



Spatial variability of methane production and methanogen communities within a eutrophic reservoir: evaluating the importance of organic matter source and quantity

Megan E. Berberich¹, Jake J. Beaulieu², Trinity L. Hamilton³, Sarah Waldo², Ishi Buffam¹

¹ University of Cincinnati, Department of Biological Sciences

² National Risk Management Research Laboratory, USEPA, Office of Research and Development, Cincinnati, OH

³ University of Minnesota, College of Biological Sciences



Freshwater CH₄ emissions (Tg C yr⁻¹)

Lakes: 53.7

Ponds: 12

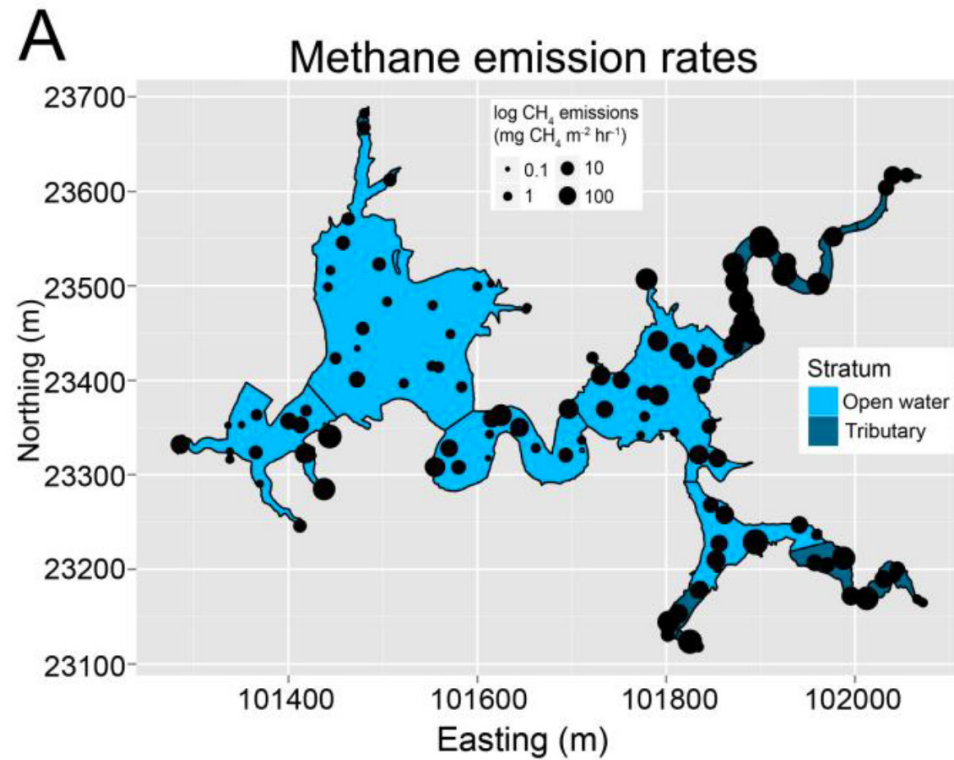
Rivers: 1.1 - 20.1

Reservoirs: 13.3 – 52.5

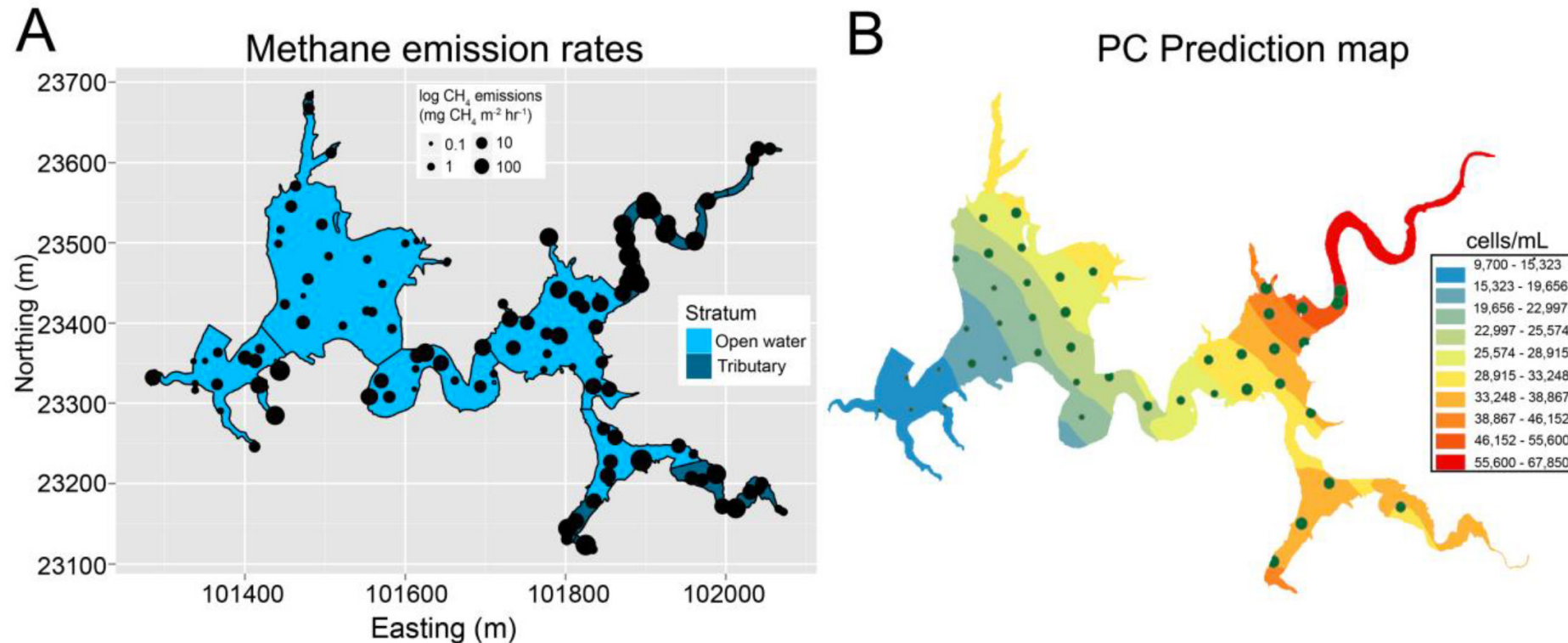
From:

Bastviken et al. (2011), Deemer et al. (2016), Holgerson and Raymond (2016), Stanley et al. (2016)

Surface methane emissions can be higher in tributary zones of reservoirs
(Beaulieu et al. 2014).



Methane emissions (Deemer et al. 2016, DelSontro et al. 2018) and sediment methane production (Duc et al. 2010, West et al. 2015) are positively correlated with variables associated with reservoir productivity.



Data from Beck et al. 2017 courtesy of Chris Nietch, USEPA

Algal OM additions to sediment incubations stimulated greater increase in CH₄ production than terrestrial OM additions (West et al. 2012).

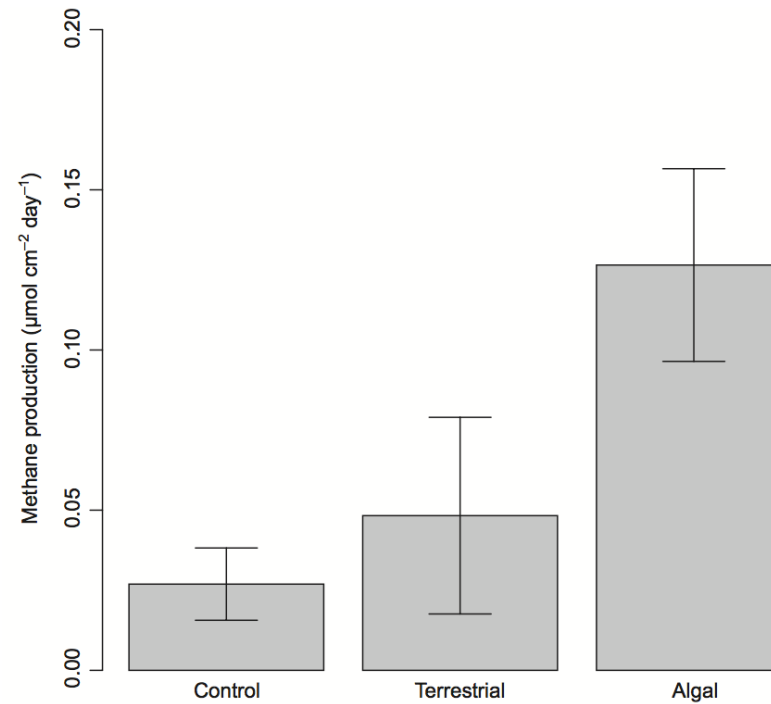
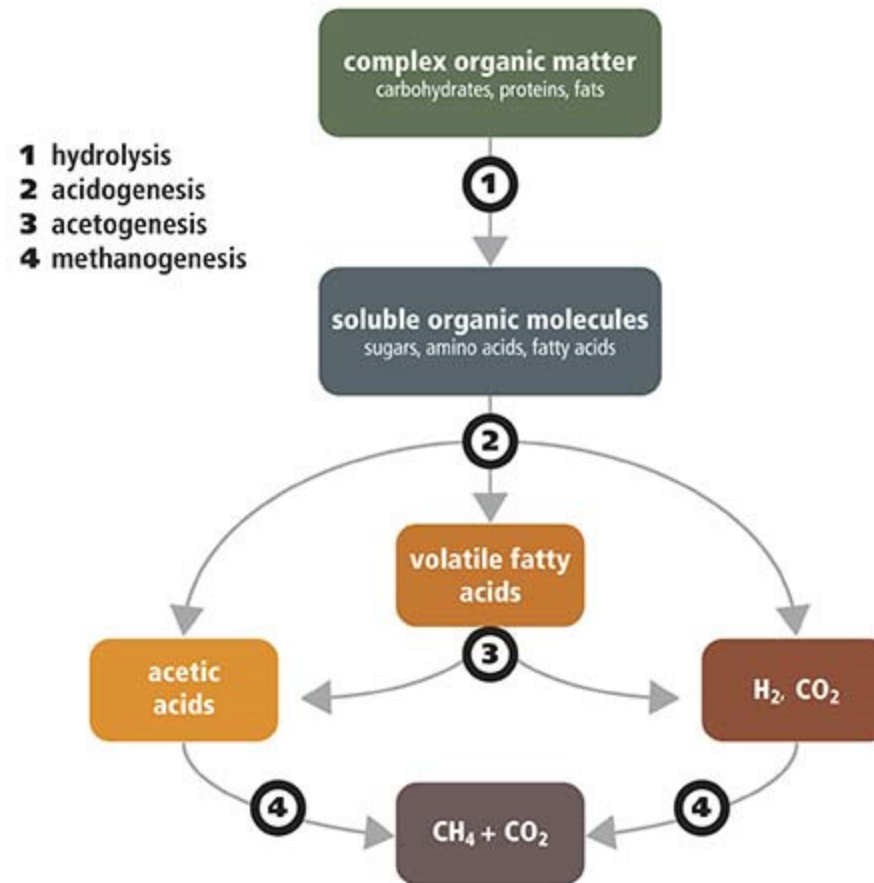
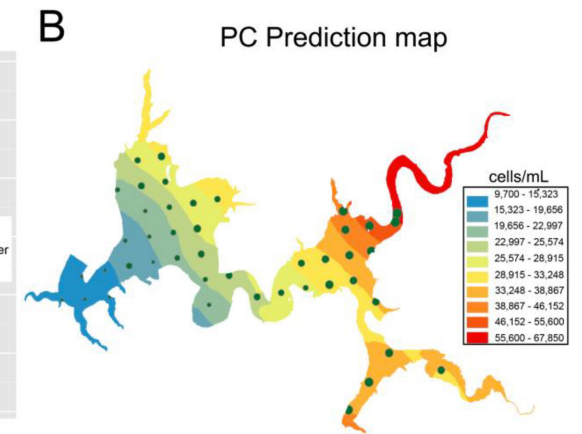
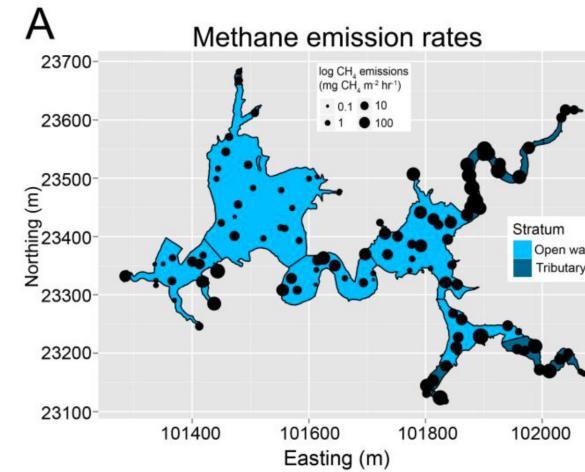
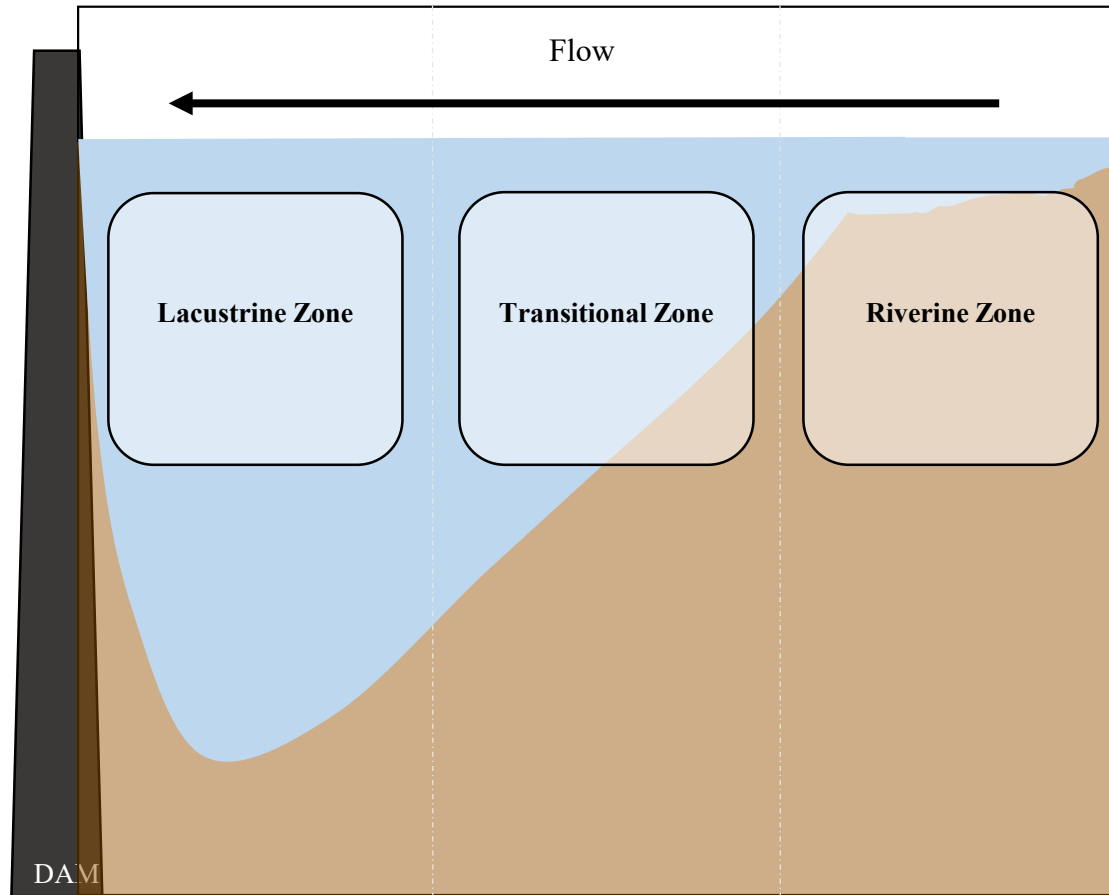


