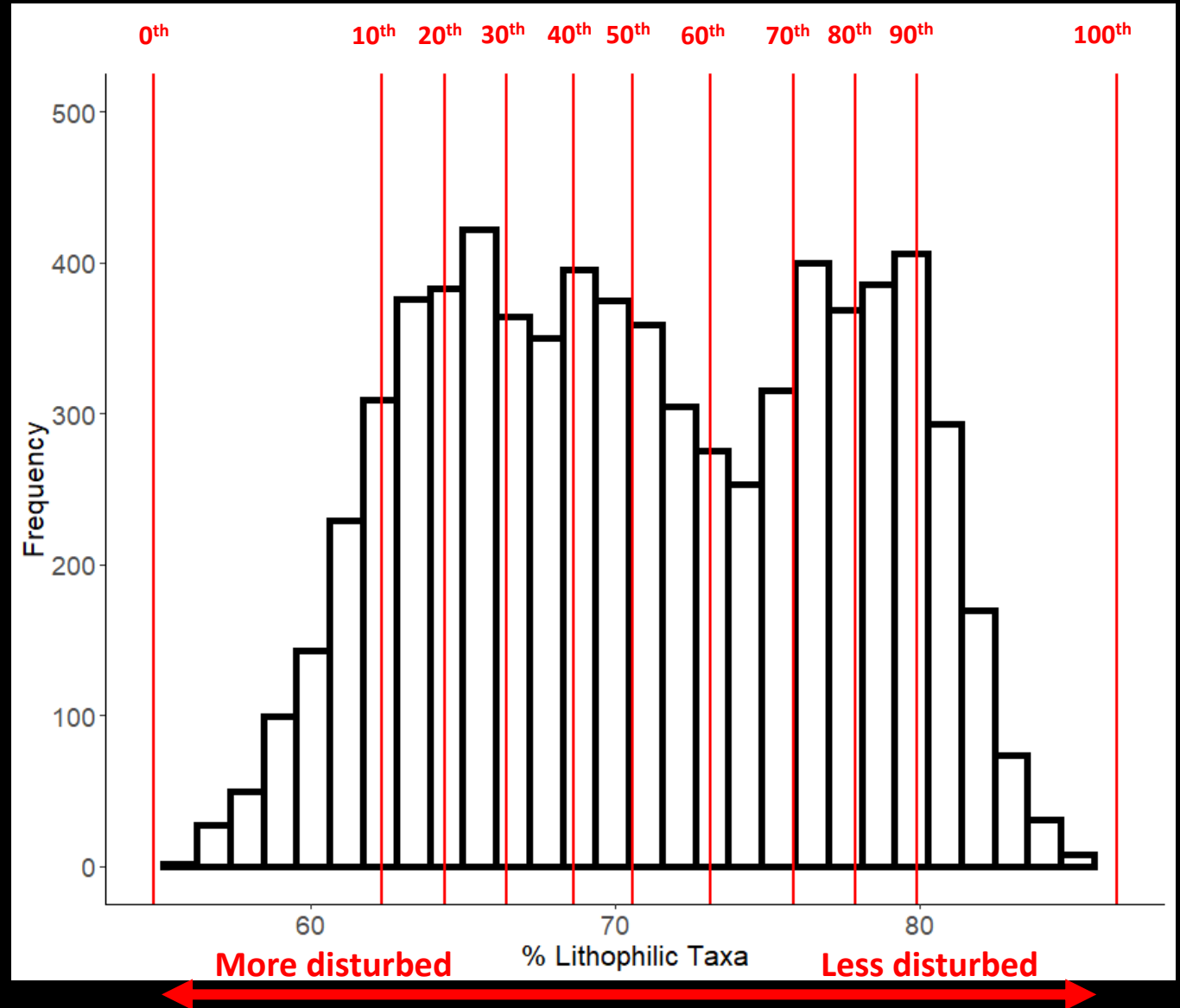
- Unable to develop MMI
- Developed different method:
  - Metric relationship with development and agriculture
  - Binned each metric into deciles (within habitat size class)
    - <10<sup>th</sup> decile = lowest metric values
    - >90<sup>th</sup> decile = highest metric values





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  - Binned each metric into deciles (within habitat size class)
    - <10<sup>th</sup> decile = lowest metric values
    - >90<sup>th</sup> decile = highest metric values
  - Calculate mean of decile scores across suitable metrics
    - Used as condition indicator

