

Biogeography of Atlantic Salmon (*Salmo salar*) in the Cape Breton Highlands

Background

Atlantic Salmon

- Anadromous salmonid; historically ubiquitous across the North Atlantic
- **Severe declines** in recent decades^{1,2}
- Threats to salmon:
 - climate change, overfishing, spawning habitat degradation, barriers to migration (i.e. dams) and genetic introgression from farmed fish³
- High degree of homing; **geographically distinct populations**; local adaptation to their specific region⁴



Figure 1 Atlantic salmon parr

Designatable Units (DUs)

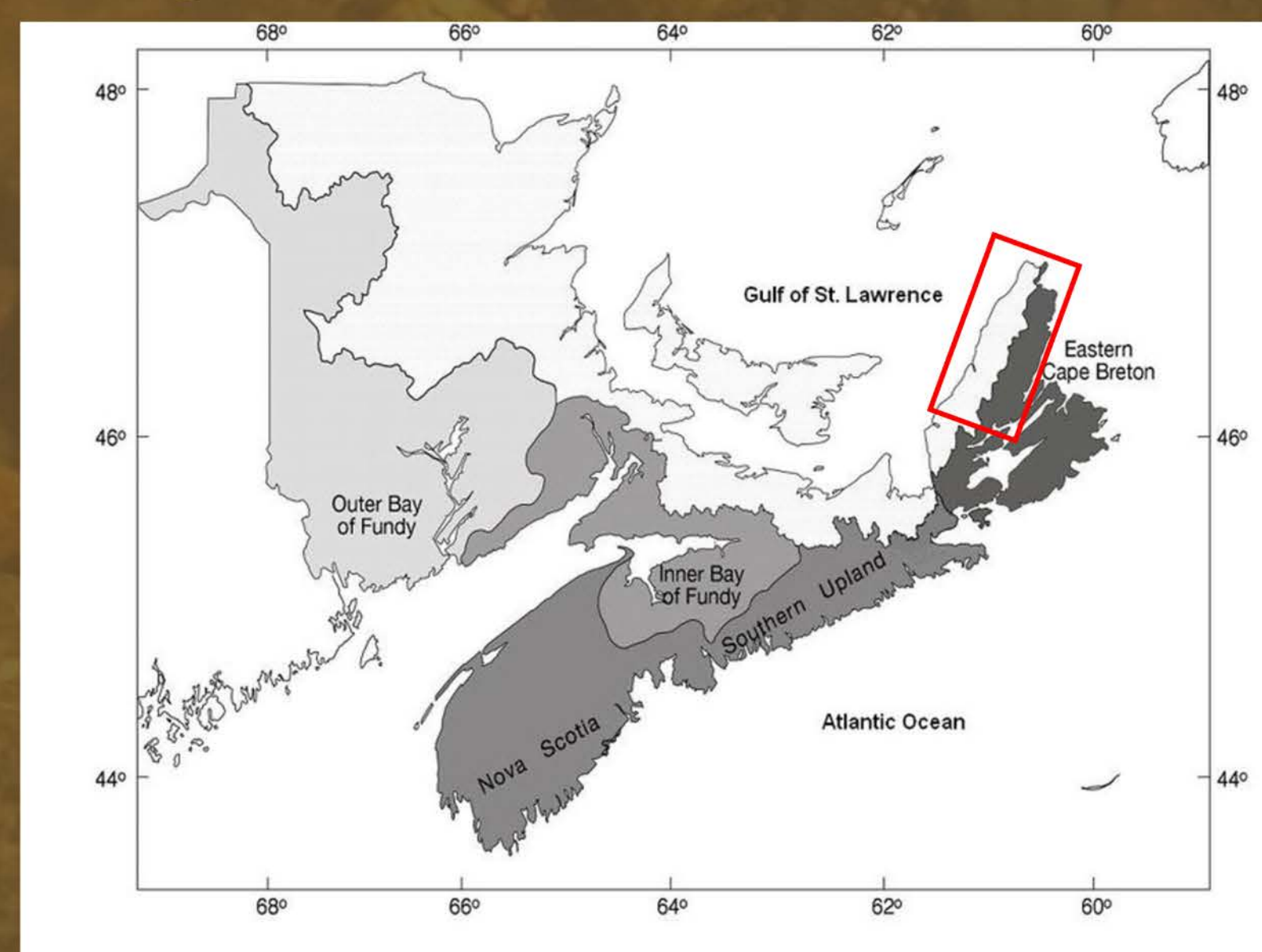


Figure 2 Map of the COSEWIC designatable units in Atlantic Canada. Photo Credit: Heather Bowlby

