Reach-scale stream restoration in agricultural streams of southern Minnesota alters structural and functional responses of macroinvertebrates



Christy Dolph OF MINNESOTA **Upper Midwest Stream Restoration Symposium** February 24th, 2014



Acknowledgements

- U.S. EPA STAR Fellowship
- MPCA 319 funds
- Coauthors:
 Sue Eggert
 Joe Magner
 Len Ferrington
 Bruce Vondracek
- Rochelle Roche
- Rhithron Associaties, Inc.
- Joel Chirhart, MPCA



Stream Restoration

- \$1 billion spent in U.S. annually
- Biodiversity effects rarely evaluated
- "If you build it, they will come"?
- Palmer et al. 2010 \rightarrow no improvements in biodiversity

Restoration: Effects beyond diversity?

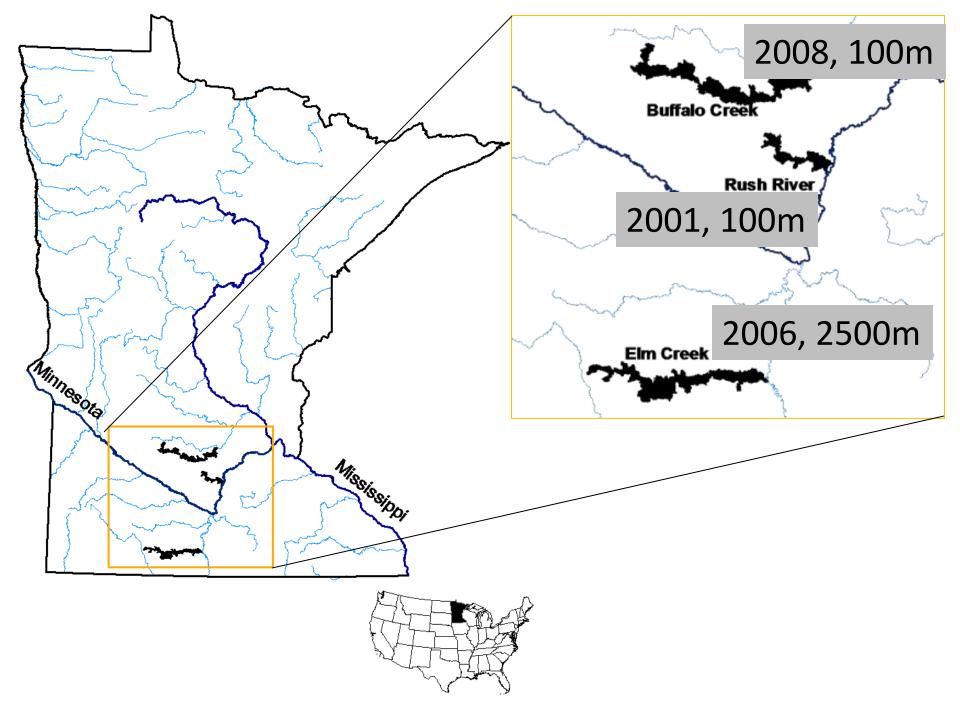
- Macroinvertebrate Secondary Production
 - Invert biomass produced over time
 - Reproduction, growth rate, survivorship, density
 - Ecosystem function: energy flow thru food webs
 - May reflect increases in habitat or food for inverts
 - May represent food resources for higher trophic levels



Objectives

I. Compare invert community structure in restored and unrestored reaches of three streams: Total richness, #EPT, IBI

2. Compare secondary production in restored and unrestored reaches of all three streams



Study System Characteristics

- 3rd order
- Land use: corn, soy, cattle, hogs
- Flashy
- High sediment loads





Stream Restoration Goals: Reduce channel erosion, improve habitat

Actions:

I.Adding boulders/wood

Redirect flow
 Strengthen banks
 Structural heterogeneity

2. Bank re-vegetation =

Stabilize banks
 Improve bank habitat

3. Engineer benches

Prevent sloughing





For each stream:

