What are the major phylogenetic lineages of rainbow trout (Oncorhynchus mykiss) and their geographic ranges? Tyler M. Breech, Janet L. Loxterman, and Ernest R. Keeley **Department of Biological Sciences**



Introduction

- The dynamic geological history in western North America has resulted in connected and separated streams over time.
- These events are thought to be a primary factor influencing the contemporary distribution of native fishes, and in many cases, has obscured the taxonomic relationship of fishes in this region.
- Rainbow trout (Oncorhynchus mykiss) are a freshwater fish species native to western North America.
- Many subspecies of rainbow trout have been hypothesized in different geographic regions.
- The relationship between taxonomic classification and major evolutionary lineages of rainbow trout is currently unknown.



Figure 1. Major subspecies of rainbow trout. Each color indicates a different subspecific region.

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Objectives

Our study seeks to investigate potential morphological and genetic/genomic differences among O. mykiss populations from across their range.

Methods

- Fish will be collected using backpack electrofishing.
- Specimens, 25-50 from each sampling site, will be photographed, weighed, measured, and have a fin clip taken.
- Common morphological characters will be assessed, as well as mtDNA sequencing of ND2 and SNP analysis.
- Principal component (PC) analyses and phylogenetic analyses will be used to compare populations both morphologically and genetically and create a rainbow trout phylogeny.



Figure 3. Example of landmarks placed on fish photos for morphometric analyses of body shape variation.