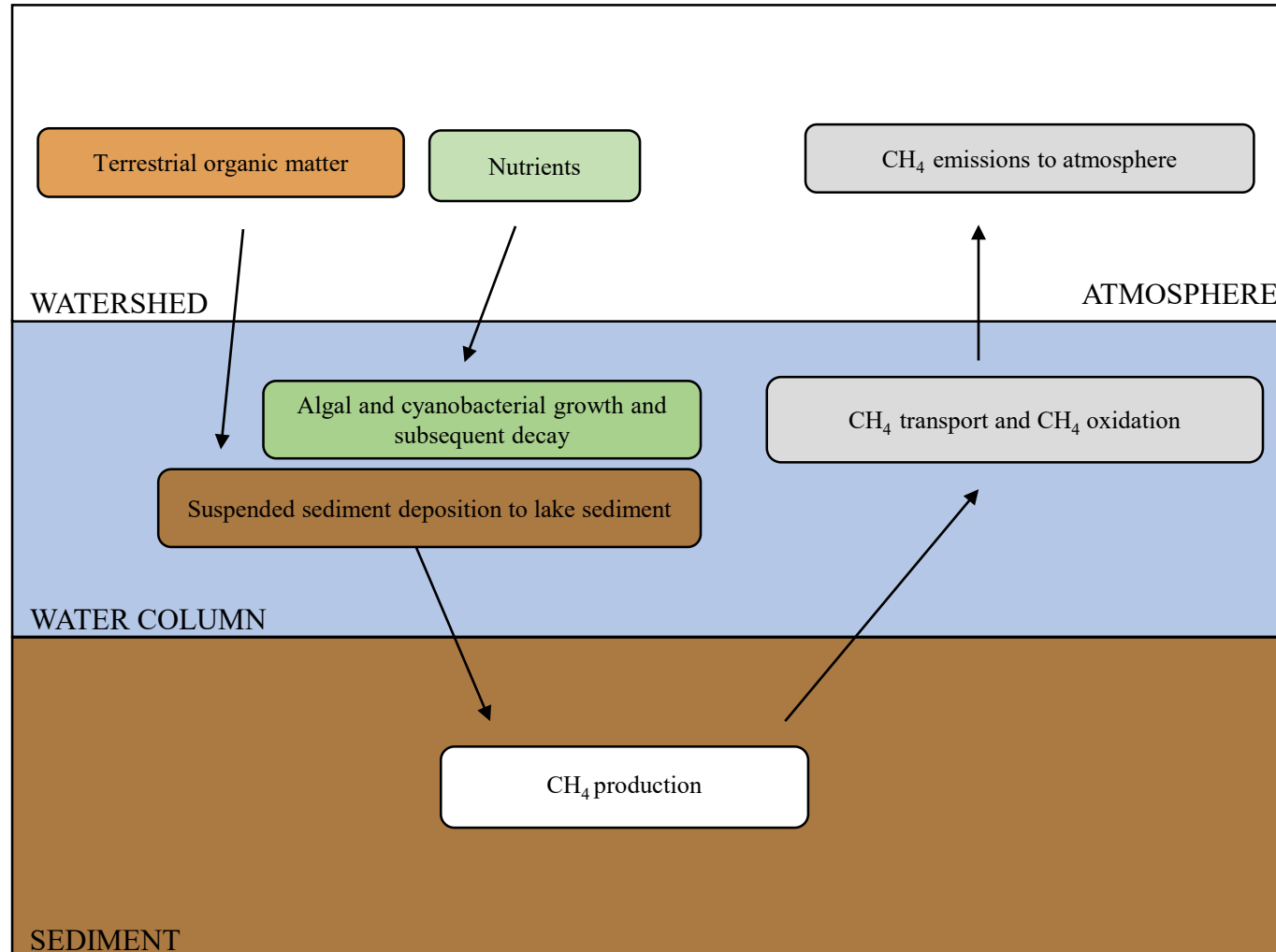
Fig. 2 Mean CH₄ production rates of Diamond Lake sediment cores with added algal biomass, added terrestrial carbon and control (one-way ANOVA, $F = 12.5$, $P = 0.007$, d.f. = 2, 6). Error bars indicate one standard deviation of the mean.

Organic matter (OM) availability is a constraint on sediment methane production.



Reservoirs can be divided into functional zones based on spatial variability (Thornton et al. 1990).





Objectives

1. Establish if **methane production rates** and **OM source and quantity** exhibit distinct variation among reservoir zones.

Objectives

1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.

Objectives

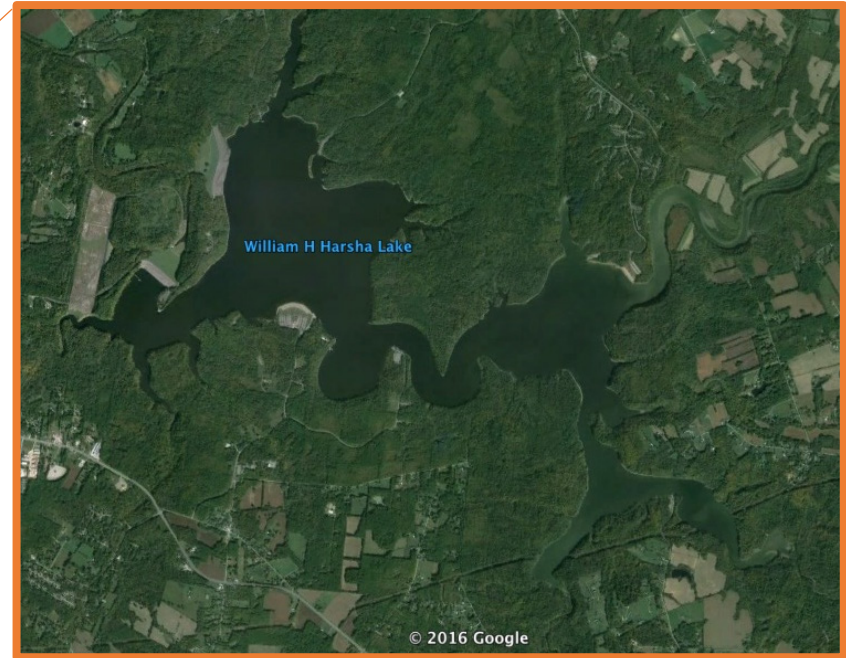
1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.
3. Describe the variations in **abundance** and **taxonomy** of **methanogen communities**.

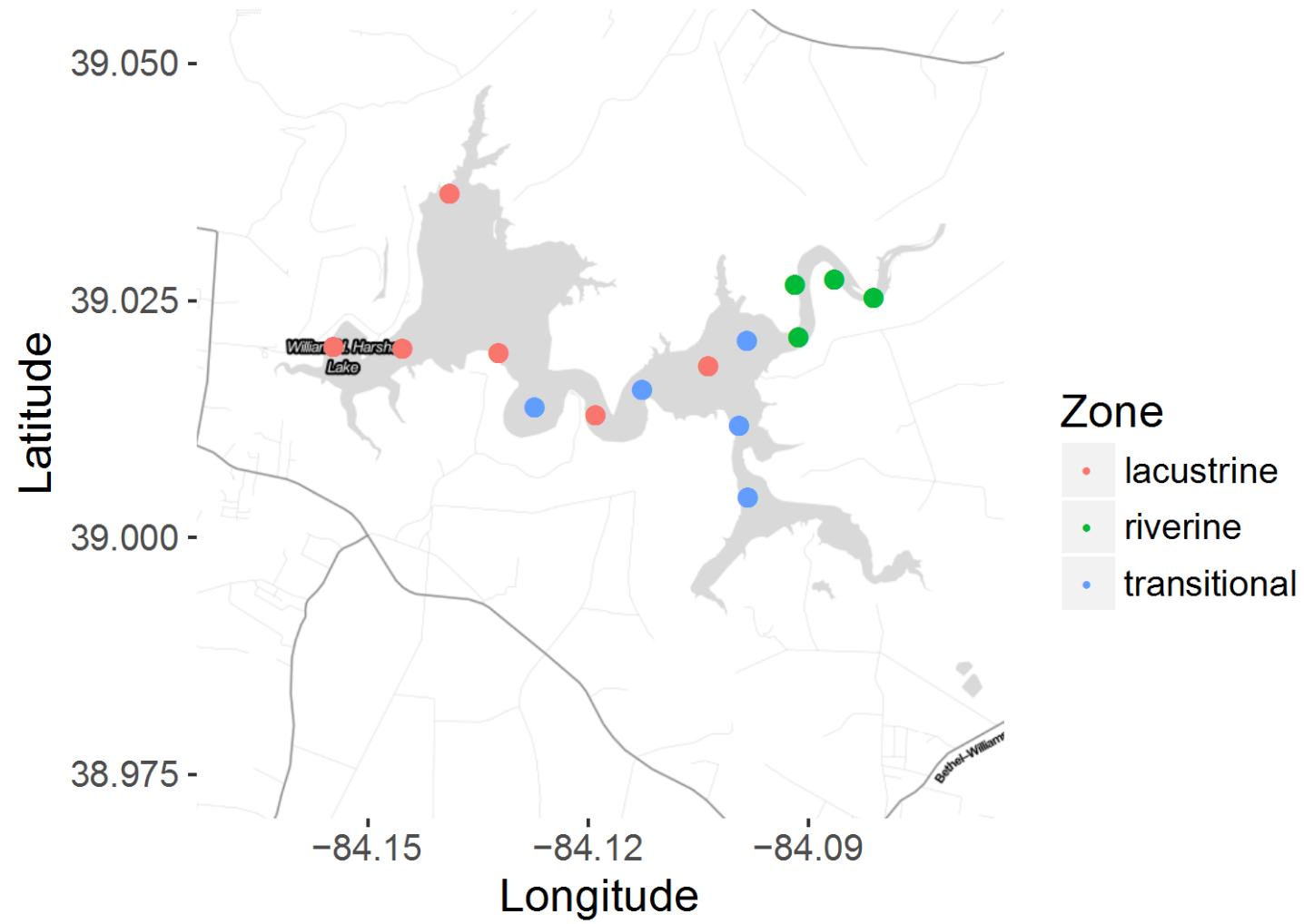
Objectives

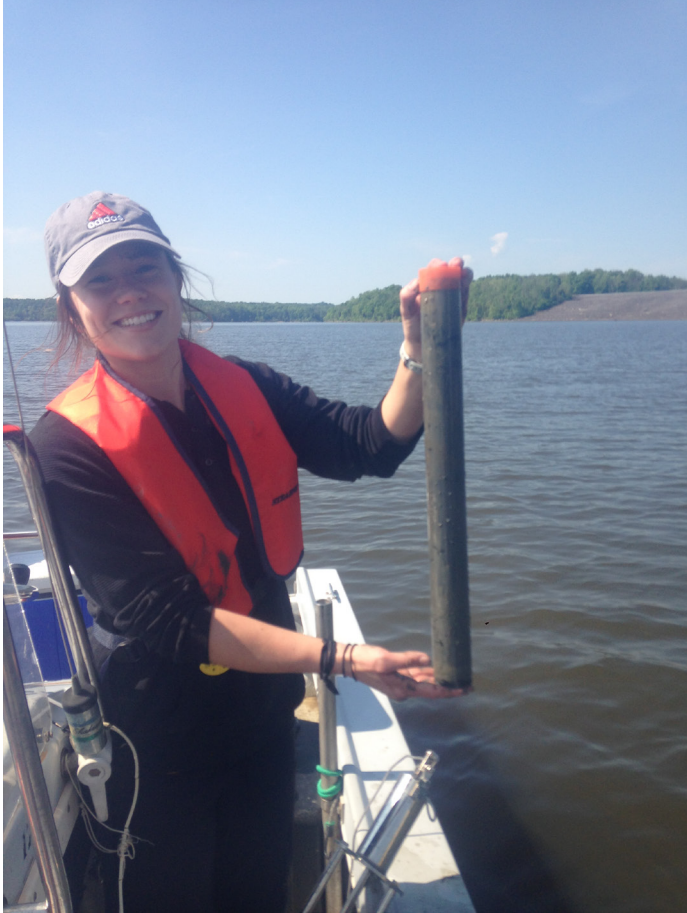
1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.
3. Describe the variations in **abundance** and **taxonomy** of **methanogen communities**.