Lower mean deciles = poorer relative condition

Higher mean deciles = better relative condition

Reach mean metric decile = 40<sup>th</sup>-50<sup>th</sup>

Region	Metric description	Decile	Decile position
NAP	Number of Cyprinidae taxa	20 <sup>th</sup> – 30 <sup>th</sup>	3
	% lithophilic taxa	40 <sup>th</sup> – 50 <sup>th</sup>	5
	% large river taxa	20 <sup>th</sup> – 30 <sup>th</sup>	3
	Number of native benthic invertivore taxa	20 <sup>th</sup> – 30 <sup>th</sup>	3
	% native individuals as Centrarchidae	60 <sup>th</sup> – 70 <sup>th</sup>	7
	% native individuals as coldwater	70 <sup>th</sup> – 80 <sup>th</sup>	8
	% native individuals as intolerant rheophilic	40 <sup>th</sup> – 50 <sup>th</sup>	5
	Number of native invertivores taxa	50 <sup>th</sup> – 60 <sup>th</sup>	6
	Number of native lotic taxa	50 <sup>th</sup> – 60 <sup>th</sup>	6
	Number of native non-tolerant benthic taxa	40 <sup>th</sup> – 50 <sup>th</sup>	5
	% native individuals as rheophilic	50 <sup>th</sup> – 60 <sup>th</sup>	6
	% native taxa as rheophilic	40 <sup>th</sup> – 50 <sup>th</sup>	5
	% native taxa as salmonid	40 <sup>th</sup> – 50 <sup>th</sup>	5
	Number of distinct Tolerant taxa	70 <sup>th</sup> – 80 <sup>th</sup>	8

