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Comparing multiple scales of land use pressures and instream stressors on headwater stream macroinvertebrates



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Overarching Theme







Shift happens in

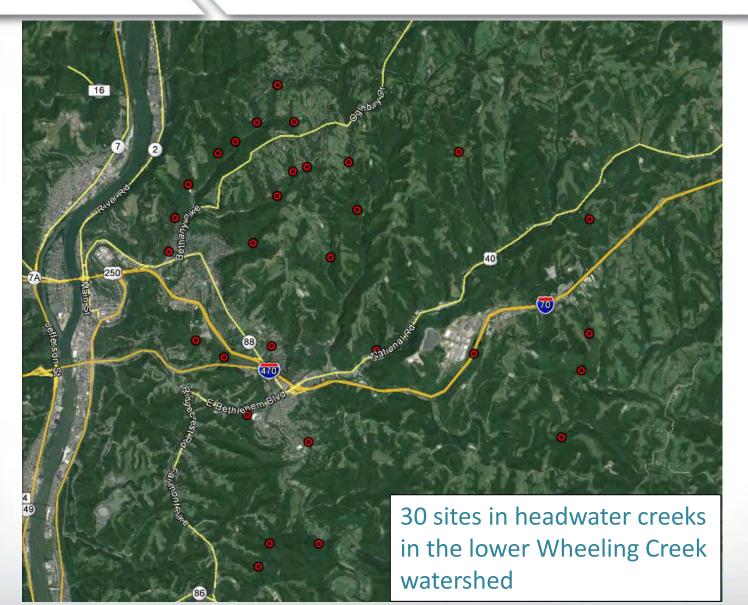
macroinvertebrate assemblages along a gradient of land uses and instream measures

SH!FT HAPPENS

https://drmikelharry.files.wordpress.com/









Reach selection

Reaches selected by similar:

- 1. Stream Size/ Catchment Area
- 2. Full streambed shading
- 3. Elevation
- 4. General lithology
- 5. Dominant soil type
- 6. Stream channel morphology