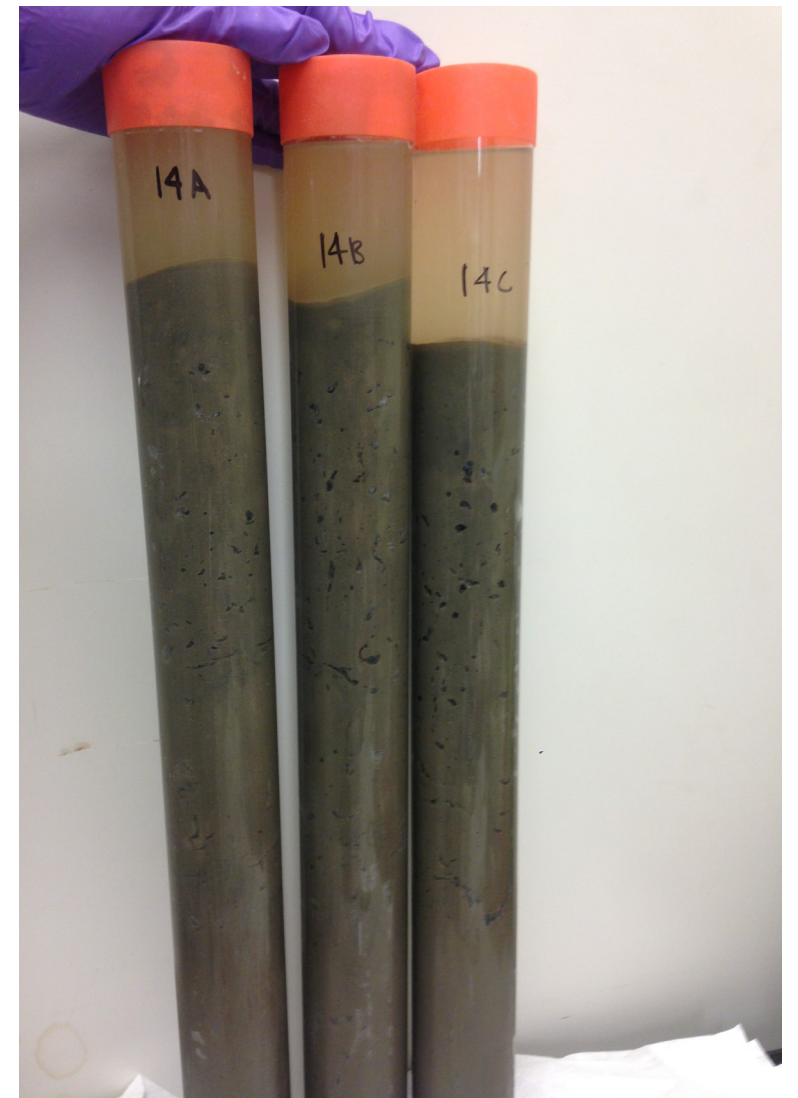
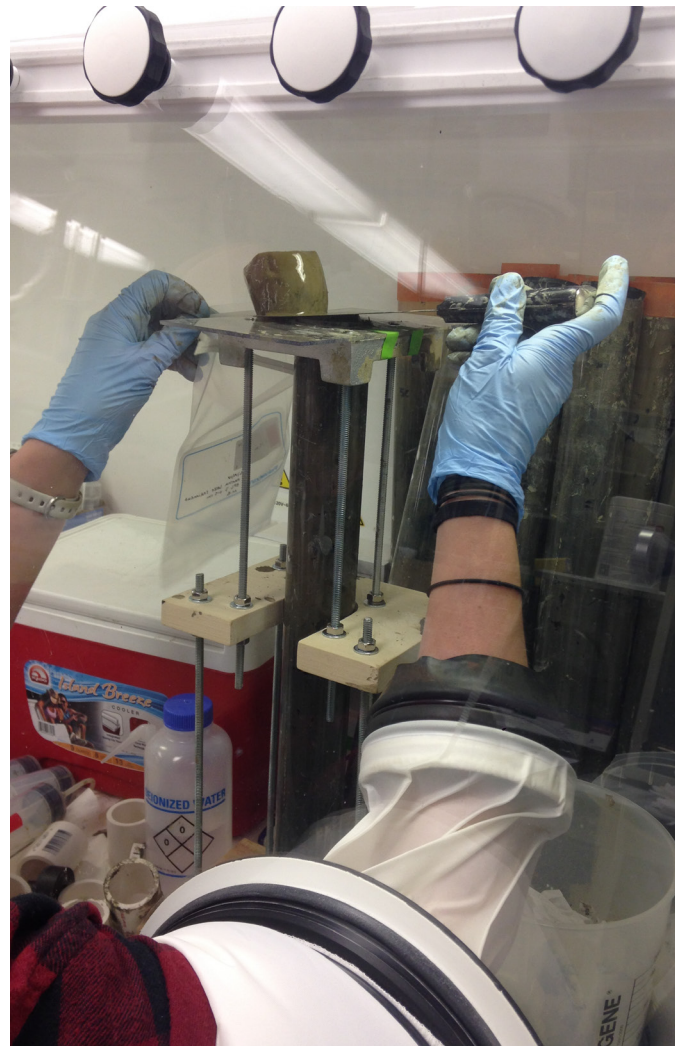
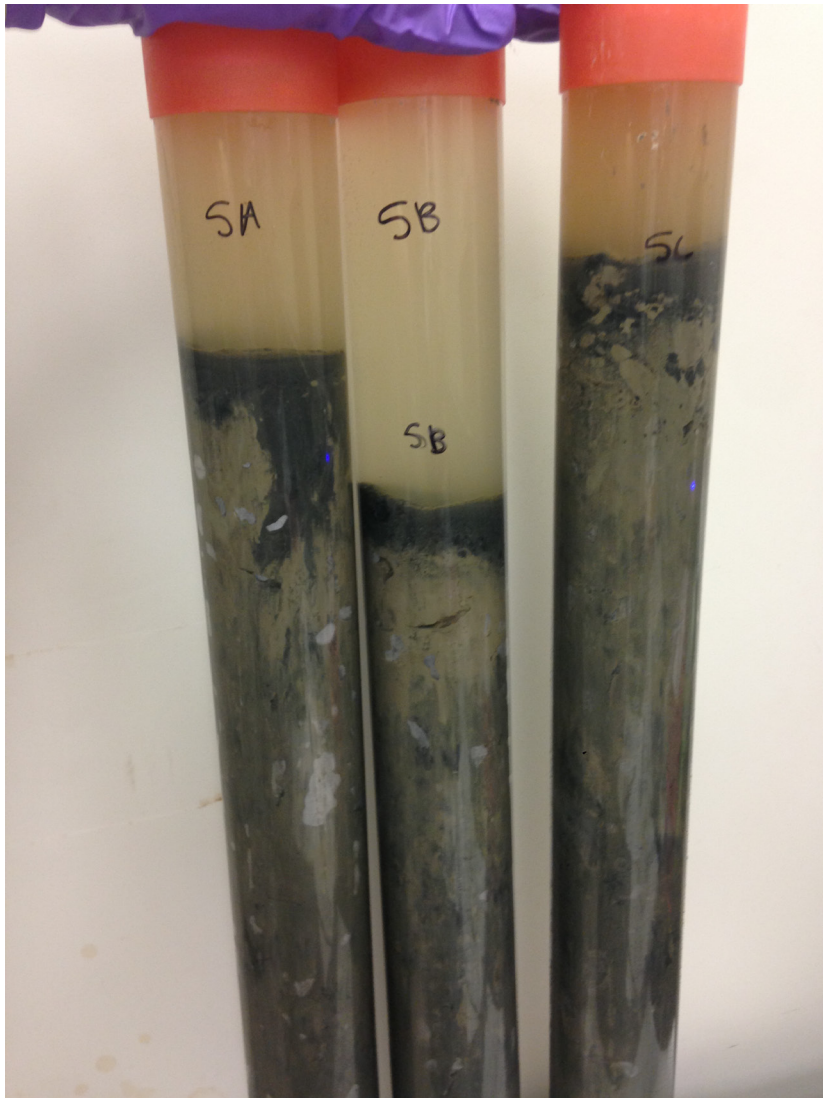
Study Site

