

INTRODUCTION

Outbreak of waterborne disease is a worldwide water quality issue, waterborne diseases cause health, social, and economic problems (Grabow, 1996).

The application of dynamic hydrologic models to simulate the pathogen loads and removal in varied aquatic ecosystems is still limited.

Rationale:

Human development has a significant impact on water quality, and quantifying this impact is critical.

Dynamic watershed-scale models are needed to understand the sources, transport and fate of pathogens and the consequences for water quality at broad regional scales and how they vary over time.

Research questions:

Are stream networks important regulators of fecal coliform transfer from source areas to critical water bodies?

Hypothesis:

We hypothesize that pathogen concentrations entering river systems from land are positively correlated with developed land use and hydrologic characteristics. Stream networks play an important role of the fate and transport of pathogen. The varied hydrologic conditions cause the difference of pathogen decay and lead to the variations of pathogen removal among aquatic ecosystems.

OBJECTIVES

To address the overarching question we will use a spatially distributed modeling approach that accounts for the location of sources, routing, and in-stream transformations.



Fig 1. Sampling points (Oyster River) and modeled domain (Lamprey River)



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Modeling the fate and transport of fecal coliform in Lamprey River Watershed

Tao Huang¹, Wilfred M Wollheim ^{1,2}, and Robert J Stewart ² ¹Department of Natural Resources and the Environment, University of New Hampshire, Durham, NH 03824, USA ²Institute for the Study of Earth, Oceans, and Space, University of New Hampshire, Durham, NH 03824, USA



Fig 7. Proportion of watershed aquatic removal



FC removal (#/yr) Value 2.41e+011 5.80e+008 5.5 2.75 0 5.5 Kilometer

Fig 8. Fecal coliform removal in the river network

SUMMARY AND FUTURE WORK

This study applied FrAMES model to simulate the fate and transport of the pathogen indicator- fecal coliform.

River networks have the ability to remove fecal coliform.

Future works include the prediction of fecal coliform concentration under future land use and climate change scenarios.

REFERENCES

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