Study Watershed and Land Use

Legend

Benthic Site Locations



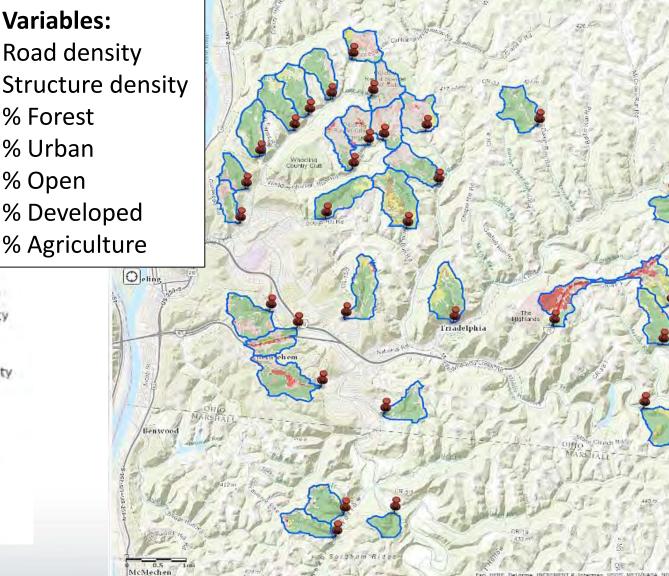
Subcatchment Areas

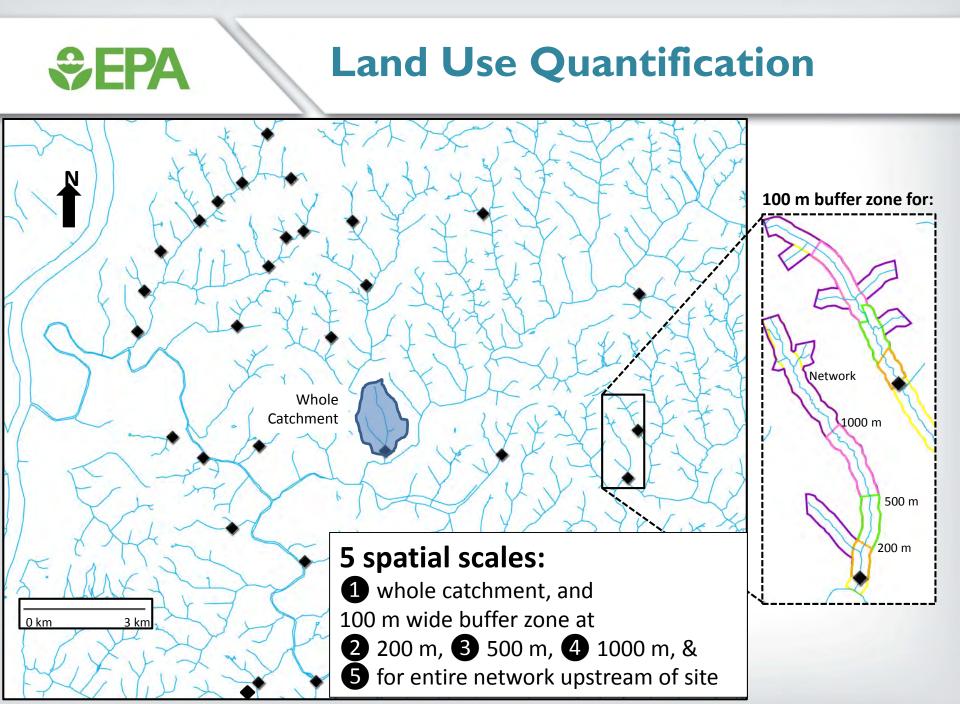


NLCD 2011

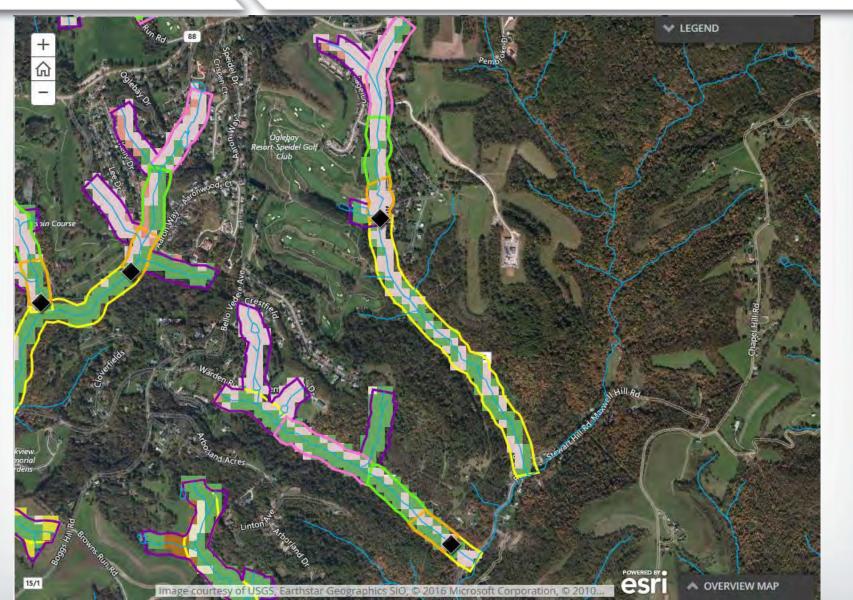


Road density % Forest % Urban % Open % Developed Subcatchment Areas Developed, Open Space Developed, Low Intensity Developed, Medium Intensity Developed, High Intensity Barren Land Deciduous Forest Evergreen Forest Mixed Forest Herbaceuous Hay/Pasture





SEPA Land Use Quantification







Sampling Parameters



Macroinvertebrates-Riffle kicknet method

\$EPA

Sampling Parameters



Macroinvertebrates-Riffle kicknet method



Habitat- RBP, channel slope, canopy cover, width

Sampling Parameters



Macroinvertebrates-Riffle kicknet method



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Sampling Parameters



Macroinvertebrates-Riffle kicknet method



Habitat- RBP, channel slope, canopy cover, width





Benthic Algae- Dominant type & mean concentration

Sampling Parameters



Macroinvertebrates-Riffle kicknet method



Habitat- RBP, channel slope, canopy cover, width





Benthic Algae- Dominant type & mean concentration



Water Chemistry- In situ & grab samples

Sampling Parameters



Macroinvertebrates-Riffle kicknet method



Habitat- RBP, channel slope, canopy cover, width





Benthic Algae- Dominant type & mean concentration



Water Chemistry- In situ & grab samples



Fecal Coliform Bacteria-Processed within 6 hours

Statistical Analyses

- Developed and compared simple and multiple linear regression models to explain and predict macroinvertebrate shifts along environmental gradients;
- Multivariate analysis (PCA, NMDS) to explore environmental and biological response patterns;
- Land use classes are spatially contagious and co-vary with instream measures, so...
 - partial correlation analysis to remove redundancy