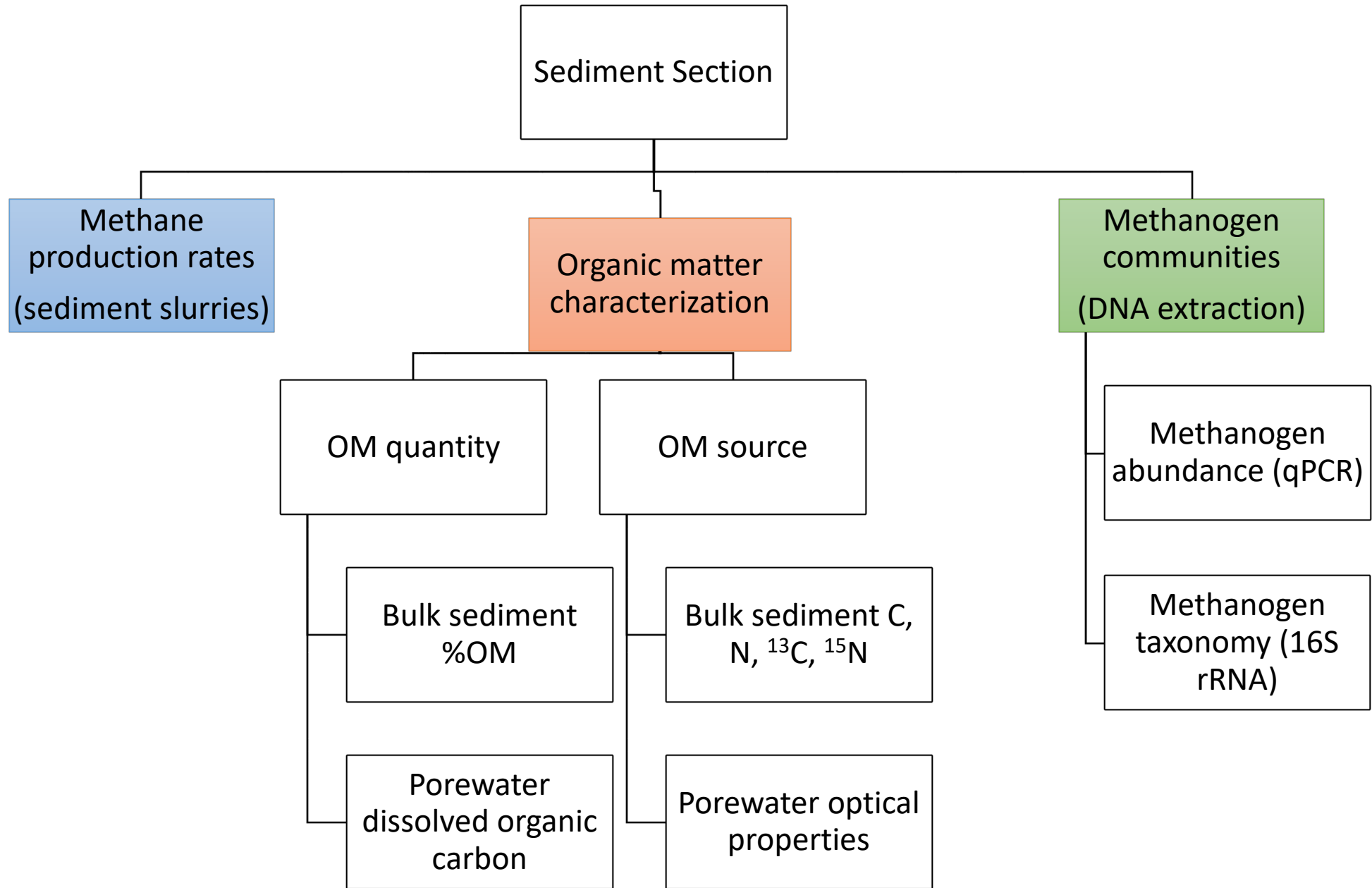
William H. Harsha (“East Fork”) Lake







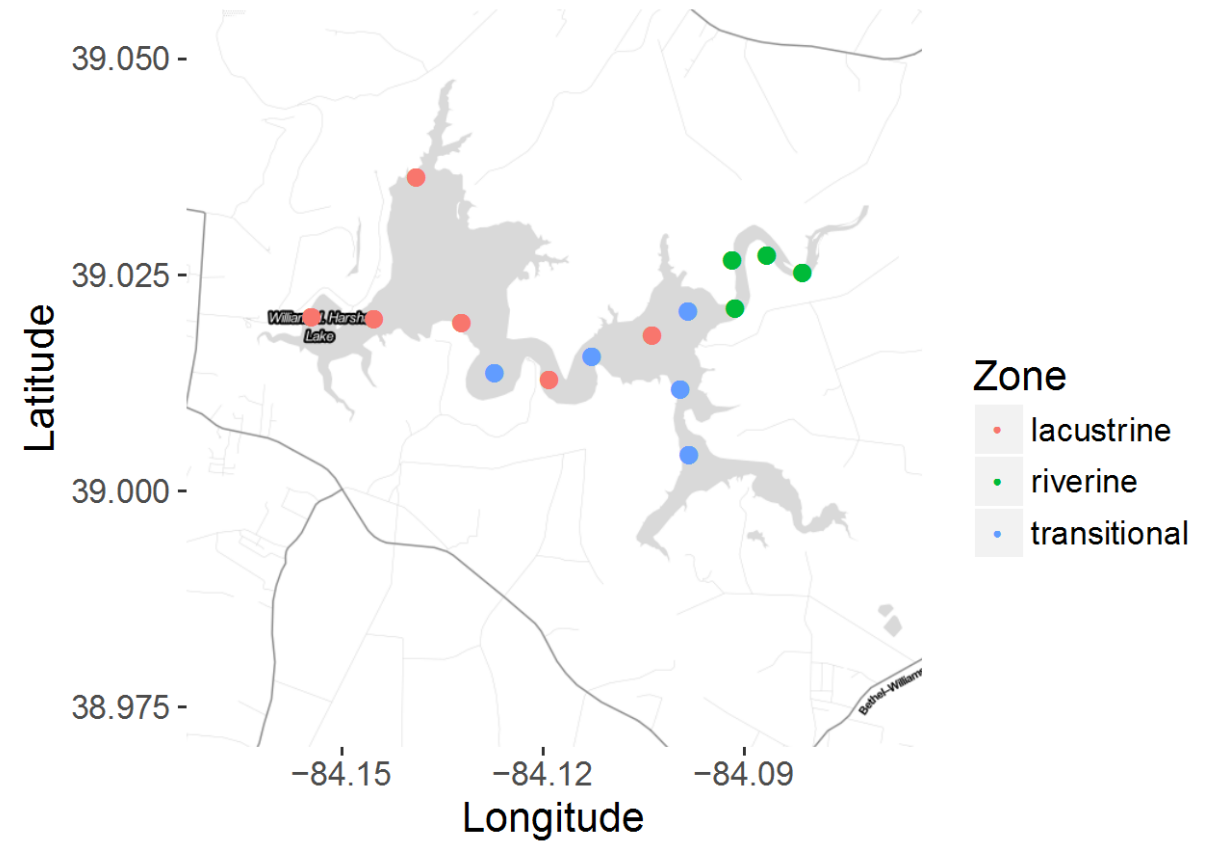
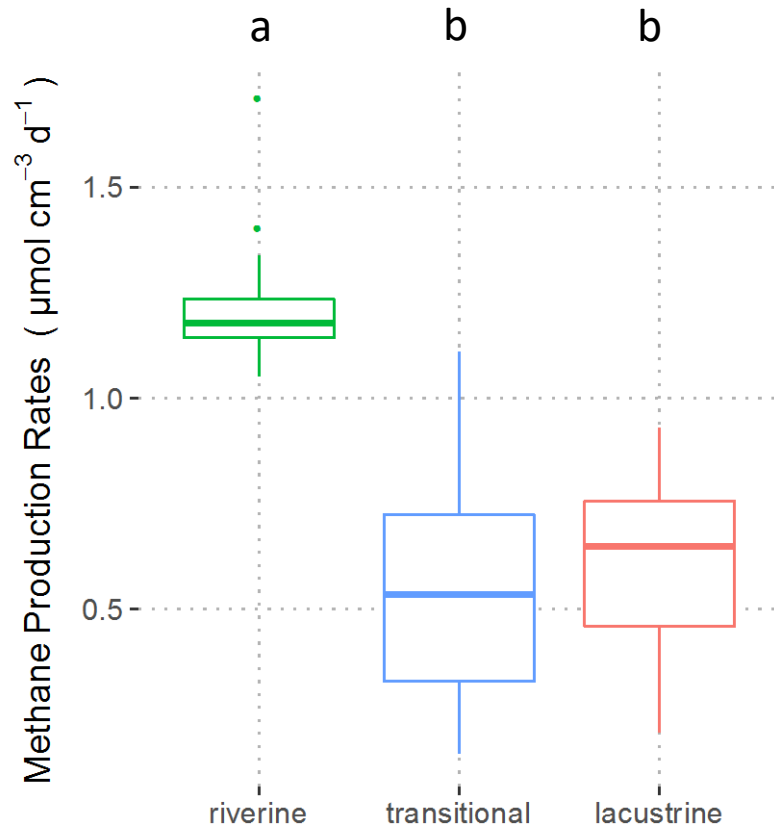




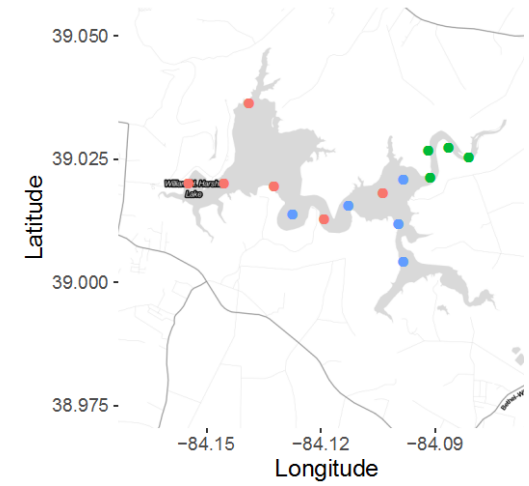
Objectives

1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.
3. Describe the variations in **abundance** and **taxonomy** of **methanogen communities**.

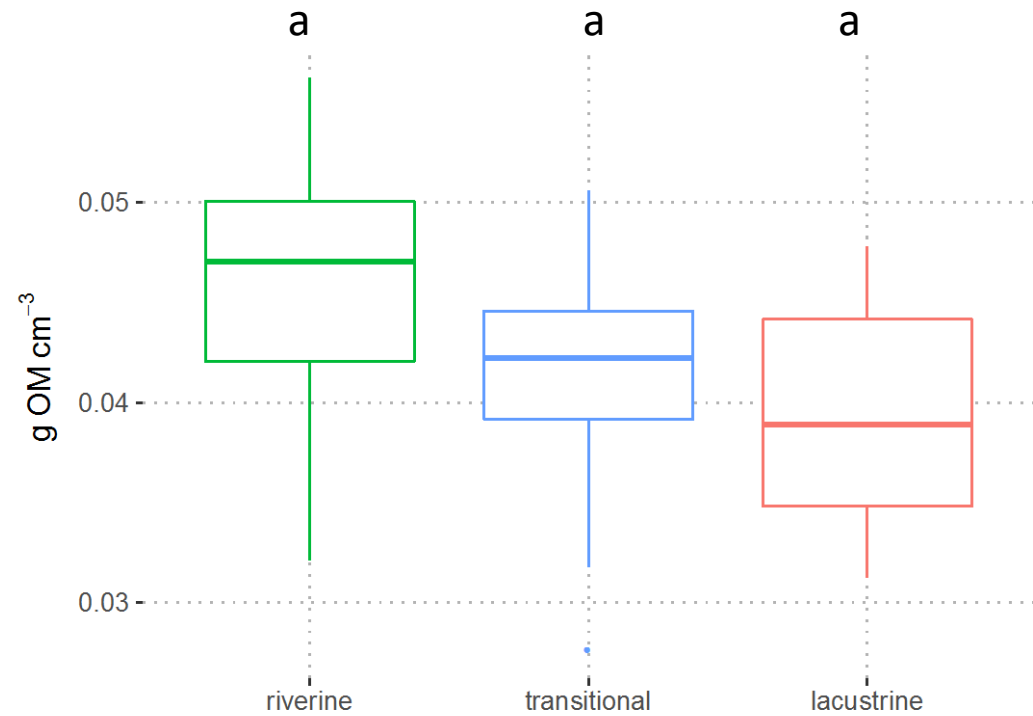
Spatial variation: Methane production rates



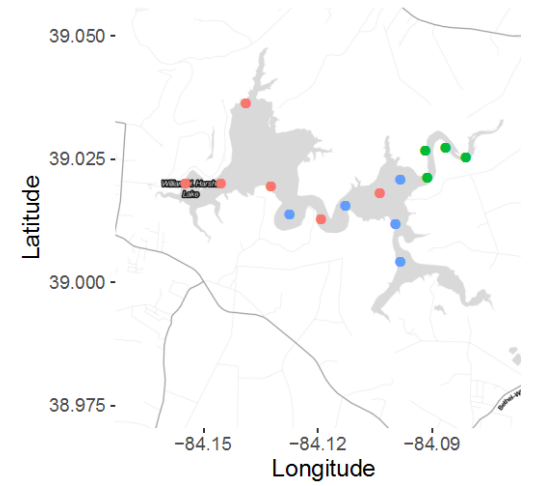
Spatial variation: Organic matter **quantity**



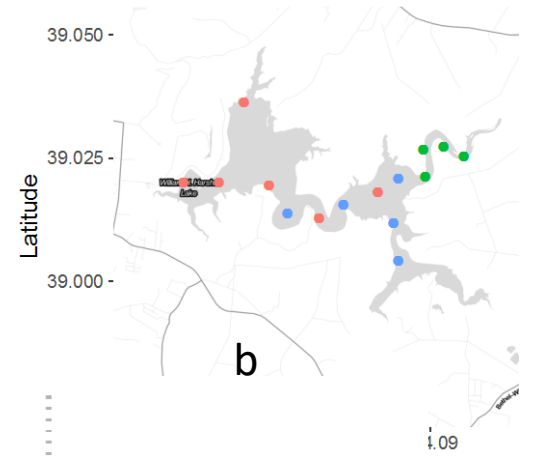
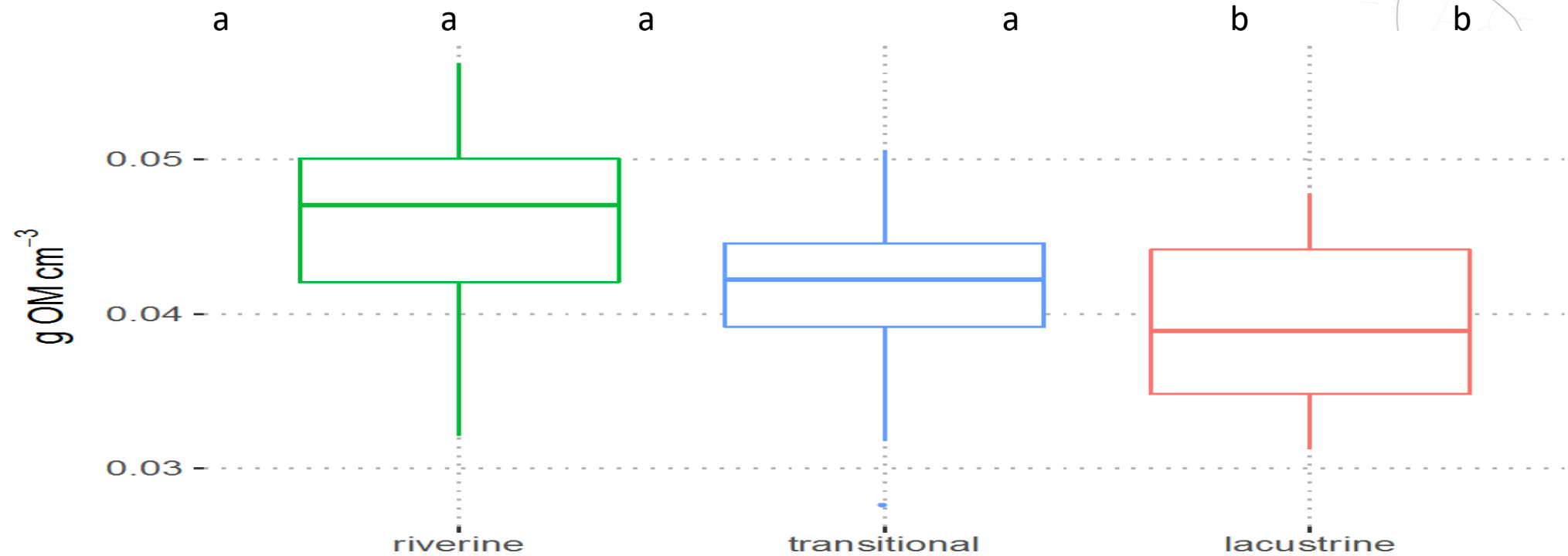
Spatial variation: Organic matter quantity