- 2010: COSEWIC divides salmon populations into 16 DUs to recognize and preserve the diversity within wild subpopulations⁵
- Salmon management differs between DUs depending on the local requirements of the population⁵

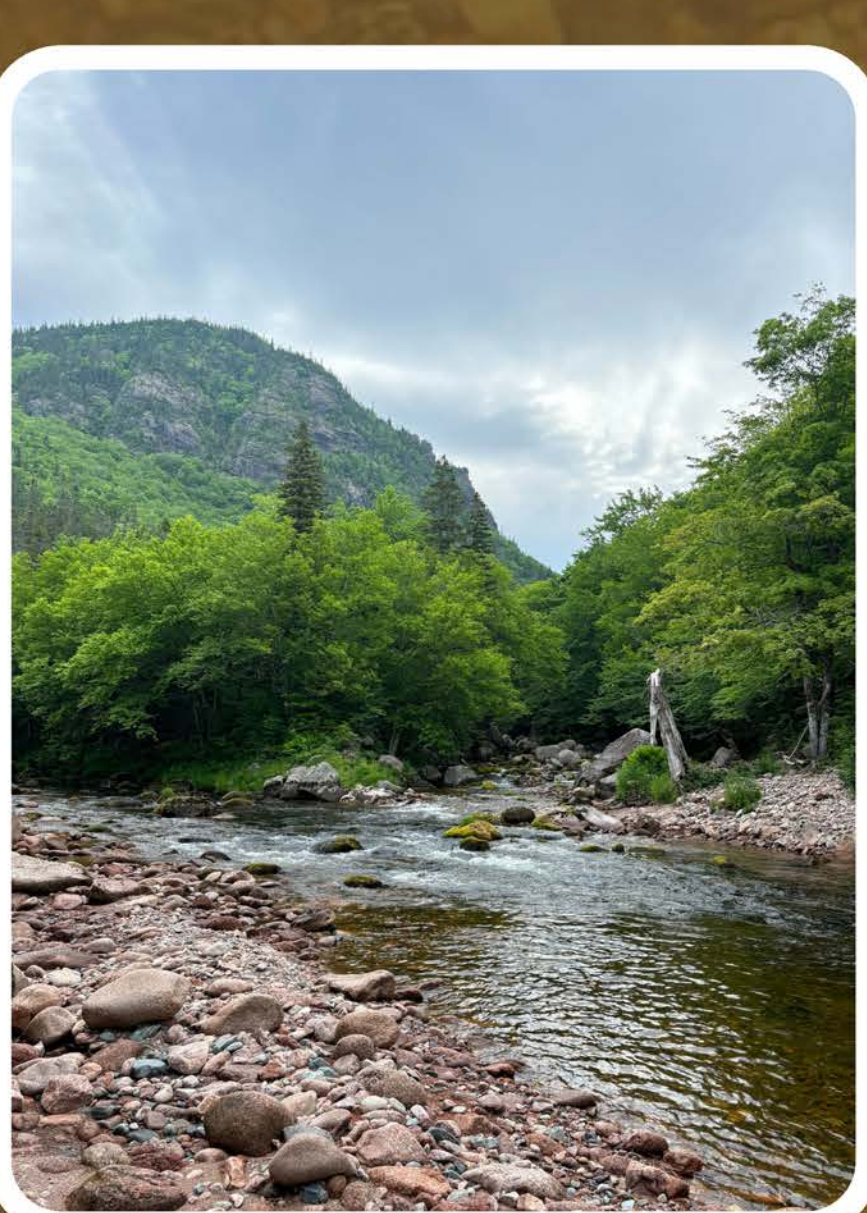


Figure 3 A river in the Highlands

The Cape Breton Highlands DUs:

- Gulf of St. Lawrence Unit:**
Conservation Status: Special Concern
- Eastern Cape Breton Unit:**
Conservation Status: Endangered⁵

The Cape Breton Highlands

- An **understudied** region in Atlantic Canada; data limited to to inaccessibility
- The exact number of rivers inhabited by salmon in the Highlands is unknown; existing distribution estimates are outdated^{6,7}
- **Effective conservation** requires **accurate data** on the distribution and abundance of wild salmon populations

Study Objectives

- 1 Update the current distribution data on Atlantic salmon in the Cape Breton Highlands
- 2 Compare the distribution and abundance of salmon between Designatable Units
- 3 Compare the species community assemblages between rivers in both Designatable Units