- stepwise-multiple regressions with low tolerance and low variance-inflationfactors
- Data were normalized by log, sqrt, or asn-sqrt transformations where necessary



Study Questions

1 Does near-stream (buffer zone) land use affect assemblages more than whole catchment-based land use?

Prediction: Near-site (200 m)-buffer zone > whole catchment

2 Does the best spatial arrangement of land use pressures better predict biological condition compared to instream measures (habitat and chemistry)?

Prediction: local instream factor > best land use indicator

3 Develop and compare strength of multivariable explanatory models based on combinations of instream and land use measures. What level of effort (field, lab, land use, combinations) is appropriate?

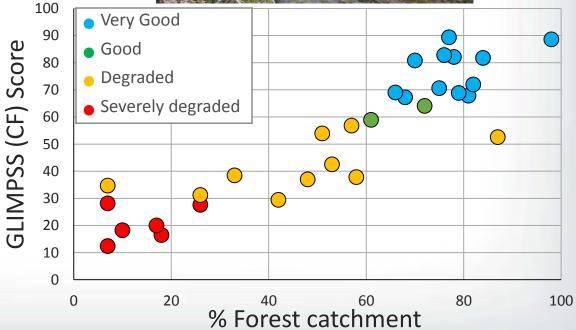




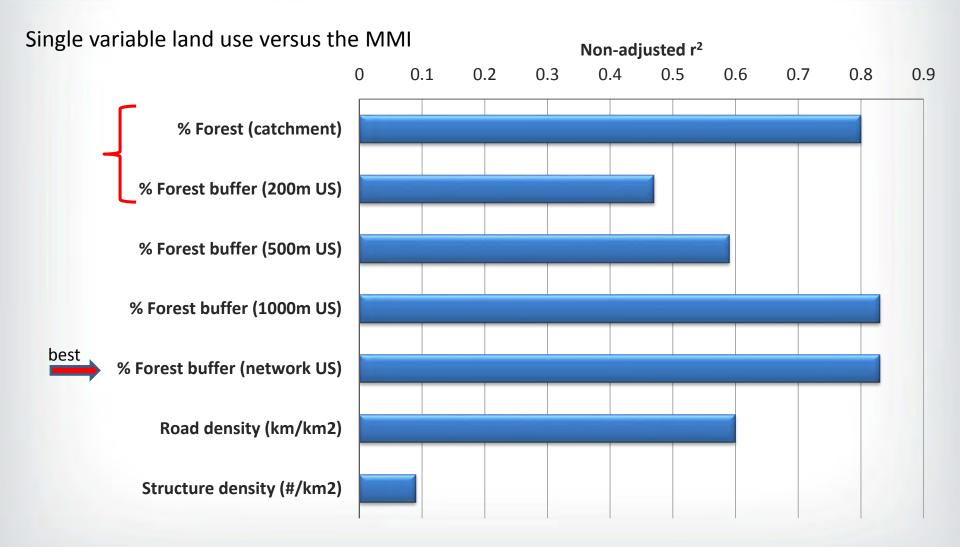
GLIMPSS (CF) scores ranged from 12 (very degraded) to 89 (very good)





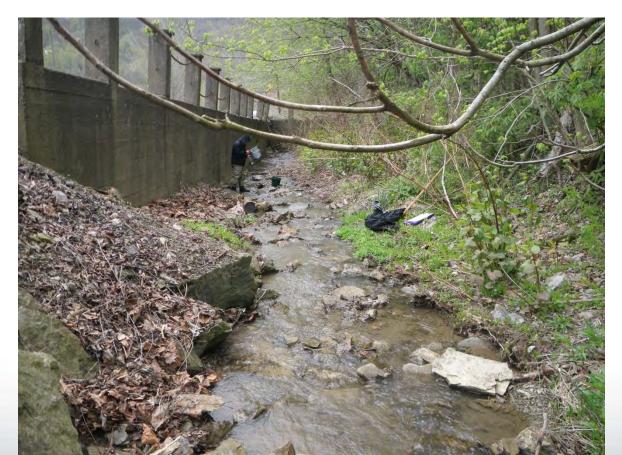


Provide Simple Linear Regression Models





Near-site (200 m) buffer zone land use explains <u>less</u> variation in macroinvertebrates than whole catchment.