- Restored Reach (100m)
- Unrestored Reach (100m)
- Two reach types separated by 100m





Macroinvertebrates: biodiversity & secondary production



6 samples: Apr-Nov, 2010







Sampling approach:

- 5 macroinvertebrate habitat types
- riffle/run
- overhanging banks
- emergent veg
- woody debris
- debris dam





- D-frame dip net
- Sample I sq. ft habitat

Taxa Richness

- 121 taxa (mostly genus IDs) total, 52 families
- 62 Diptera
- 19 Ephemeroptera
- 13 Coleoptera
- II Trichoptera
- 6 Plecoptera
- 4 Hemiptera
- 3 Odonata
- I Amphipoda
- I Collembola

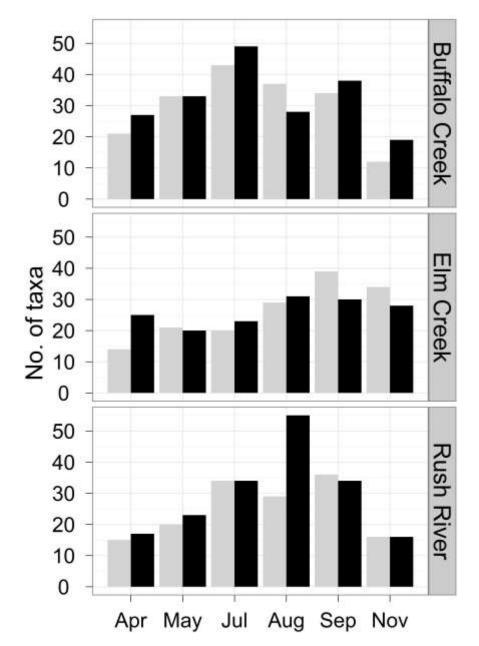








Total taxa richness



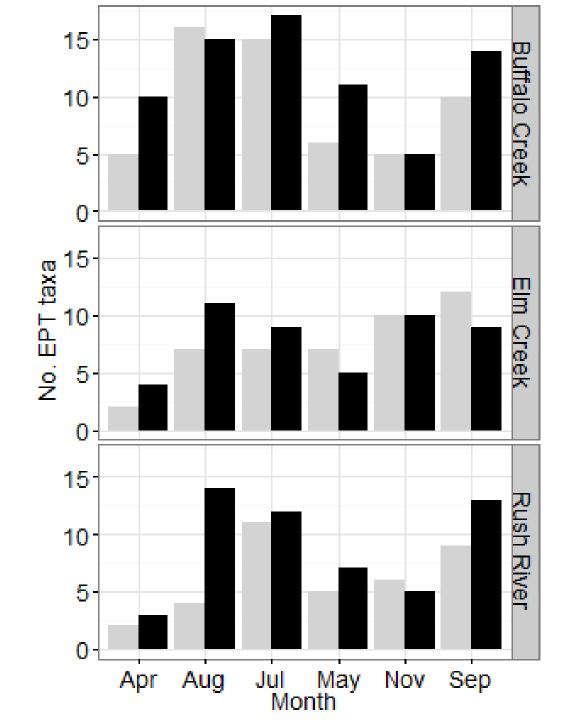
Unrestored Restored

No significant difference in richness between restored and unrestored reaches

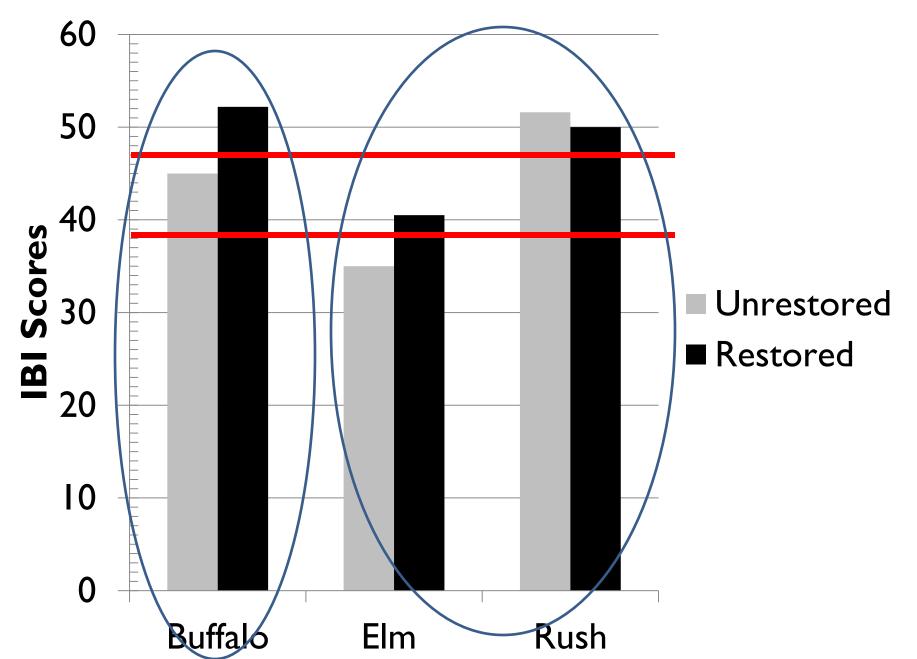
of EPT taxa

Unrestored Restored

Controlling for site and month effects, **restored reaches yielded 2 additional EPT taxa** compared to unrestored reaches, on average



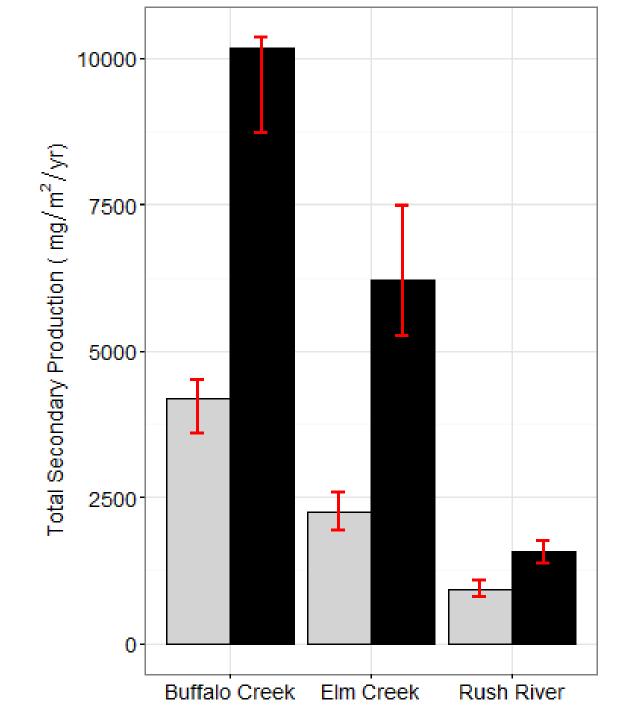
IBI Scores: Prairie & Forest, Low-Gradient