Results

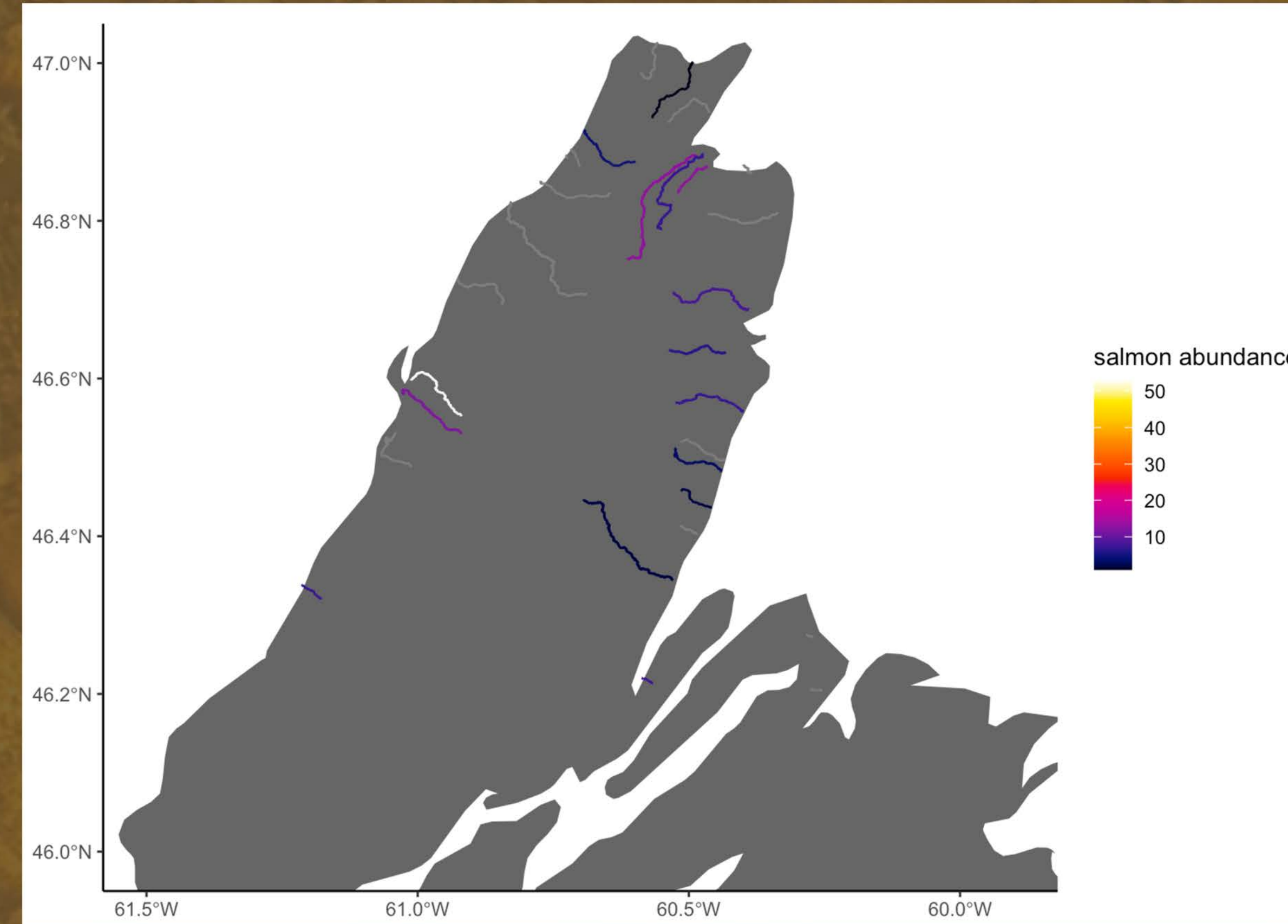


Figure 4 Salmon distribution and abundance across sampled rivers in the Cape Breton Highlands. Grey rivers indicate that salmon were absent