BULK SEDIMENT fraction of OM



Spatial variation: Organic matter quantity

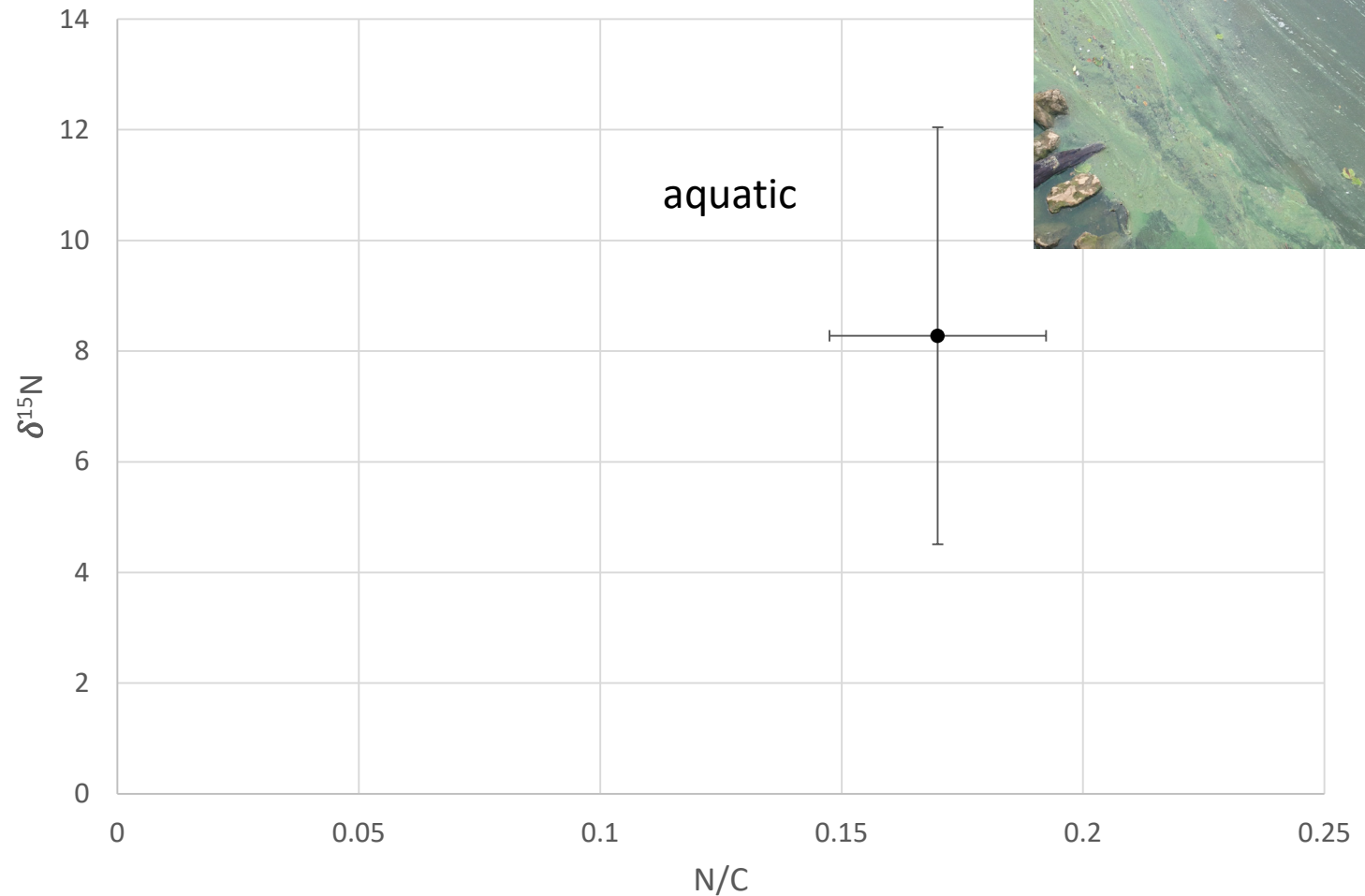


BULK SEDIMENT fraction of OM

DISSOLVED fraction of OM

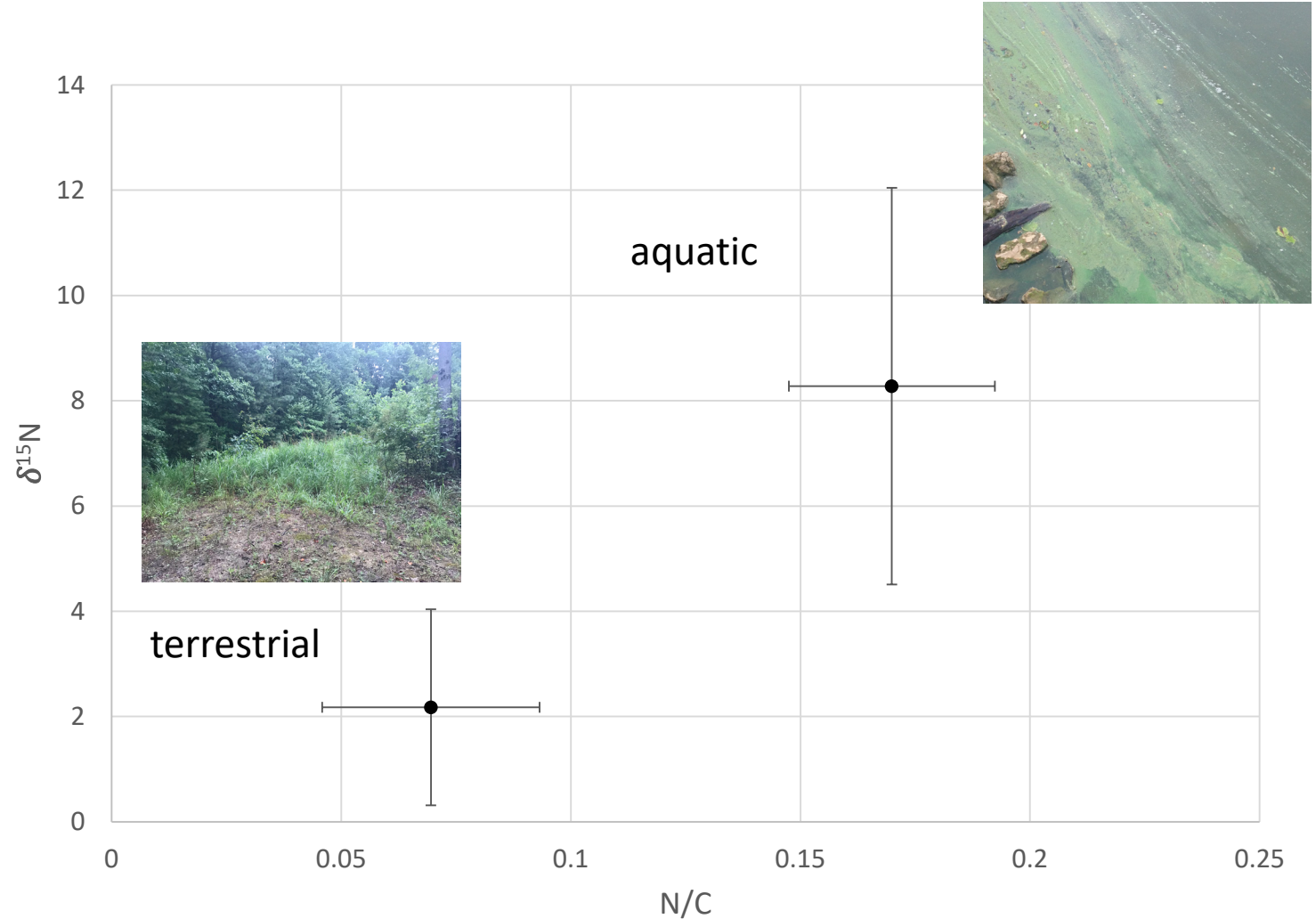
Spatial variation: Organic matter **source**

Mixing model to estimate the proportion of sediment OM that is from algal-derived sources

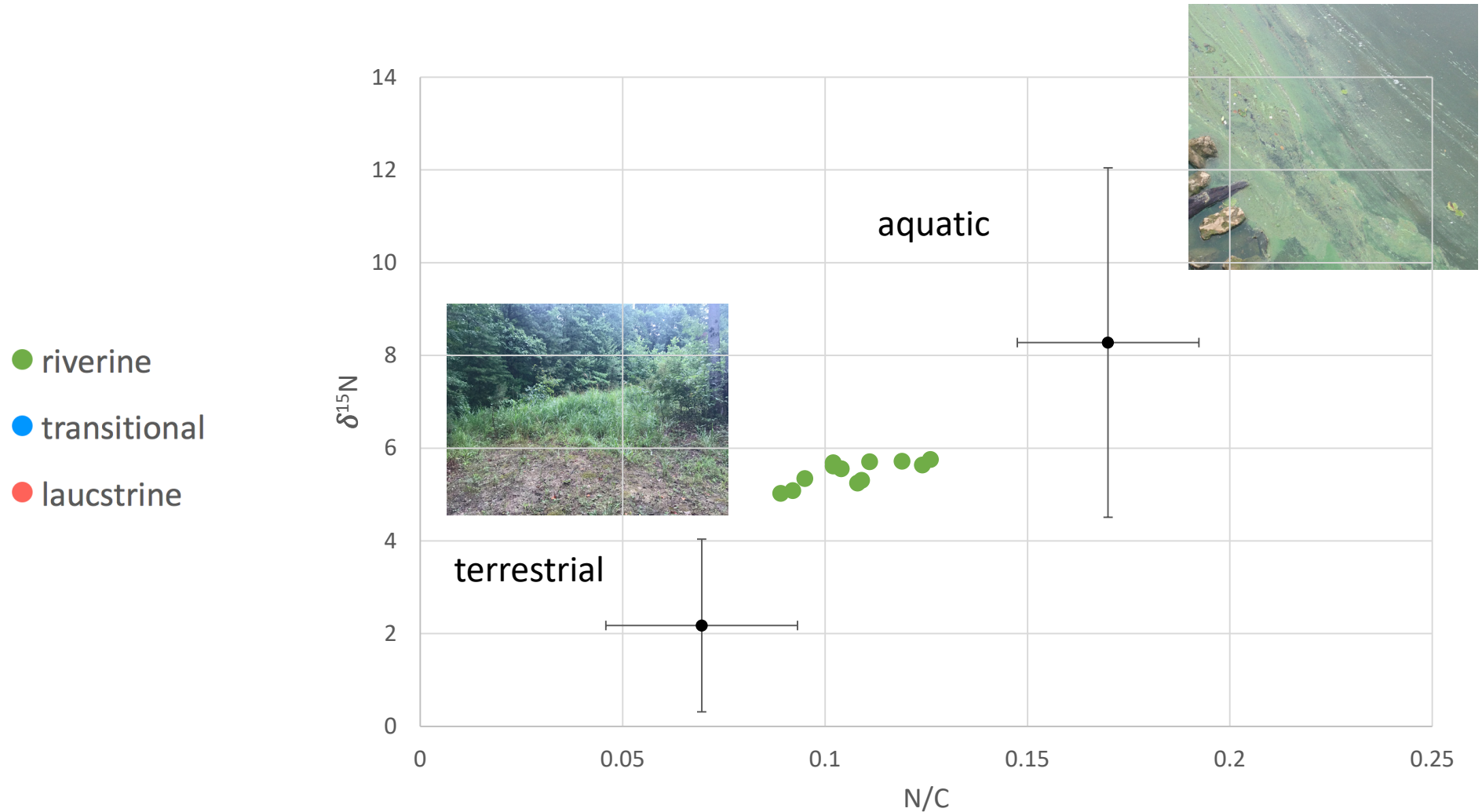


Spatial variation: Organic matter **source**

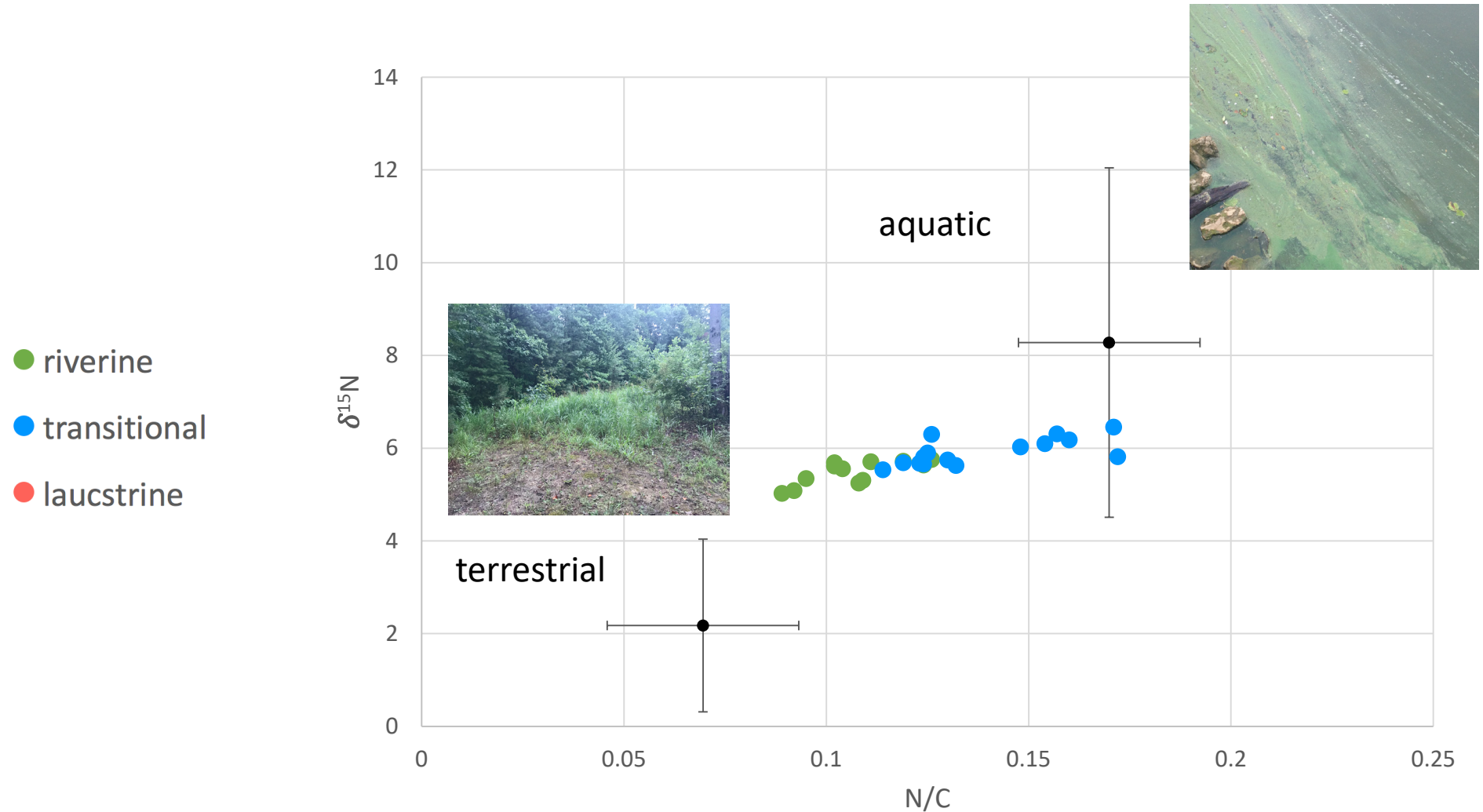
Mixing model to estimate the proportion of sediment OM that is from algal-derived sources



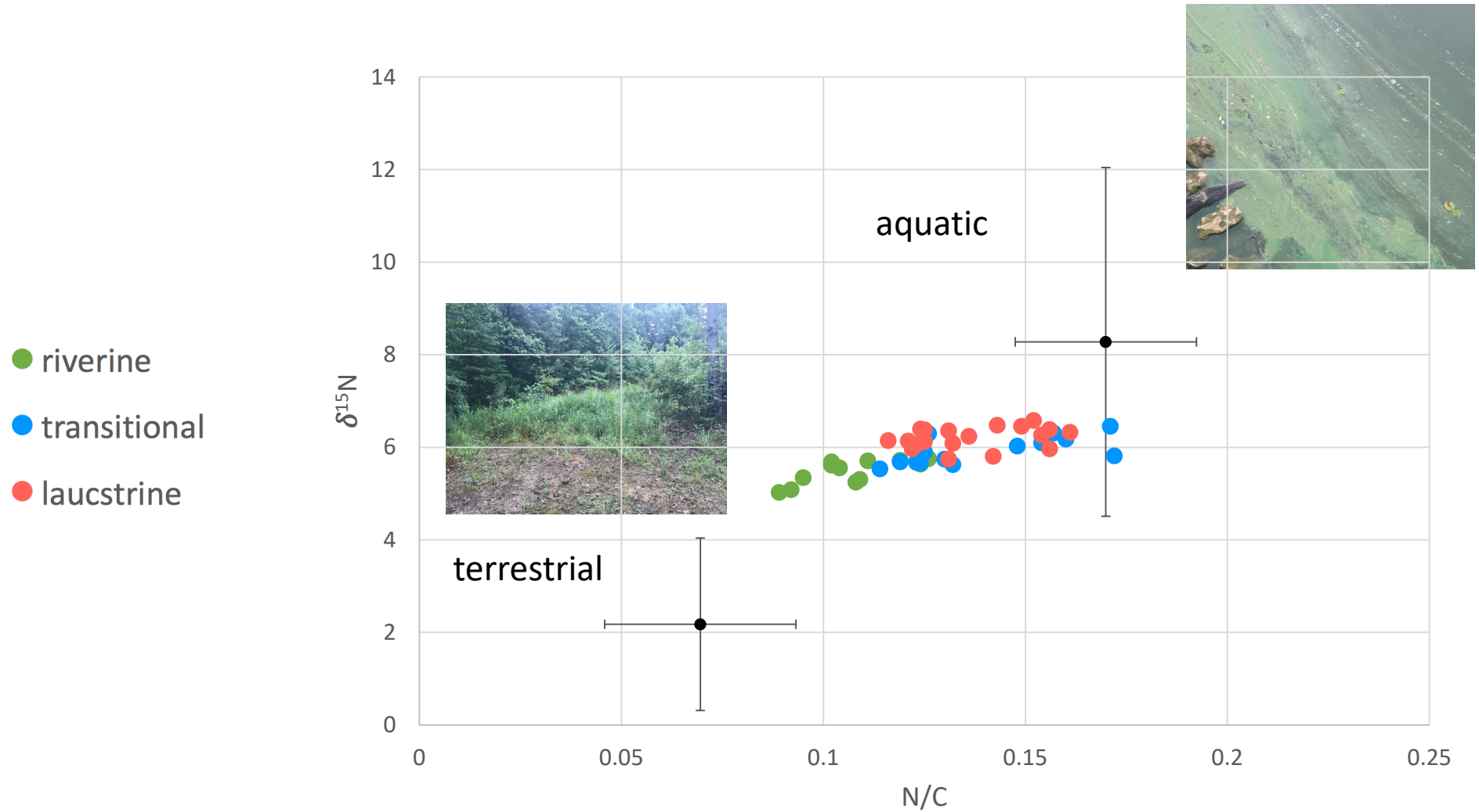
Spatial variation: Organic matter **source**