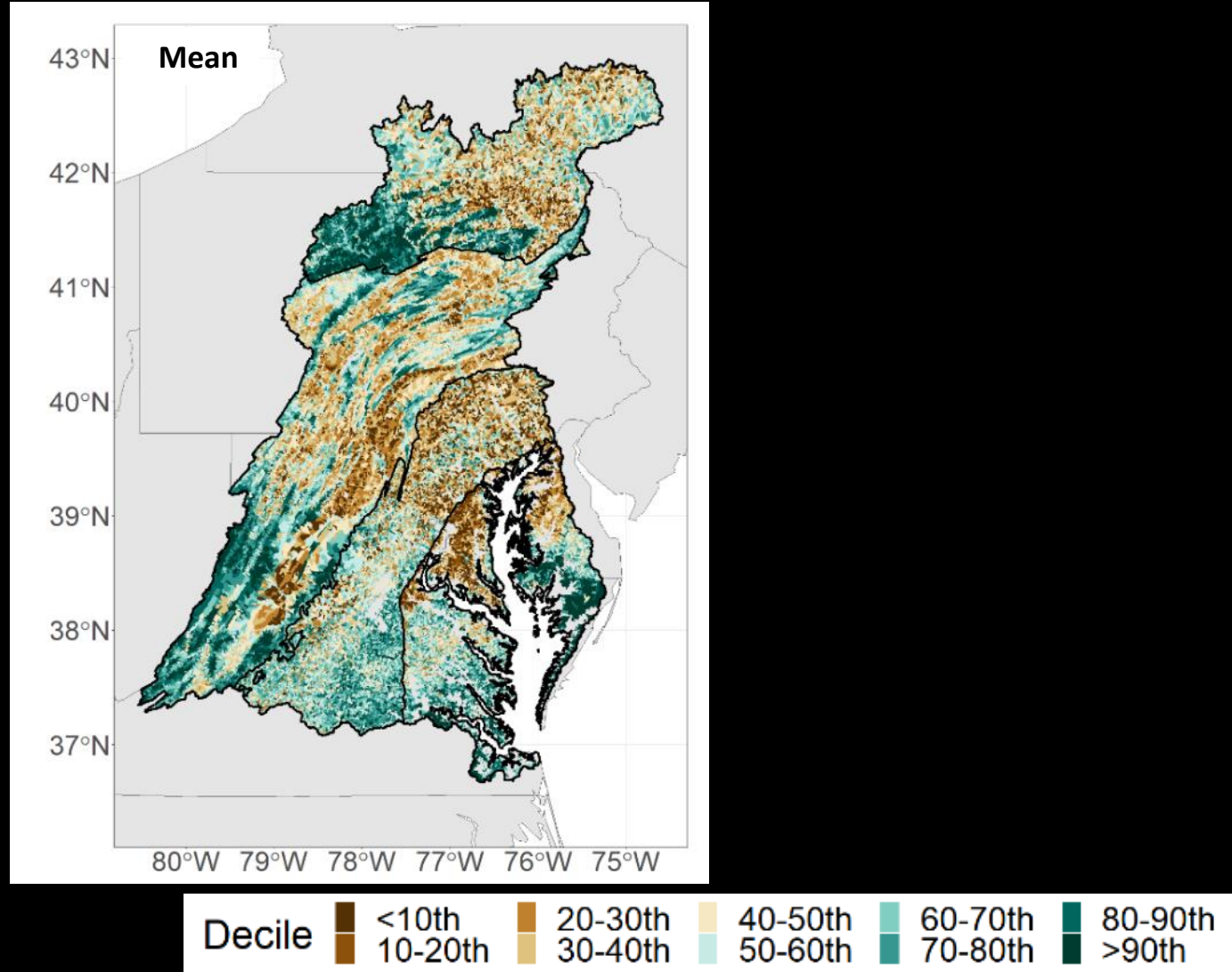


# Results – Community (2016 only)

Mean of metric  
deciles for each  
aggregated  
ecoregion

Lower deciles =  
poorer relative  
condition

Higher deciles =  
better relative  
condition



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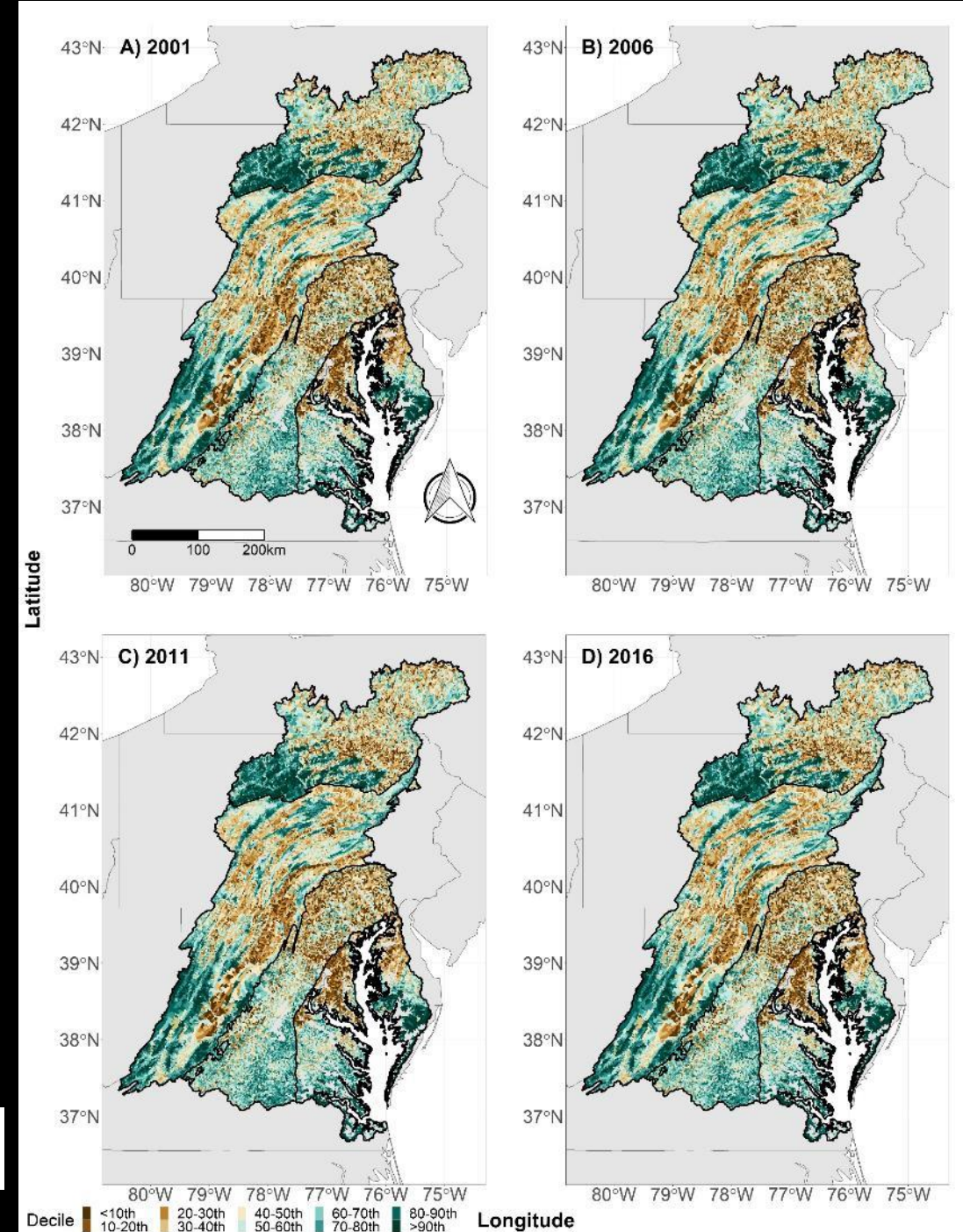
Higher deciles =  
higher SE



## Results – Community (4 years)

Have predicted values and uncertainty (SEs) for 4 periods where NLCD were available