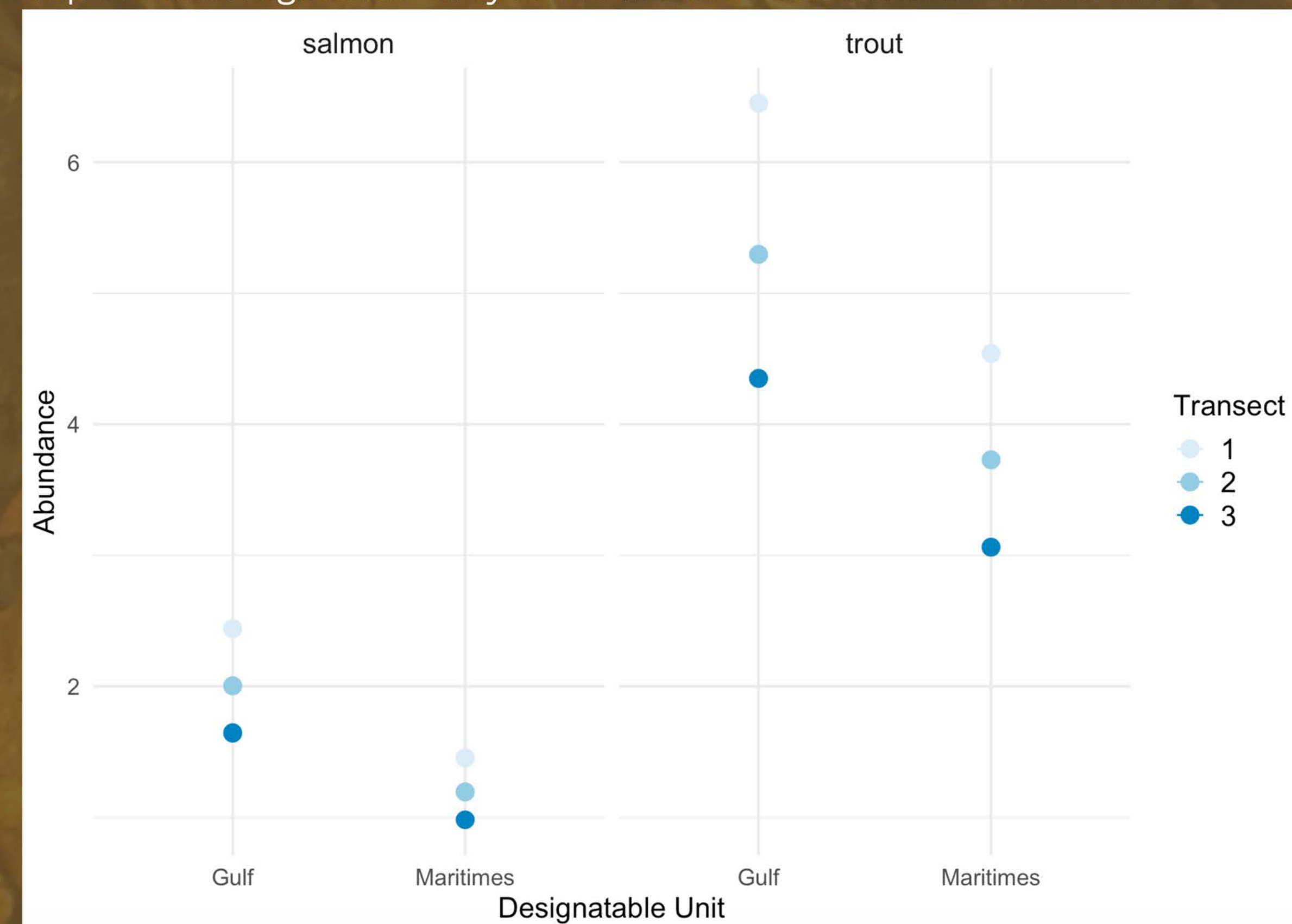


Figure 5 Predicted salmon and trout abundances for each transect of each DU based on a GAM (Poisson distribution), p-value = 0.204