Study Questions

1 Does near-stream (buffer zone) land use affects assemblages more than whole catchment-based land use?

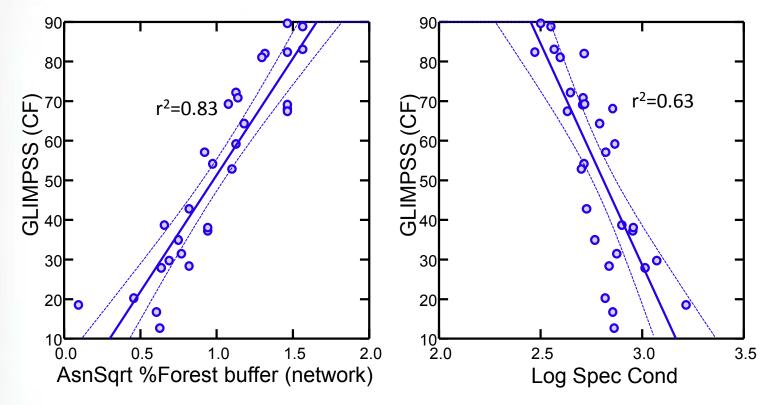
Prediction: near-site (200 m) buffer zone > whole catchment

2 Does the best spatial arrangement of land use pressures better explain biological condition compared to instream measures (habitat or chemistry)?

Prediction: local instream factor > best land use indicator

3 Develop and compare strength of multivariable explanatory models based on combinations of instream and land use measures. What level of effort (field, lab, land use, combinations) is appropriate?

Question **2** Simple Linear Regression Models



Land Use vs. Biology

SEPA

- 1. %forest buffer (network) (r²=0.83)
- 2. % forest buffer (1000m) (r²=0.83)
- 3. % forest catchment (r²=0.80)
- 4. % developed catchment (r²=0.80)

Instream vs. Biology

- 1. Spec. Cond. (r²=0.63)
- 2. Hardness (r²=0.58)
- 3. Chloride (r²=0.48)
- 4. Total habitat score (r²=0.46)



Prediction 2 refuted since land use classes were stronger single predictors than any single instream variable.