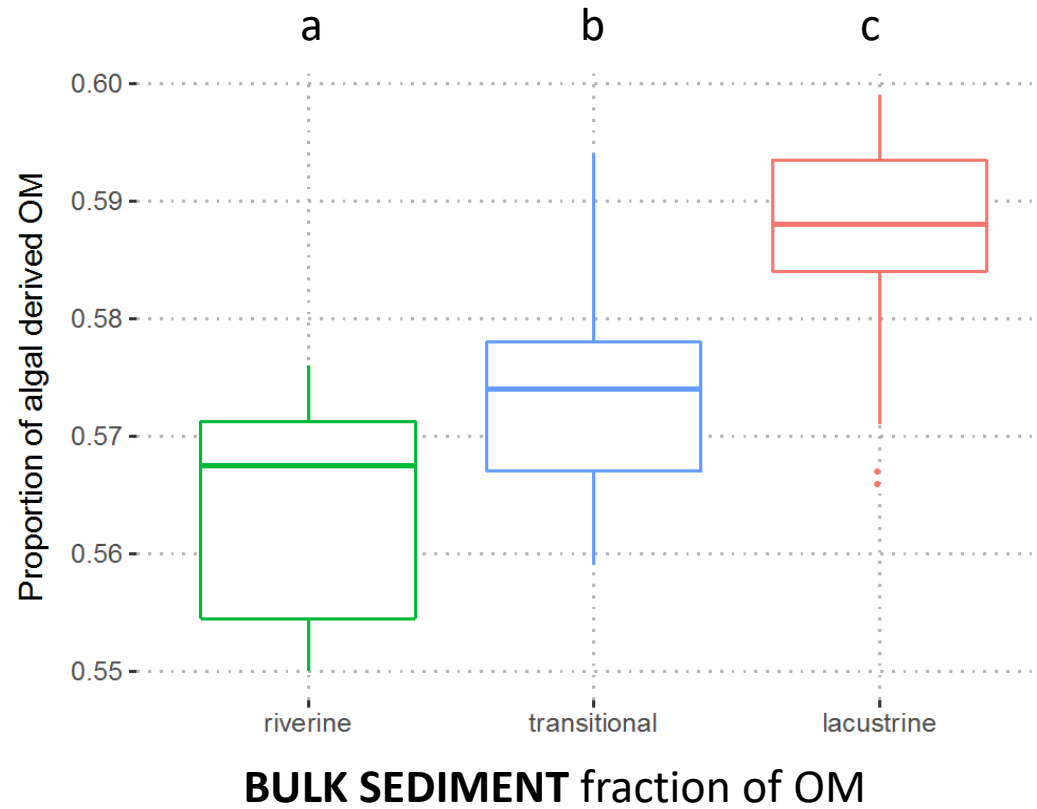
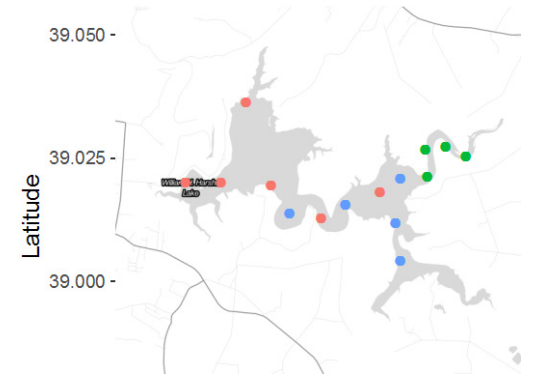
Spatial variation: Organic matter **source**



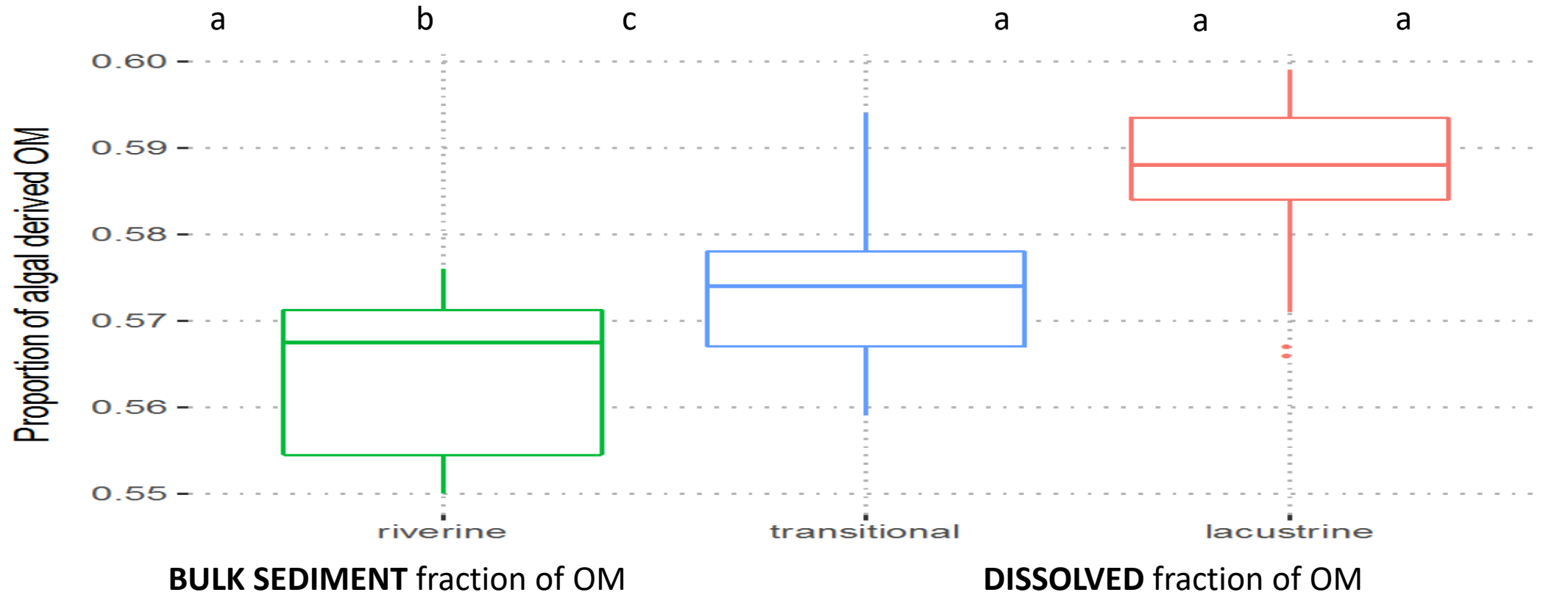
Spatial variation: Organic matter **source**



Spatial variation: Organic matter **source**



Spatial variation: Organic matter **source**



Objectives

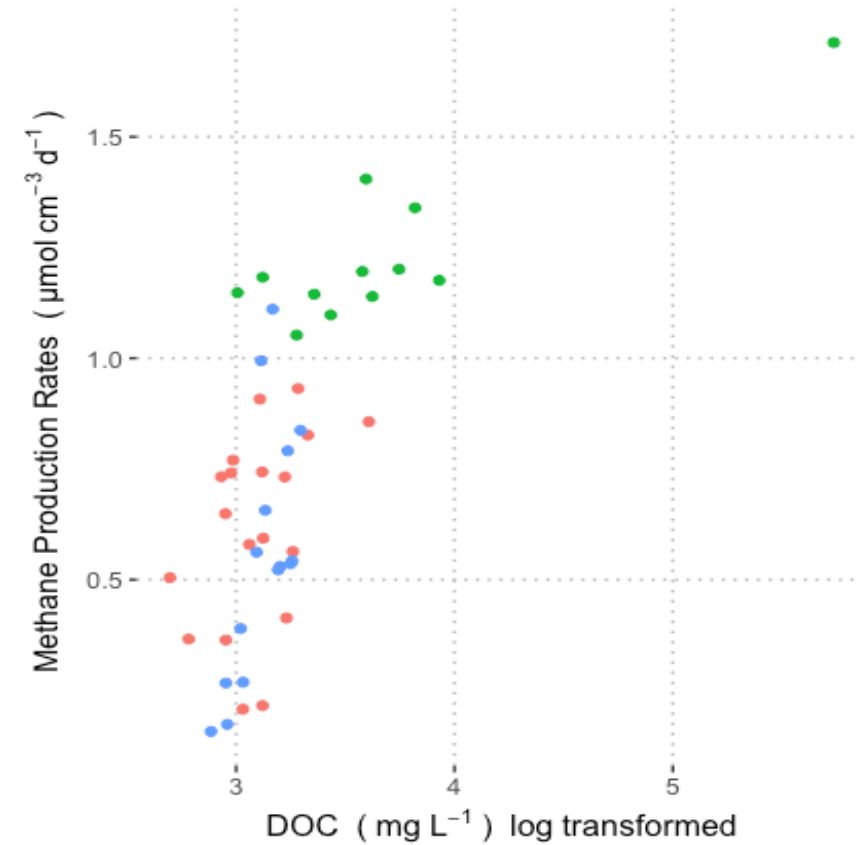
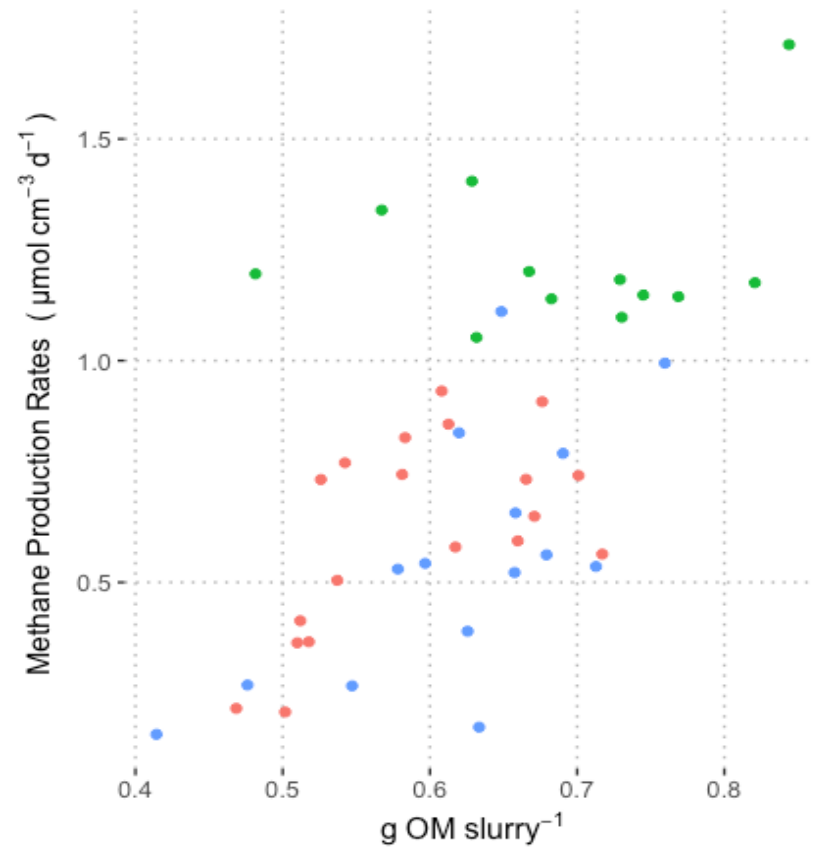
1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.
3. Describe the variations in **abundance** and **taxonomy** of **methanogen communities**.