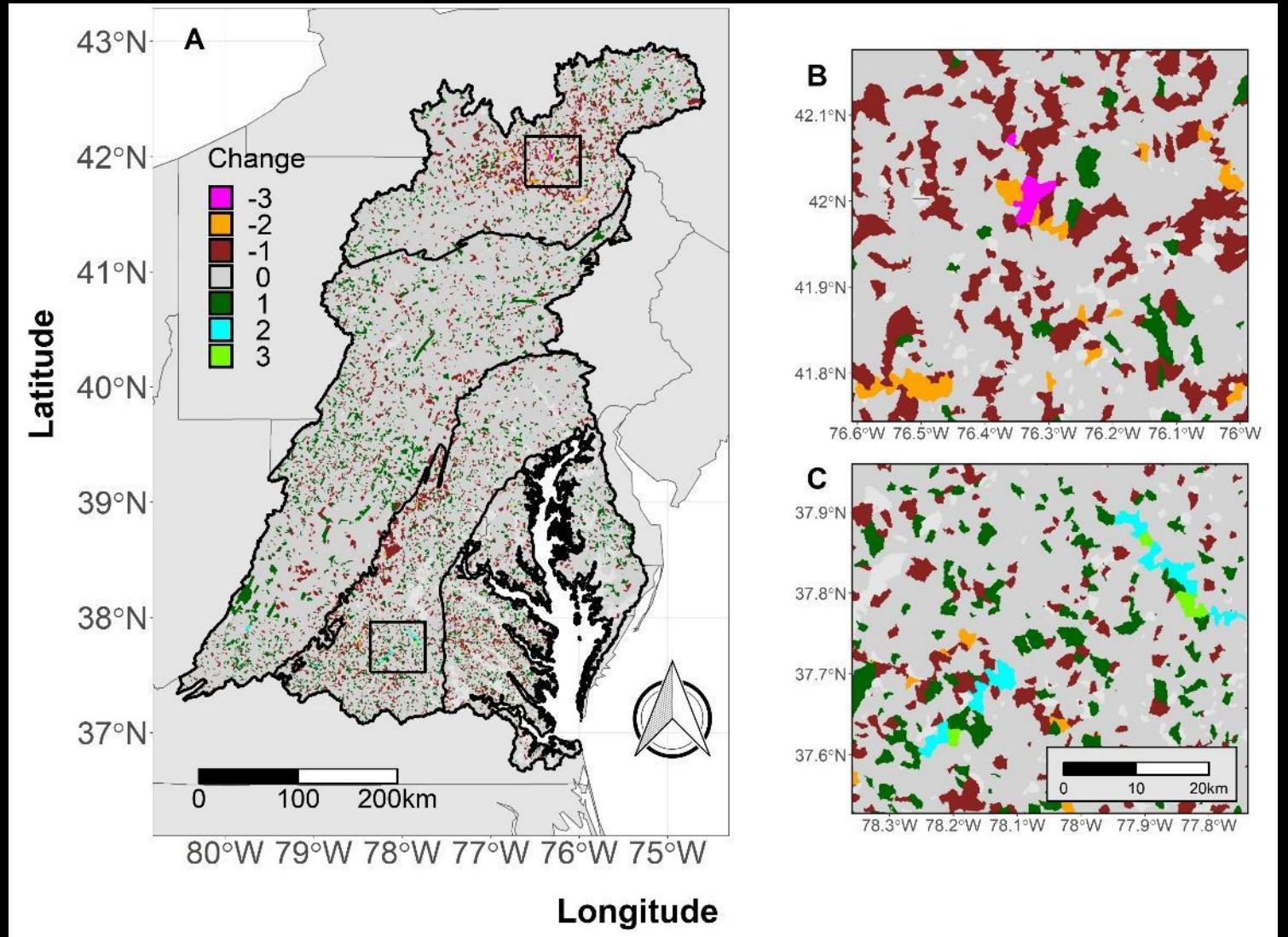
Allows a change between periods analysis