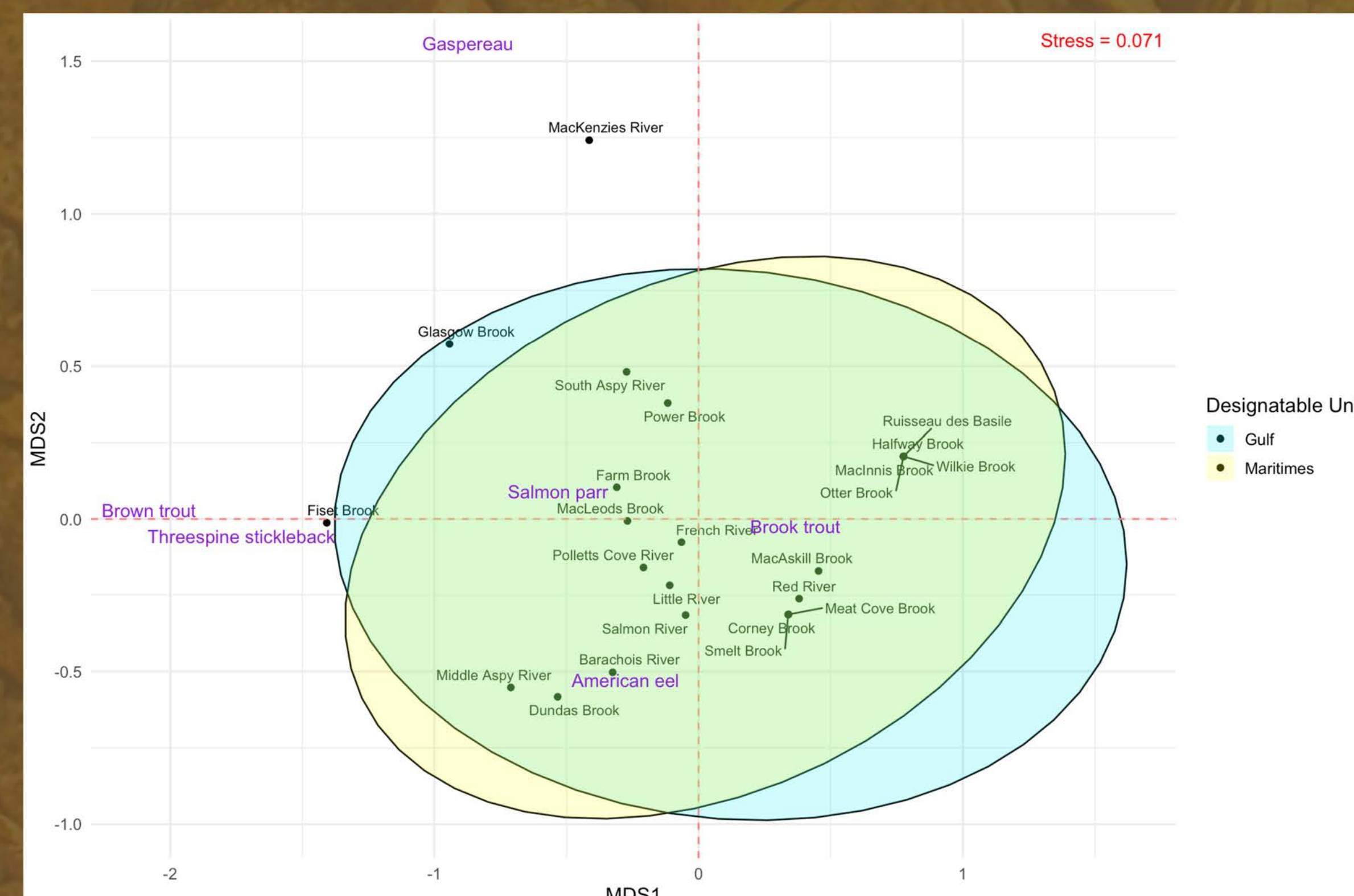


Figure 6 Community composition of Highlands rivers between DUs based on NMDS analysis. Elipses represent each DU at a 95% confidence interval

Methods

Site selection

- 24 rivers selected
- **Criteria:** river size (width and depth), gradient, accessibility to surveyors, even distribution across the area of interest.
- Rivers with limited data from the DFO, Parks Canada, or other published reports were prioritized