H1: Organic matter **quantity** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.48
Conditional R ²	0.87

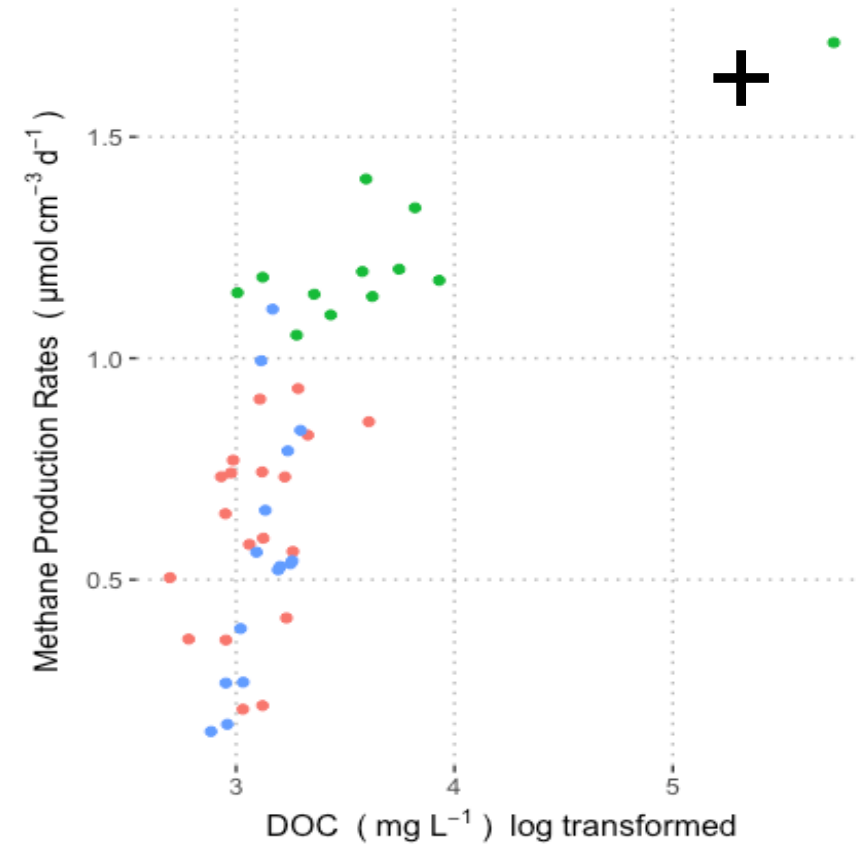
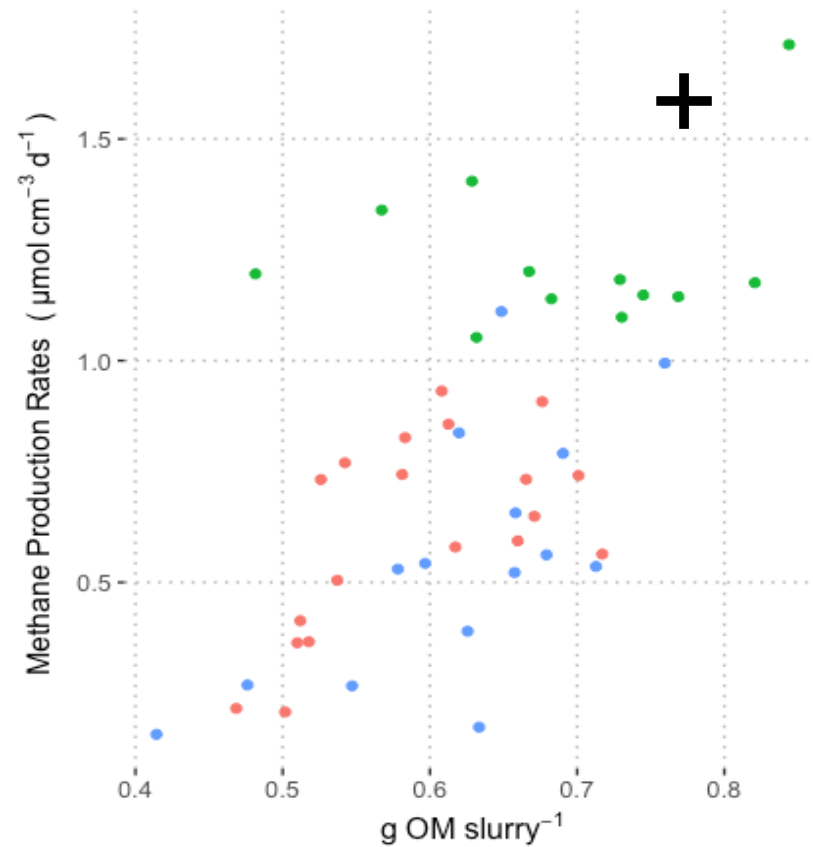
H1: Organic matter **quantity** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.48
Conditional R ²	0.87



H1: Organic matter **quantity** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.48
Conditional R ²	0.87

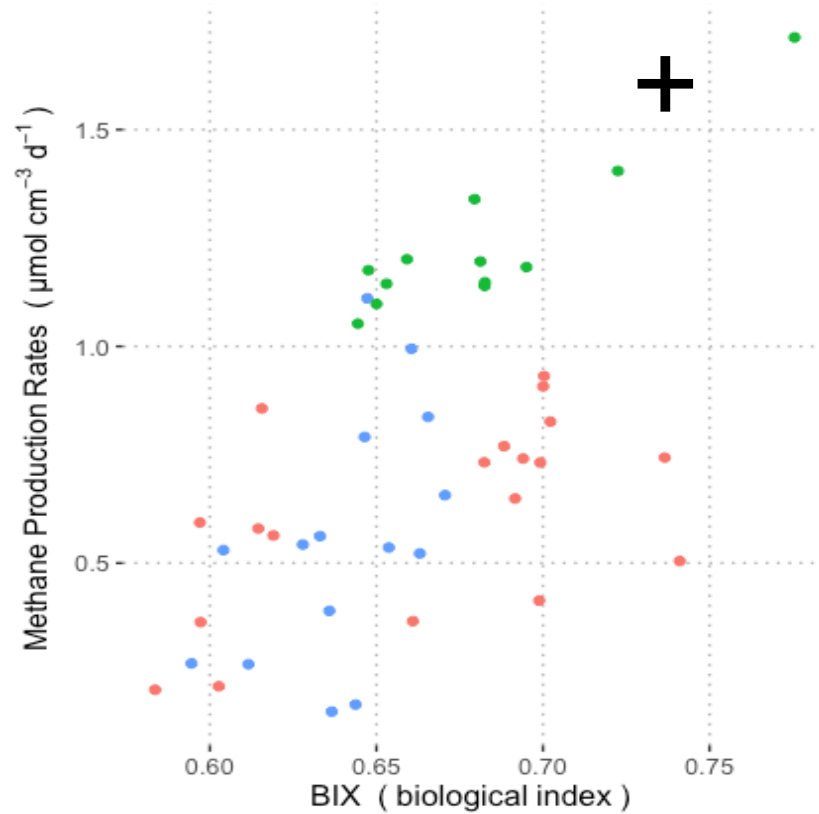


H2: Organic matter **source** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.33
Conditional R ²	0.85

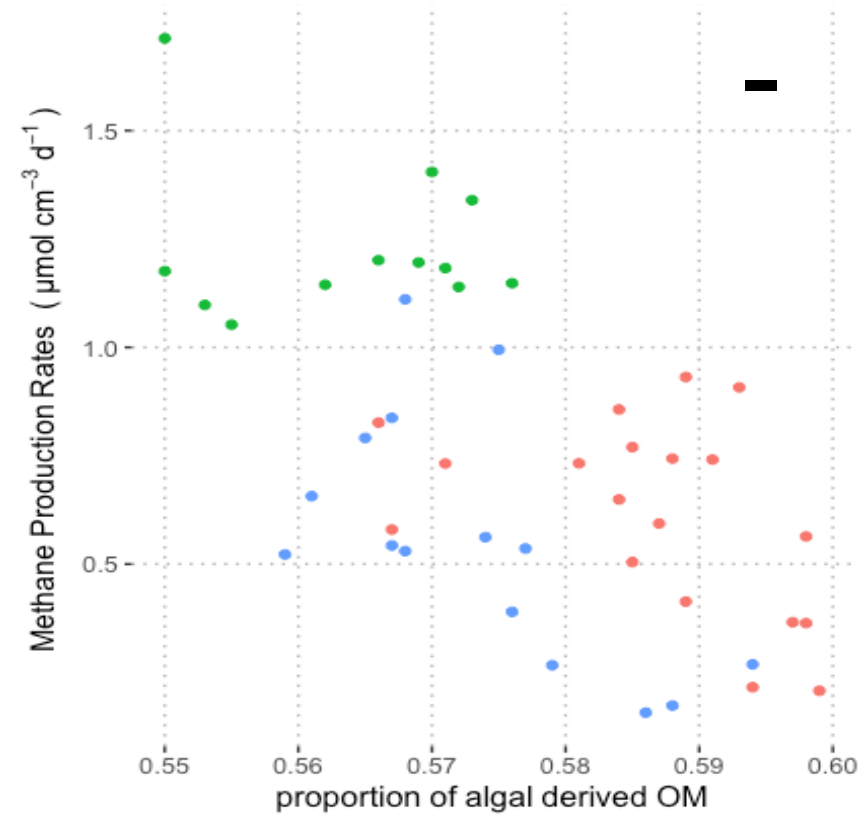
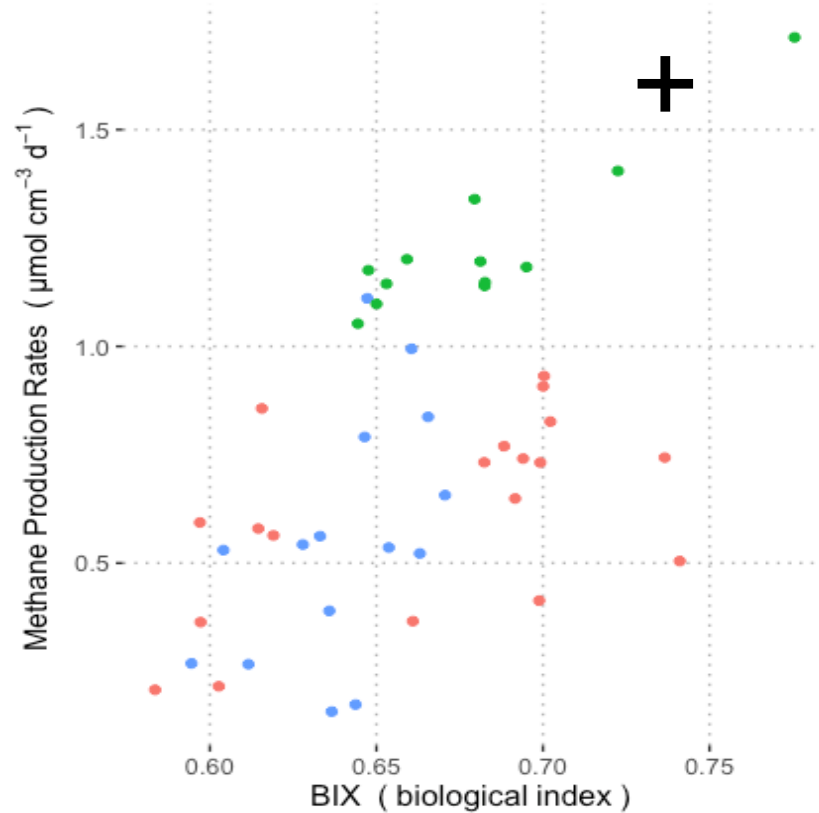
H2: Organic matter **source** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.33
Conditional R ²	0.85



H2: Organic matter **source** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.33
Conditional R ²	0.85



H3: The combination of OM **quantity and source** is the best predictor of methane production rates in Harsha Lake.

Marginal R ²	0.70
Conditional R ²	0.89

Model summaries:

Model	Hypothesis	AICc	Rank	Model probability w_i	Evidence ratio	Marginal R ²	Conditional R ²
H1	quantity	-13.4	3	0.003	370.6	0.48	0.87
H2	source	-15.6	2	0.008	123.8	0.33	0.85
H3	quantity + source	-25.2	1	0.989		0.70	0.89

Objectives

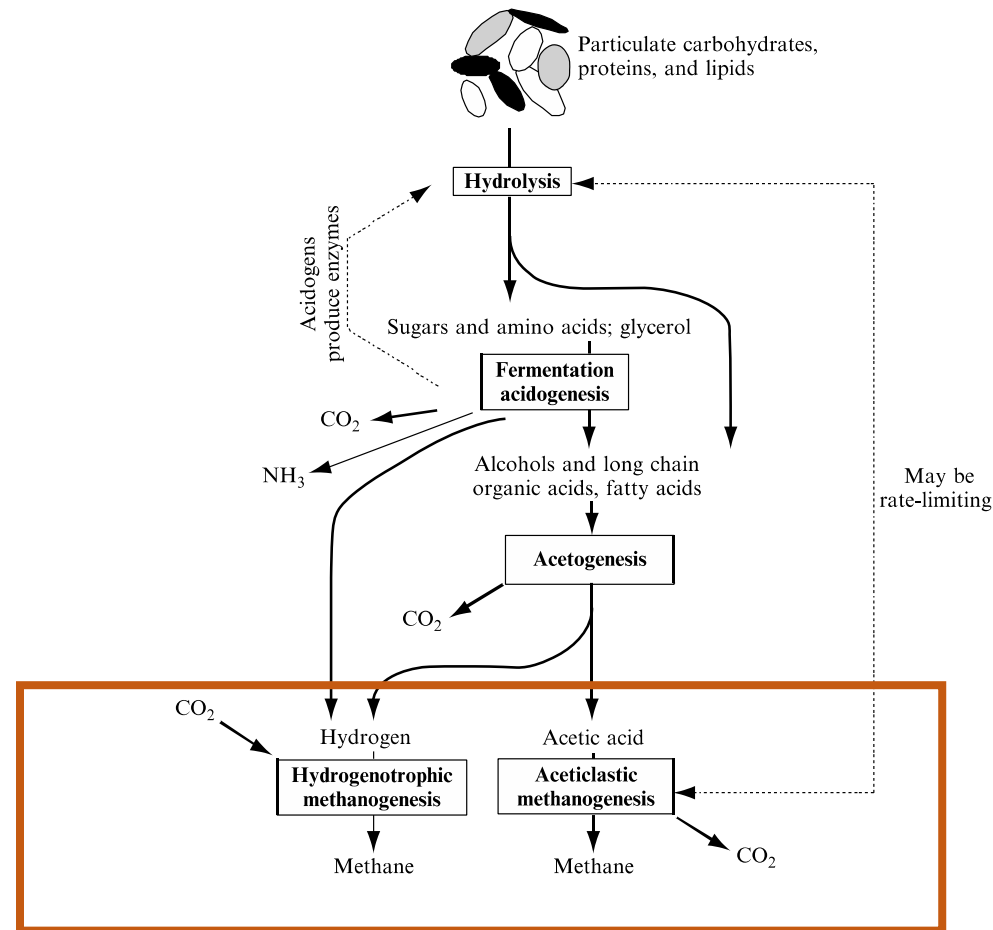
1. Establish if **methane production rates** and **OM source** and **quantity** exhibit distinct variation among reservoir zones.
2. Determine the relative **influence** of **OM source** and **OM quantity** on methane production rates.
3. Describe the variations in **abundance** and **taxonomy** of **methanogen communities**.