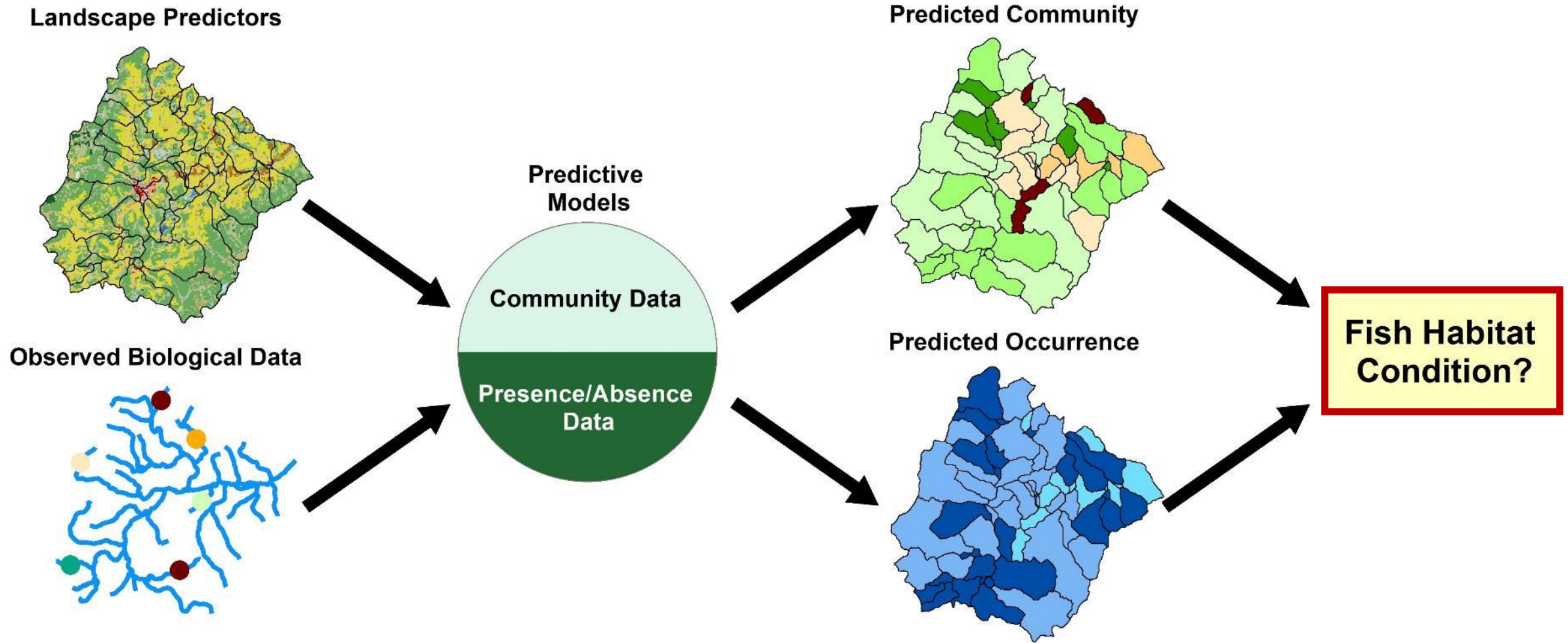
# Results - Community

## Change 2001 to 2016:

- When displayed spatially results can be used to identify locations with changes
- Those with a large positive change may be areas that have responded to management actions (future research need)
- Those with large negative changes may be areas requiring better management or restoration