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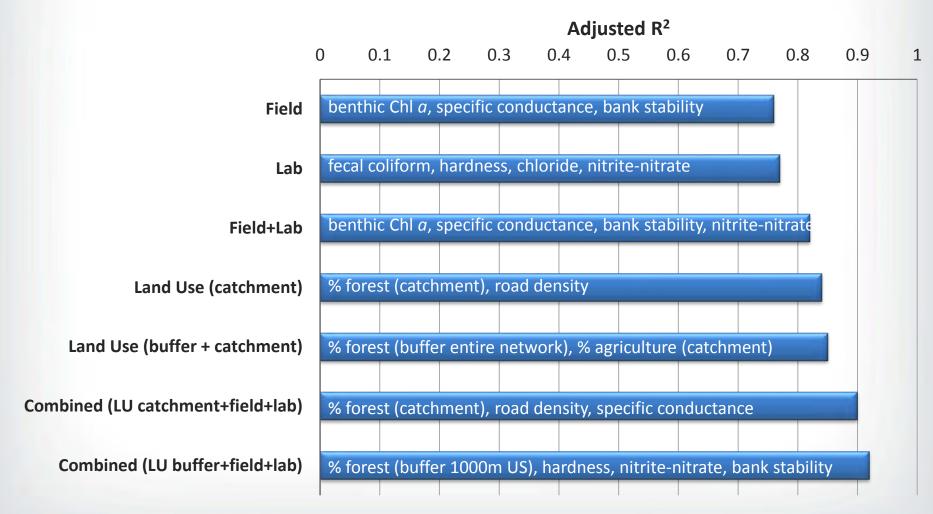
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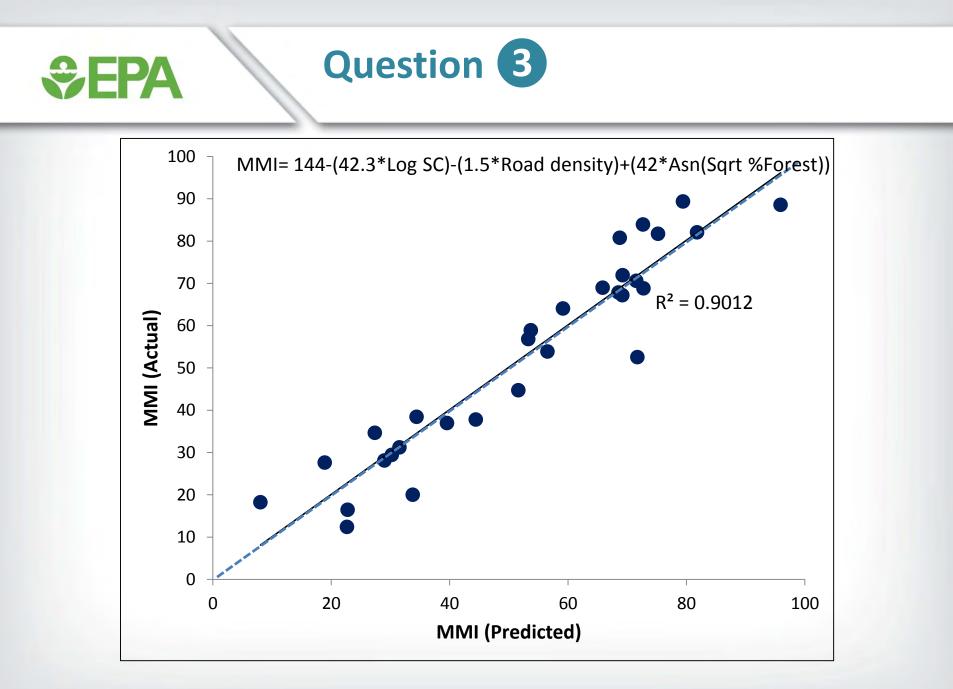
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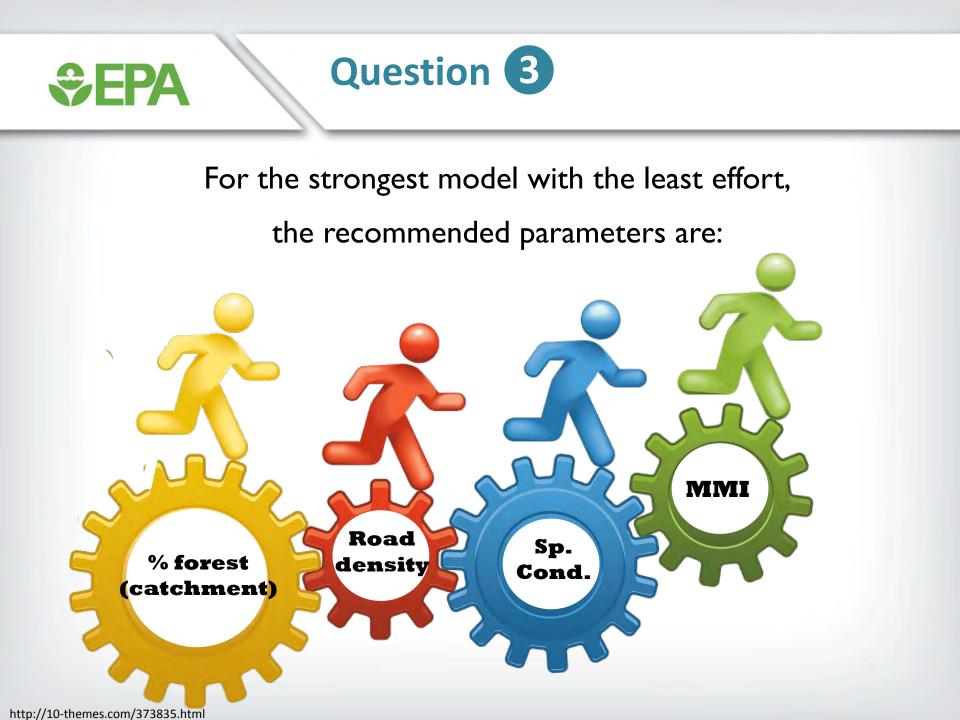
What level of effort (field, lab, land use, combinations) is appropriate?