Microbial decomposition of OM

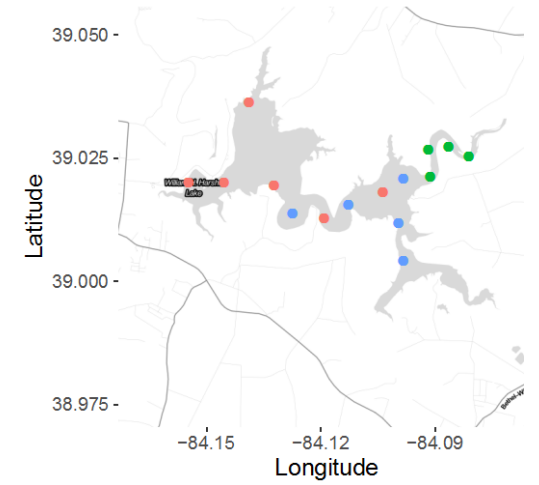
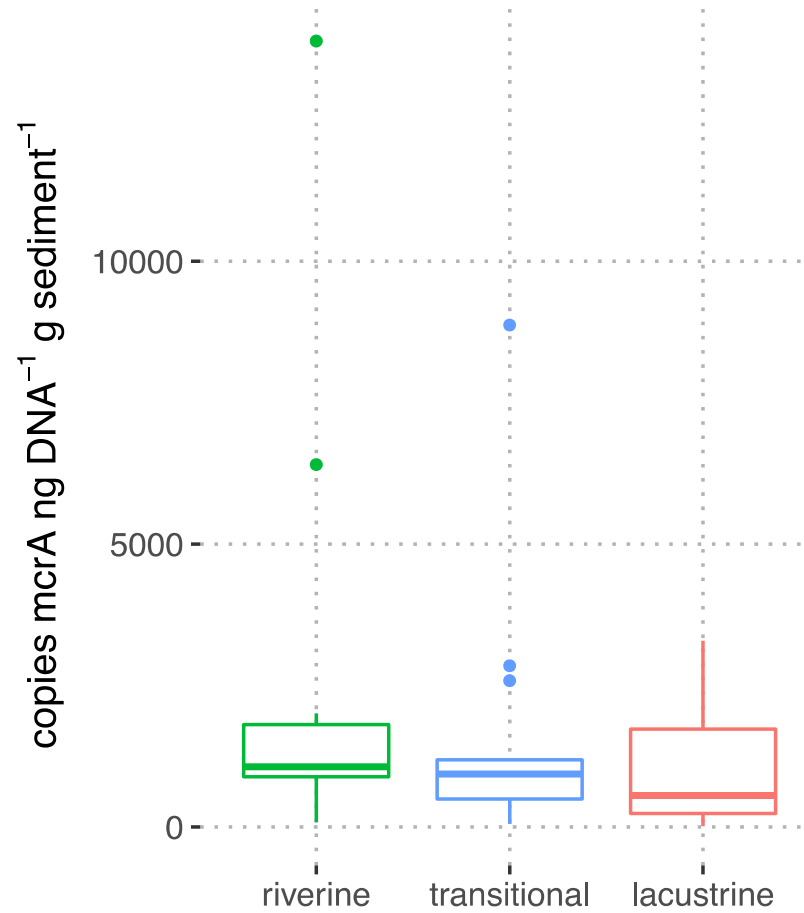
Primary fermenting bacteria

Anaerobic oxidizing bacteria

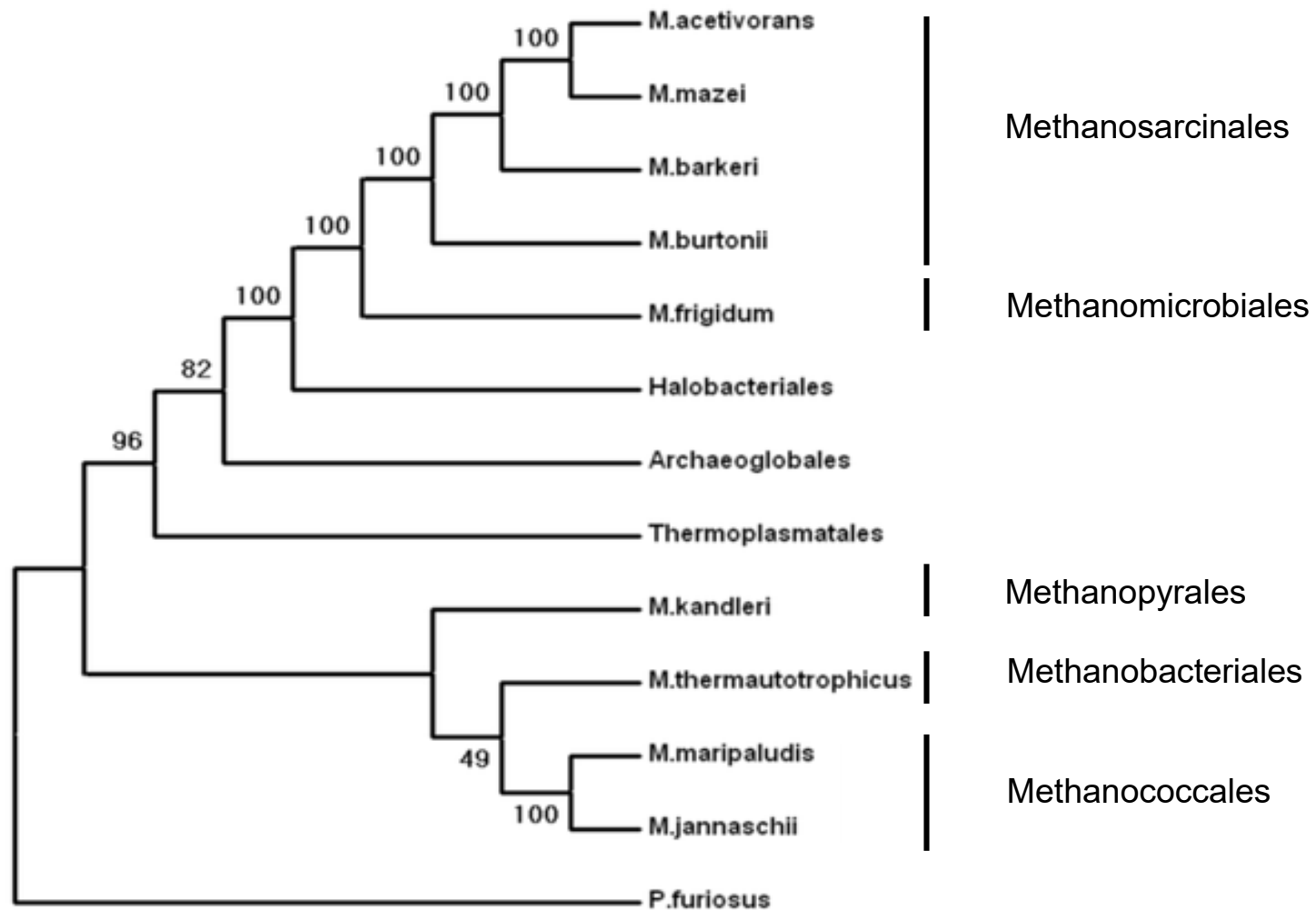
Methanogenic archaea



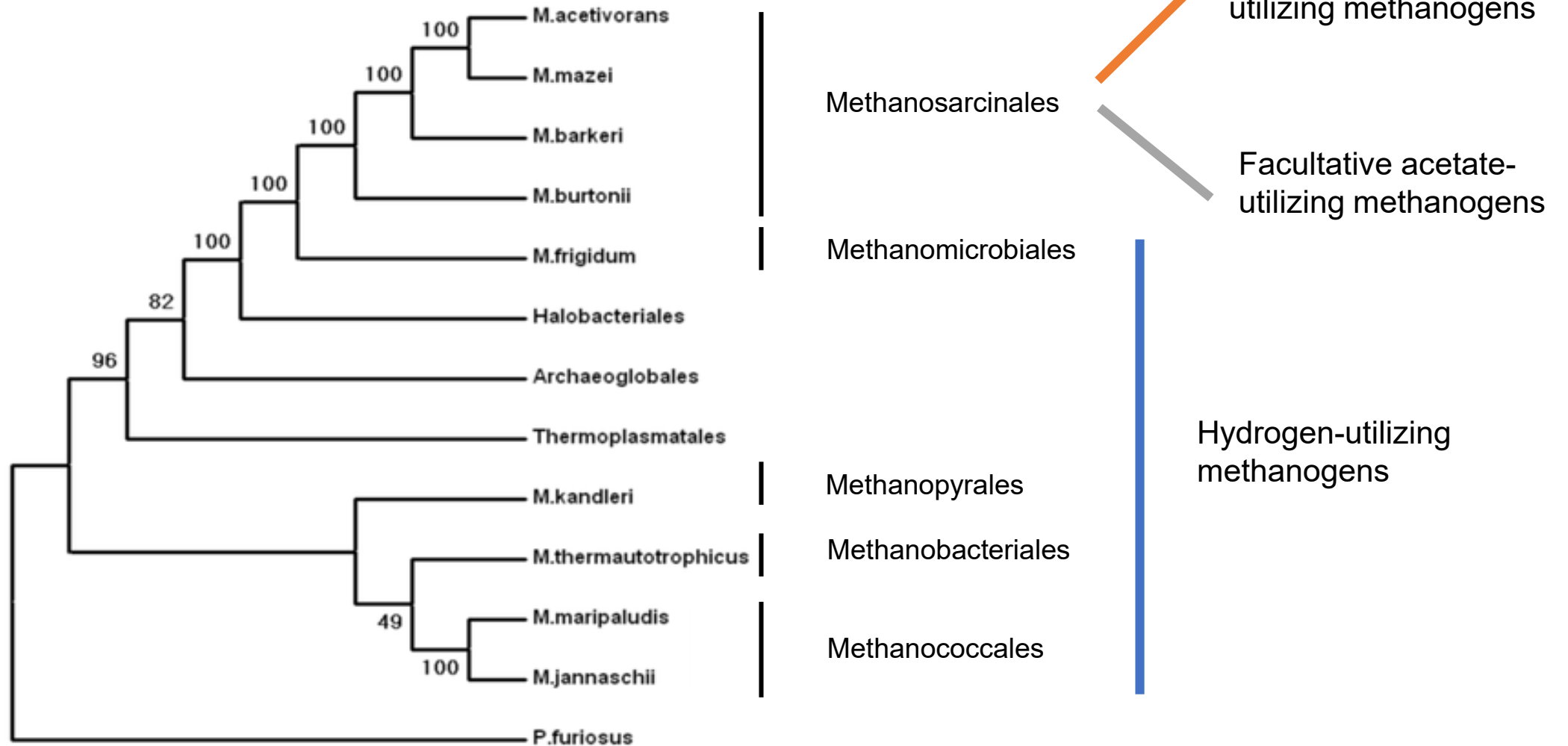
Spatial variation: Methanogen abundance



Methanogen taxonomy



Methanogen taxonomy



Relative abundance of methanogen genera by zone

Methanomicrobiales; Methanoregula	47.7	69.5	66.9
Methanosarcinales; Methanosaeta	19.6	15.2	16.6
Methanobacteriales; Methanobacterium	15.1	9.6	7.3
Methanomicrobiales; Methanolinea	2.8	6.4	4.8
Methanosarcinales; Methanosarcina	15.4	0.5	0.4
Methanomicrobiales; Methanoline	1.4	1.7	3.6
Methanomicrobiales; Methanospirillum	4.3	0.8	1.3
Methanocellales; Rice_Cluster_I	0.9	0.1	0.3
Methanomicrobiales; SMS-sludge-7_unclassified	0.3	0.1	0.5
Methanosarcinales; Candidatus_Methanoperedens	0.4	0.8	0
Methanomicrobiales; Methanomicrobiales_unclassified	0	0.4	0.4
Methanosarcinales; Methanolobus	0.8	0	0
Methanomicrobiales; Methanomicrobiales_unclassified	0.4	0.1	0.2
Methanosarcinales; Methanomethylovorans	0.6	0	0
Methanobacteriales; Methanobrevibacter	0.4	0	0
	riverine	transitional	lacustrine

Relative abundance of methanogen genera by zone

Methanomicrobiales; Methanoregula	47.7	69.5	66.9
Methanosarcinales; Methanosaeta	19.6	15.2	16.6
Methanobacteriales; Methanobacterium	15.1	9.6	7.3
Methanomicrobiales; Methanolinea	2.8	6.4	4.8
Methanosarcinales; Methanosarcina	15.4	0.5	0.4
Methanomicrobiales; Methanoline	1.4	1.7	3.6
Methanomicrobiales; Methanospirillum	4.3	0.8	1.3
Methanocellales; Rice_Cluster_I	0.9	0.1	0.3
Methanomicrobiales; SMS-sludge-7_unclassified	0.3	0.1	0.5
Methanosarcinales; Candidatus_Methanoperedens	0.4	0.8	0
Methanomicrobiales; Methanomicrobiales_unclassified	0	0.4	0.4
Methanosarcinales; Methanolobus	0.8	0	0
Methanomicrobiales; Methanomicrobiales_unclassified	0.4	0.1	0.2
Methanosarcinales; Methanomethylovorans	0.6	0	0
Methanobacteriales; Methanobrevibacter	0.4	0	0
	riverine	transitional	lacustrine

Conclusions

- The riverine zone was different in its methane production rates, sediment characteristics, and methanogen composition
- Quantity and source of OM combined better predict methane production rates than either quantity or source alone
- The proportion of algal-derived OM in the bulk sediment was negatively correlated to methane production rates.

Acknowledgements



Advisor

Dr. Ishi Buffam

Committee

Dr. Jake Beaulieu

Dr. Trinity Hamilton

Lab Members

Jeremy Alberts

Mark Mitchell

Alicia Goldschmidt

Chelsea Hintz

Sarah Handlon

Undergraduates

Madison Duke

Kaitlin Henn

Caroline Tran

Field Crew

Kit Daniels

Madison Duke

Dr. Sarah Waldo

Dr. Xuan Li

Dr. Joel Allen

EPA Help

Karen White

Dr. Mike Elovitz

Dr. Aabir Banerji

Funding

EPA Traineeship Program

Ohio WRC Grant

Weiman/Benedict

WISE Program

ASLO travel award