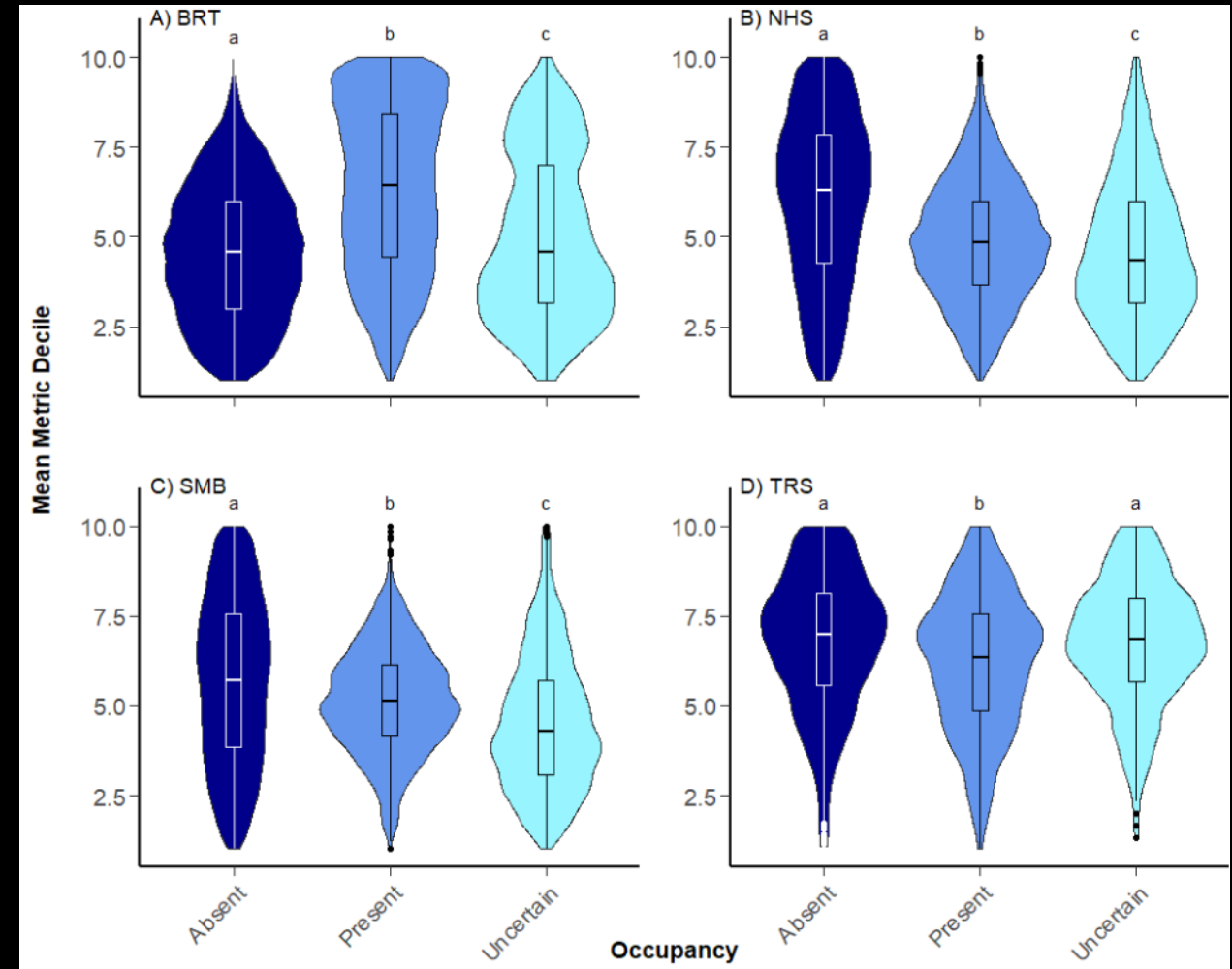
# Results – Community vs. Species

- Do community and species predictions align?
  - Expect mean metric decile and sensitive species to align

	Community	Species
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Strength	Useful for overall condition	Detailed insight into key species
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# Results – Community vs. Species

- Do community and species predictions align?
  - Expect mean metric decile and sensitive species to align
- Brook Trout predicted presences were in reaches with a higher mean deciles
- All other three species showed opposite pattern with predicted presences being higher in reaches with lower mean deciles