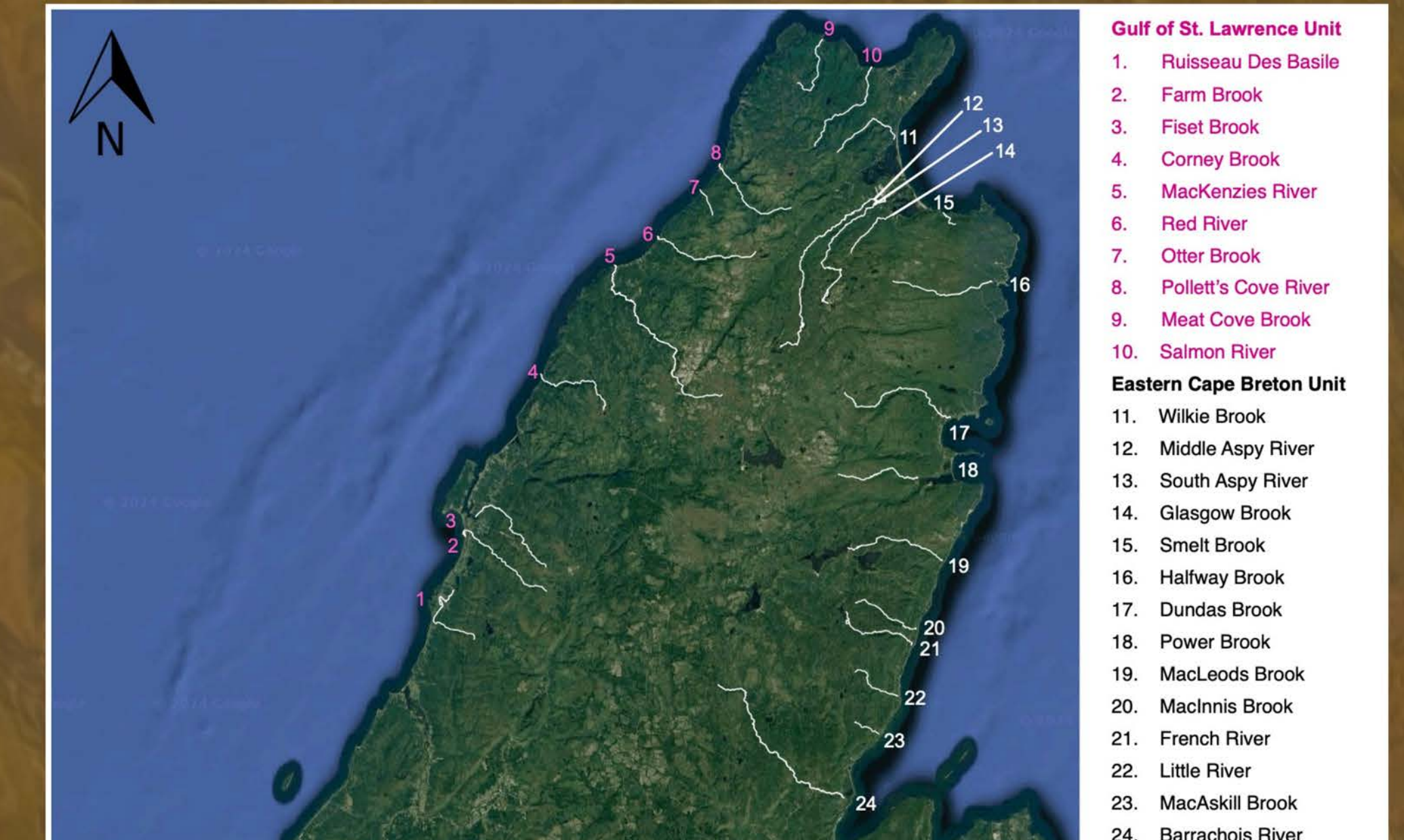


Figure 7 Map of the Cape Breton Highlands illustrating the rivers surveyed in June 2024 (white). The legend indicates the designatable unit in which each river belongs.



Figure 8 Electrofisher operator and dip netter surveying Polletts Cove River

Field Methods

- 3 transects of ~300 meters were surveyed with electrofisher (Figure 8); environmental metrics recorded
- Captured specimens were identified by species and fork length
- Genetic samples (upper caudal fin clippings) were collected from Atlantic salmon

Data Analysis

- NMDS and perMANOVA: compared species assemblages between DUs
- General additive model (GAM; poisson distribution) with ANOVA test: compared salmon abundance between DUs
- General linear model (GLM; Poisson distribution) tested environmental metrics with species abundance to determine predictors of salmon habitat suitability

Discussion

- Of the 24 rivers sampled, **13 systems** supported Atlantic salmon parr
- Population abundance and distribution are **similar** between DUs
- Community assemblages between DUs **similar**, despite differences in habitat

Significance

- Effective allocation of conservation resources requires accurate, up-to-date knowledge of the habitats occupied by Atlantic salmon
- In the event of the collapse of salmon stocks in large rivers, smaller systems may be the final strongholds of genetically diverse annually spawning fish
- This study emphasizes the consideration of small, isolated populations in the broader context of habitat restoration and conservation research

Acknowledgements

I would like to thank Josh Roland and Morgan Piczak for spending many hours in the field contributing to this project. Thank you also to the entire Lennox Lab for assisting throughout the analytical process of this project. This research would not have been possible without contributions from the Nancy Witherspoon Research Award and a grant from the Nova Scotia Freshwater Fisheries Research Cooperative

References