Stepwise Multiple Regression Models





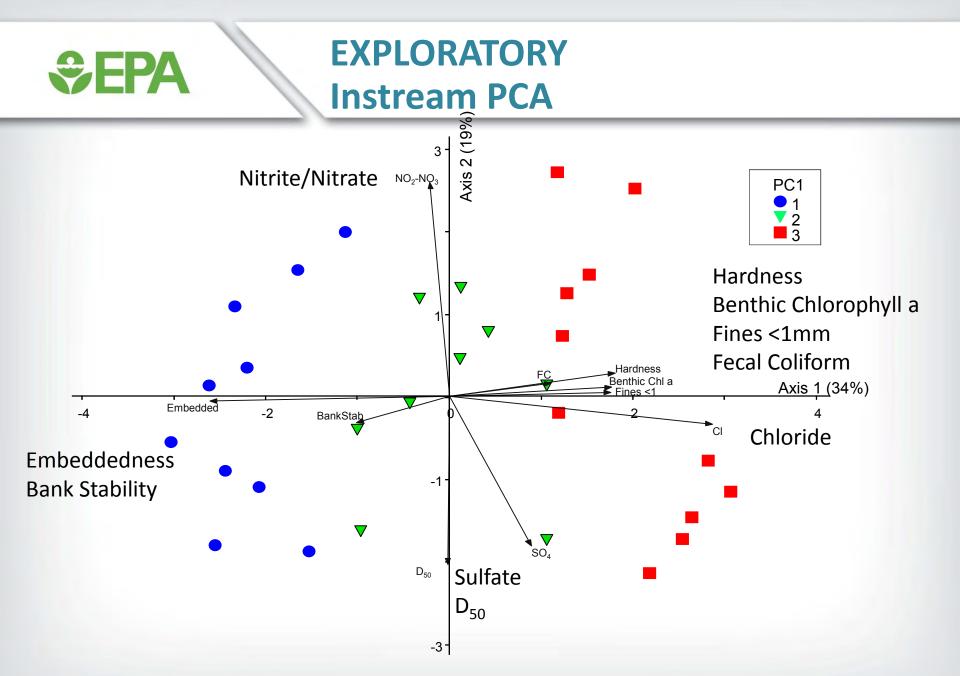




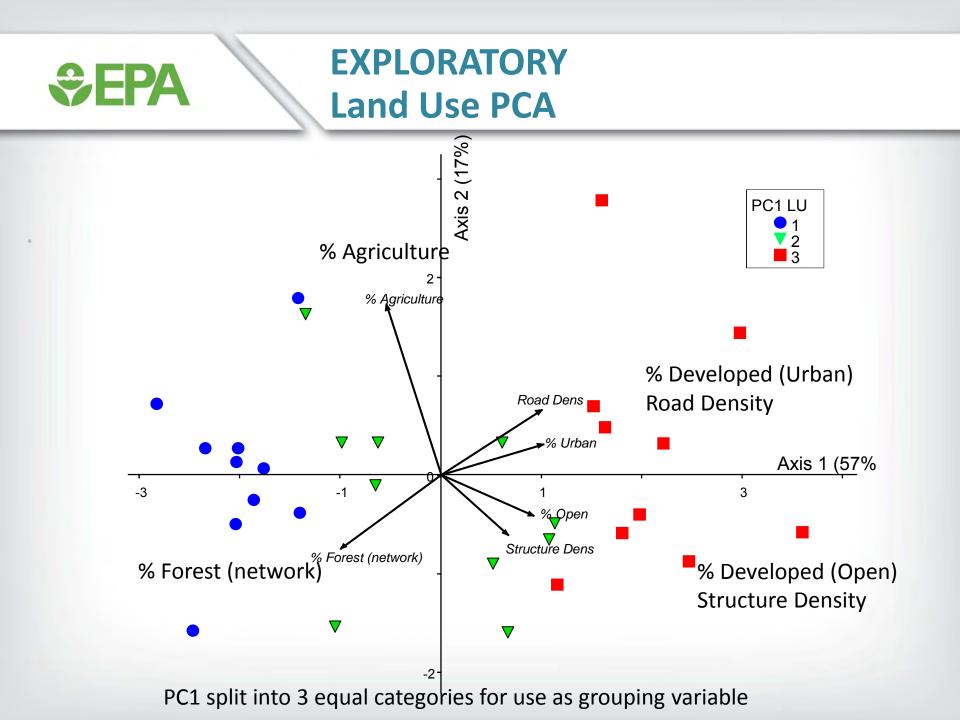
Digging Deeper: Exploratory Analyses



http://conservationmagazine.org

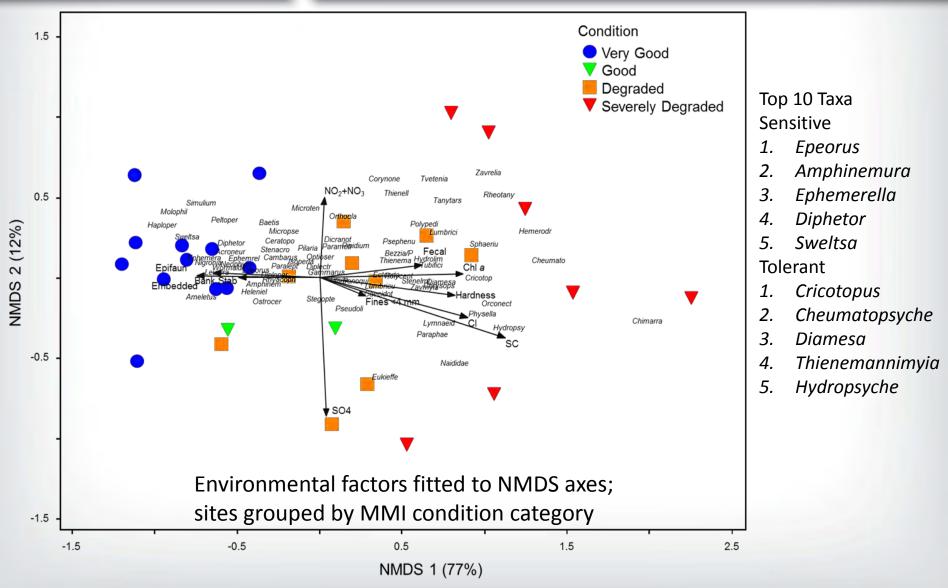


PC1 split into 3 equal categories for use as grouping variable



NMDS Ordination Community Structure

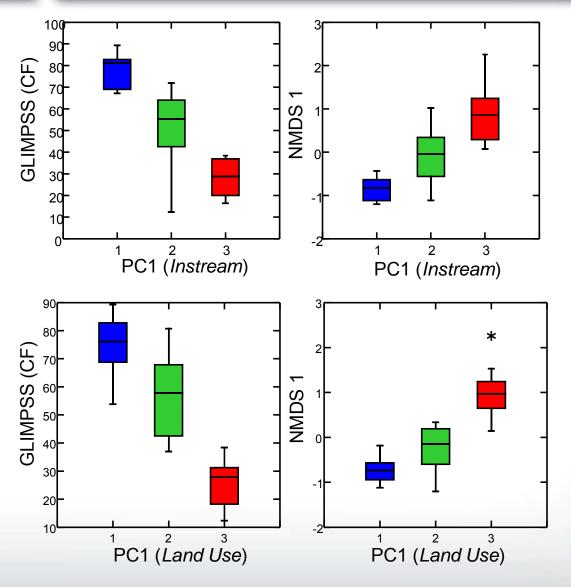
SEPA



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Biological Response

Macroinvertebrate MMI and NMDS 1 were similarly responsive to both *Instream* and *Land Use* stressor gradients (PCA) in the multivariate world



Conclusion

Landscape models can be used for prediction or simple explanation, but resources managers are often tempted to use without biological data.



Local models should never replace biological assessments.

GIS-based models can provide managers with decision tools but are deficient without accompanying instream data.

Our models could help target areas for more intensive monitoring, prioritization of conservation areas, and/or selection of reference sites.



QUESTIONS???



Thank you to EPA R3 employees Frank Borsuk and Don Evans, interns Trevor Dunn and Lyndsey Burton, and the EPA R3 Laboratory