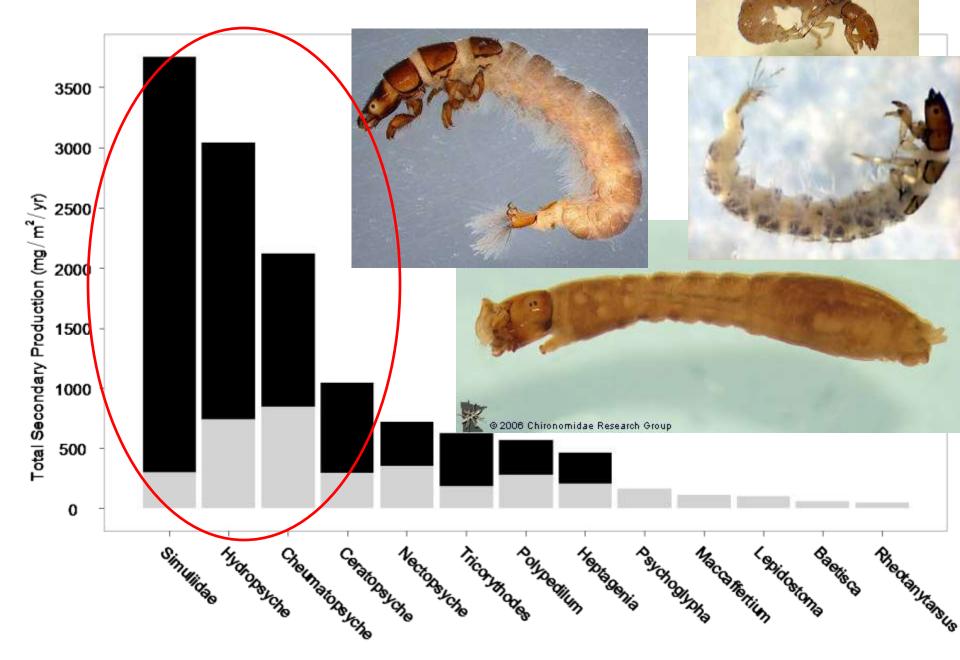
Secondary Production

Production 2-3 x higher in restored reaches

> Unrestored Restored



Dominant Taxa: Buffalo Creek

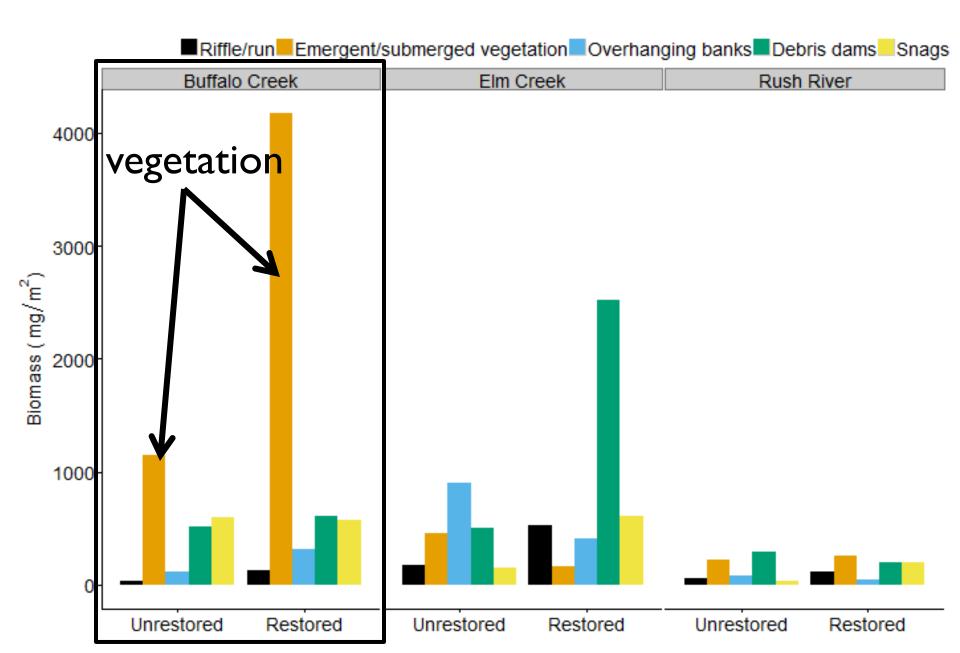


Conclusions

- No difference in taxa richness \rightarrow larger-scale drivers may limit sensitive species
- Reach scale restoration may have effects beyond total taxa richness (#EPT, IBI, production)
- Higher production of some taxa may indicate more stable habitat or higher quality food resources
- Conservation implications: production effects limited to dominant, tolerant taxa?