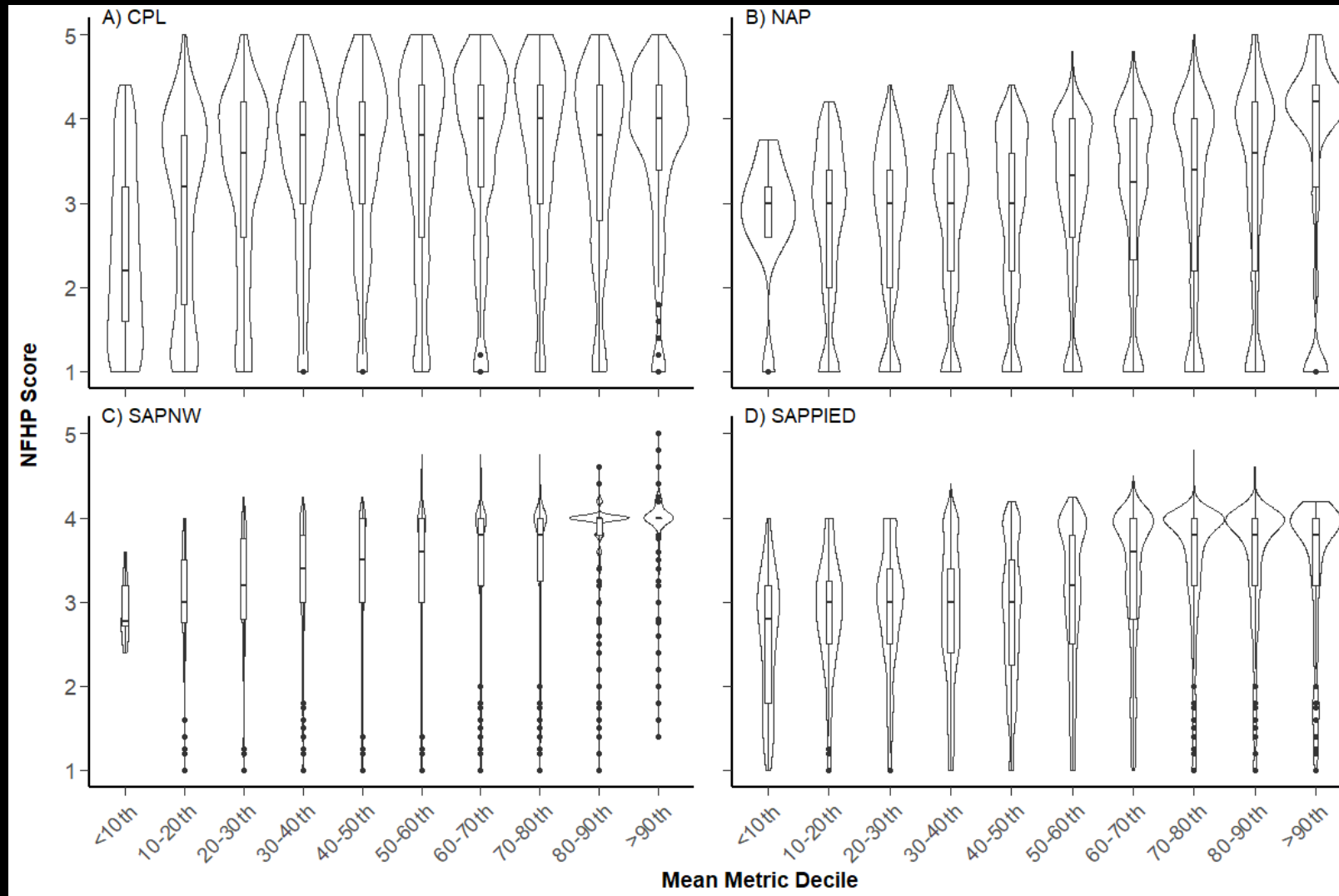


# Comparison to NFHP



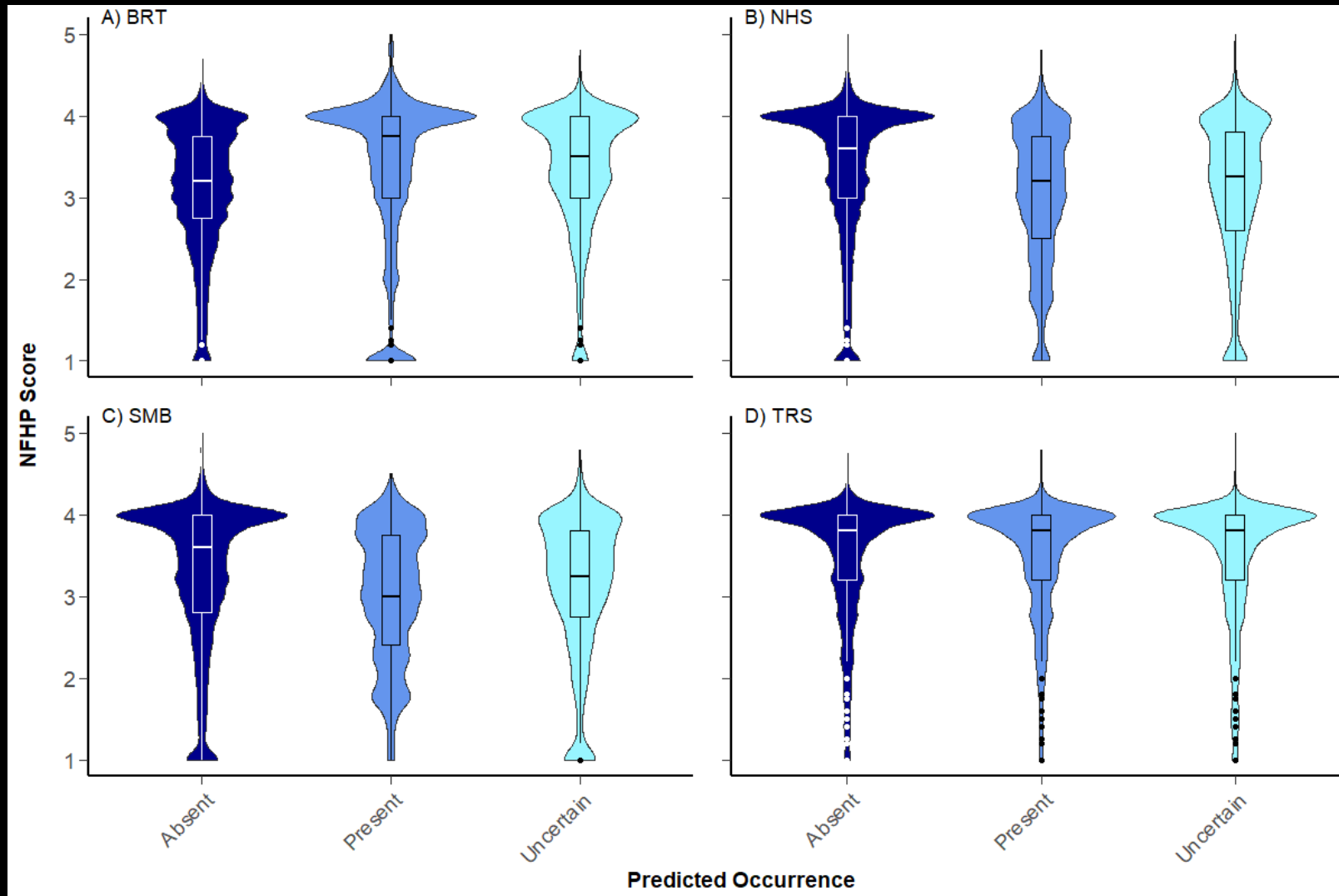
# Community vs NFHP

Mean decile and NFHP ratings generally agree but high variability



## Species vs NFHP

Brook Trout (BRT) agrees with NFHP, other sensitive species do not, all show high variability