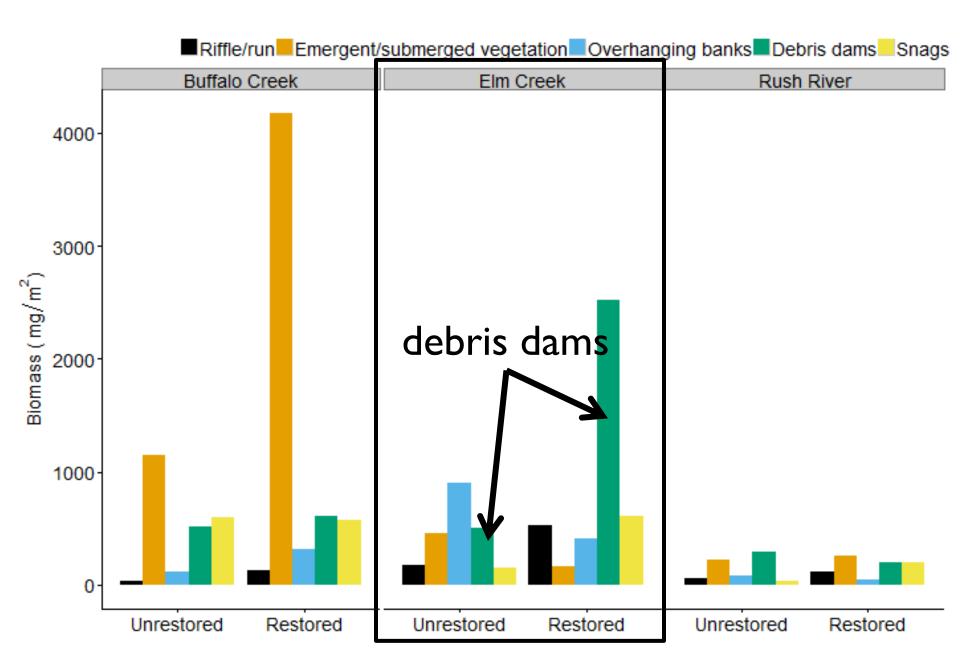


Biomass by Habitat Type

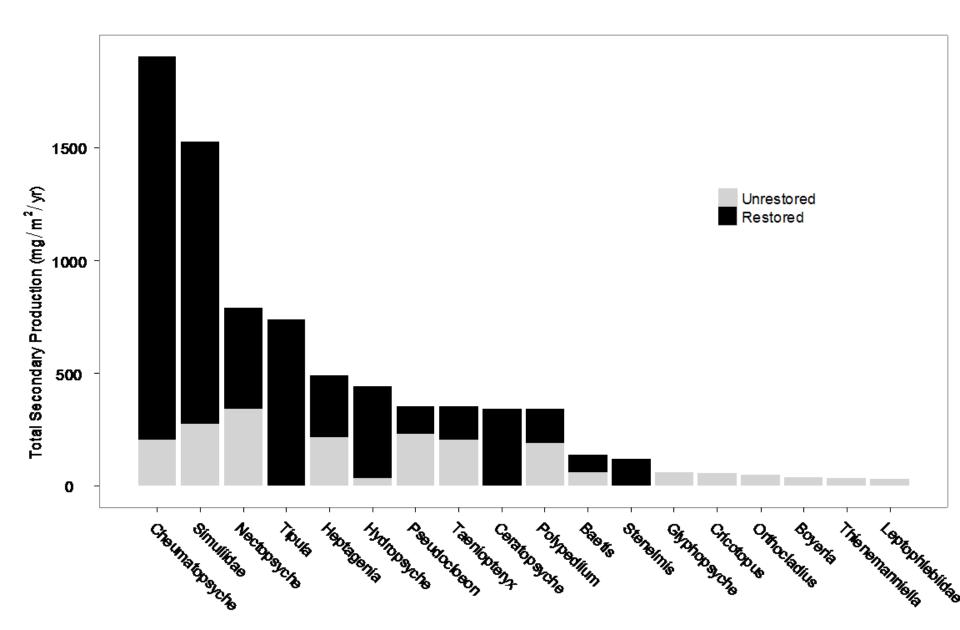




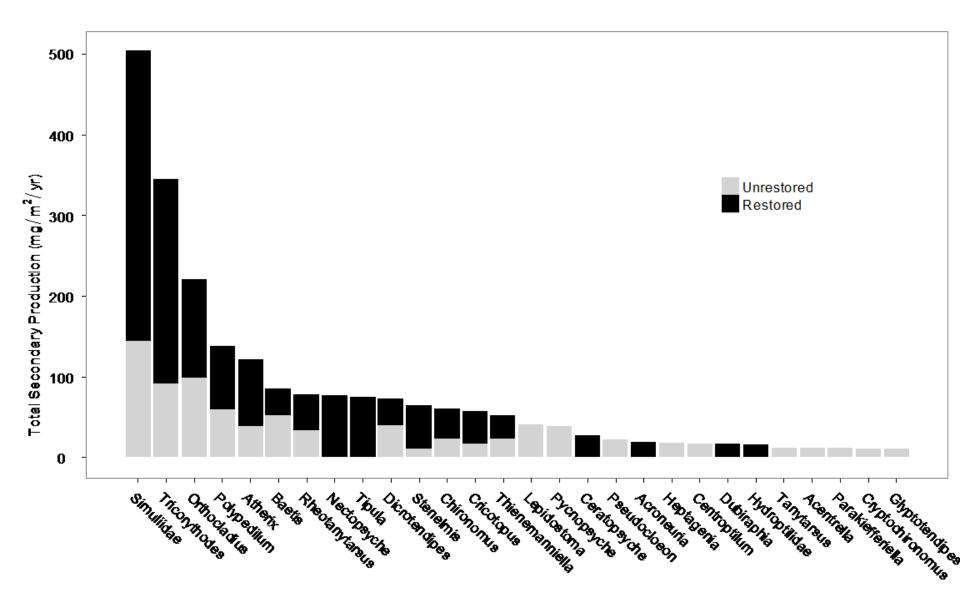
Biomass by Habitat Type



Dominant Taxa: Elm Creek



Dominant Taxa: Rush River



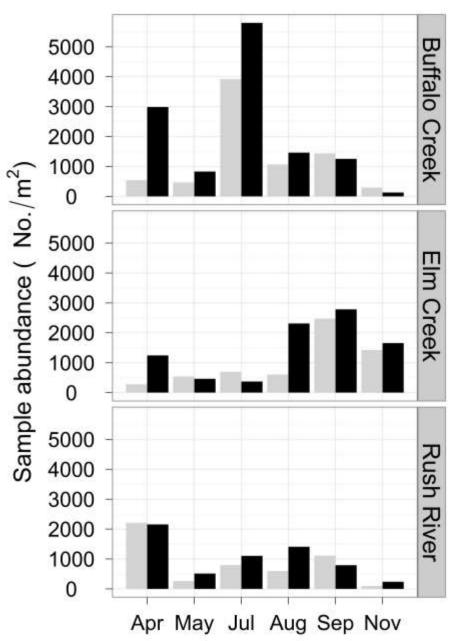




In-stream Habitat

Habitat	Buffalo Creek		Elm Creek		Rush River	
	U	R	U	R	U	R
% embedded	86 %	64%	63%	40 %	24%	28%
% sand	70%	48%	30%	24%	54%	44%
% gravel	8%	24%	52%	56%	10%	16%
% cobble	8%	20%	12%	14%	20%	30%

Invertebrate abundance



Unrestored Restored

No significant difference in <u>mean</u> <u>abundance</u> between restored and unrestored reaches

Biomass

Mean biomass per visit:

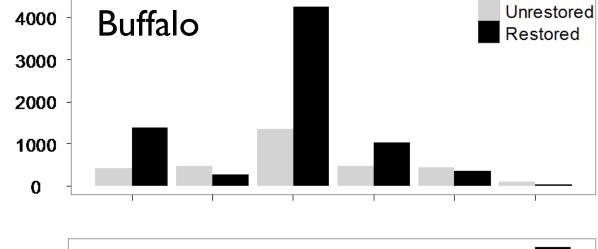
Restored = 760 mg/m²

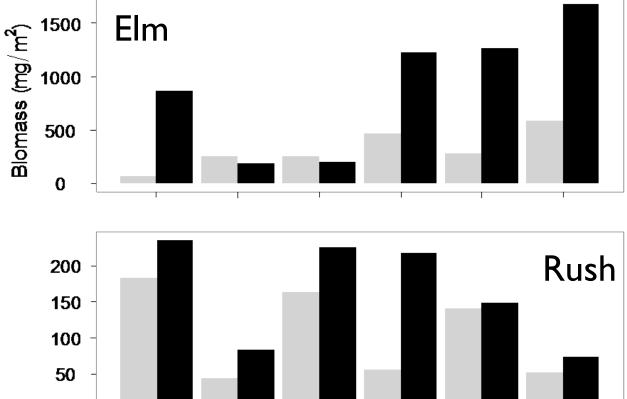
Unrestored = 320 mg/m²

0

Apr

May





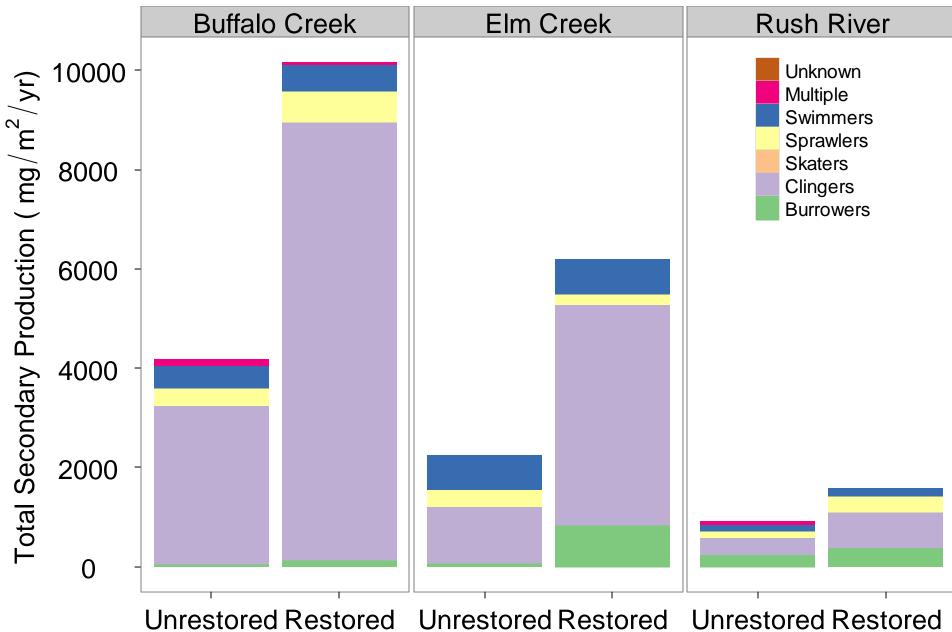
Jul

Aug

Sep

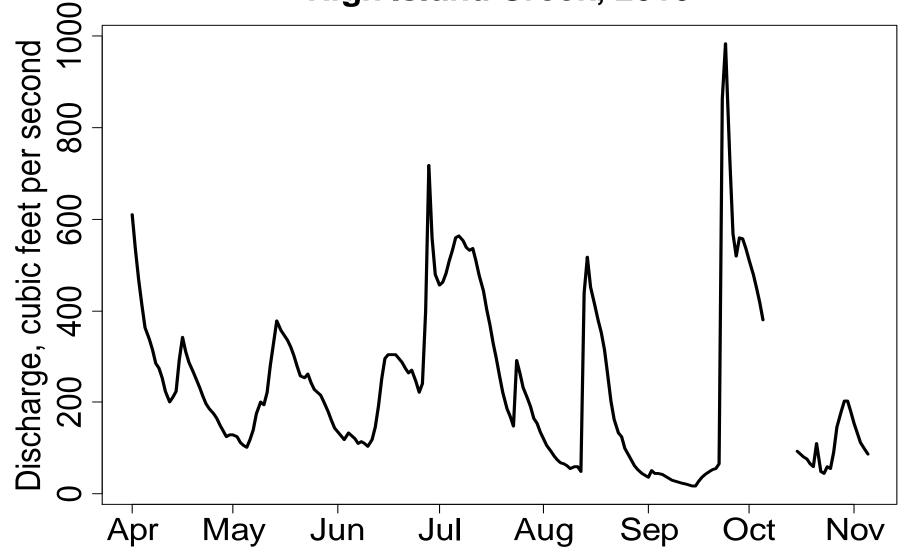
Nov

Production by Habit Groups



Drainage & Discharge

High Island Creek, 2010



Methods

<u>Biodiversity</u> = Taxa richness (most common taxonomic unit = genus)

<u>Secondary production</u> = Biomass over time

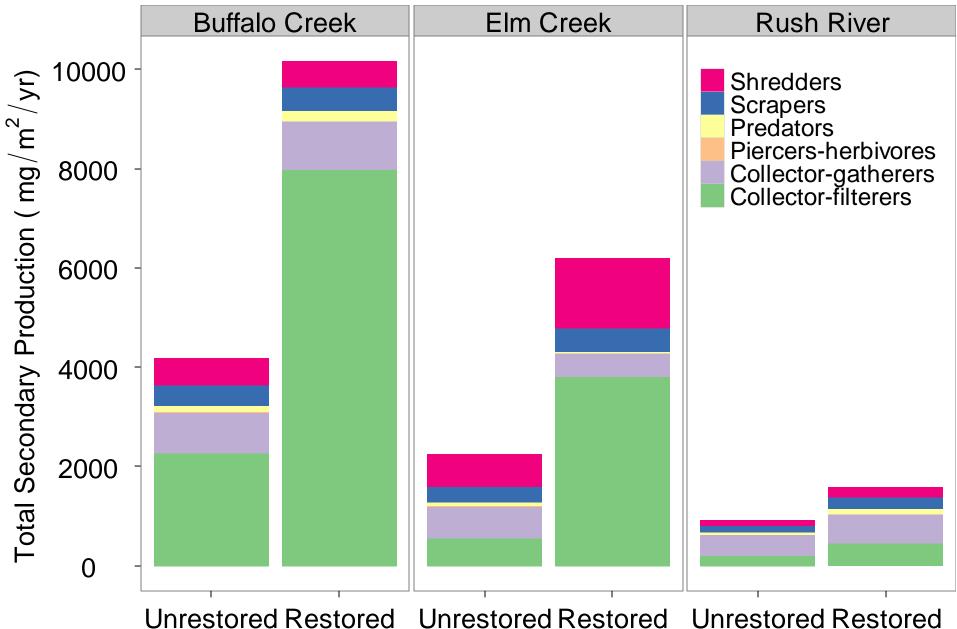
- I. Measure length of each specimen (extra work!)
 2. Use existing length-mass regression equations to determine mass
- 3. Calculate production (P) from mass using one of three methods:

Methods

in R

- Size-frequency method
- Instantaneous Growth Method
- Published P/B ratios

Production by Functional Feeding Groups



Density

UnrestoredRestored

Controlling for site and month effects, **density was 258 individuals/m2 higher in restored reaches** compared to unrestored reaches, on average