## Summary

- Examined individual species habitat suitability
- Developed approach to identify habitat condition for 4 regions based on community metrics and mean deciles
- Community and species levels agreed for one sensitive species (Brook Trout); other two species may be less sensitive than USEPA suggests
- Using both approaches simultaneously may offer more insight into reach conditions
- Results from both approaches generally agree with the NFHP, but lots of variability was found



## Future Work

- Migrate to a 1:24,000 map scale framework
- Investigate finer resolution land use (e.g., 1 m) and summarization methods (e.g., distance-weighting)
- Incorporate future climate and land use scenarios

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- Investigate finer resolution land use (e.g., 1 m) and summarization methods (e.g., distance-weighting)
- Incorporate future climate and land use scenarios
- Develop a data visualization web-based application

Request feedback on development of this application to increase usefulness/relevance

## Data Visualization

- Static Storymap – only narrative with embedded figures and tables

[Endocrine Disrupting Compounds in the Chesapeake \(arcgis.com\)](#)

- Interactive Dashboard – users can query and select data that figures and tables display

[EcoSHEDS | USGS](#)



# Acknowledgements

- **Thank you to data providers**
- **Thank you to Karen Blocksom (EPA)**
- **Thank you to Stephen Faulkner and Richard Walker (USGS Chesapeake Regional Assessment Team)**
- **Support was provided by the USGS Fisheries Program and the Chesapeake Bay Program**

## **The following programs provided data for the Fish Database:**

- New York Department of Environmental Conservation
- Pennsylvania Department of Environmental Protection
- Pennsylvania Fish and Boat Commission
- West Virginia Department of Environmental Protection
- West Virginia Department of Natural Resources
- Maryland Department of the Environment
- Maryland Department of Natural Resources; Resource Assessment Service
- Maryland Department of Natural Resources; Fishing and Boating Services
- Anne Arundel County (MD) Department of Public Works
- Baltimore County (MD) Department of Environmental Protection and Sustainability
- Frederick County (MD) Sustainability and Environmental resources
- Howard County (MD) Storm Water Management Division
- Montgomery County (MD) Department of Environmental Protection
- Virginia Department of Game and Inland Fish
- Virginia Commonwealth University
- Virginia Department of Environmental Quality
- Fairfax County (VA) Department of Public Works & Environmental Sciences
- National Parks Service Shenandoah National Park
- Susquehanna River Basin Commission
- United States Environment Protection Agency
- United States Geological Survey

# Thank you!

## Contact Information:

### Assessment of fish habitat:

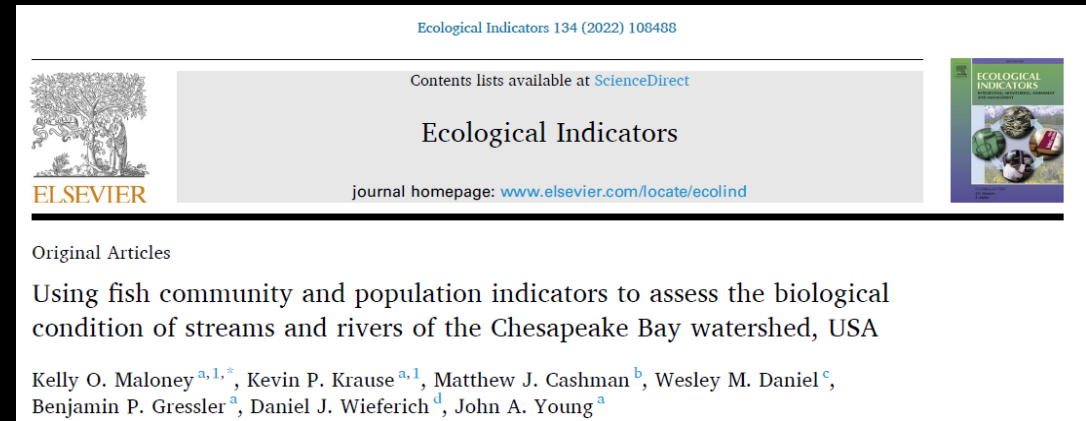
Kelly Maloney ([kmaloney@usgs.gov](mailto:kmaloney@usgs.gov))

Kevin Krause ([kkrause@usgs.gov](mailto:kkrause@usgs.gov))

## Data Visualization:

Kelly Maloney ([kmaloney@usgs.gov](mailto:kmaloney@usgs.gov))

Stephanie Gordon ([sgordon@usgs.gov](mailto:sgordon@usgs.gov))



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Data releases: <http://dx.doi.org/10.5066/P9B4BMAG>

<http://dx.doi.org/10.5066/P9D6JU4X>

<http://dx.doi.org/10.5066/P9C1PX4P>

### Science Summary:

[Assessing the habitat conditions to support freshwater fisheries in the Chesapeake Watershed | U.S. Geological Survey \(usgs.gov\)](#)