Impacts of managed gene flow on disease resistance in hatchery Chinook salmon

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This information is preliminary and is subject to revision. It is being provided to meet the need for timely best science.

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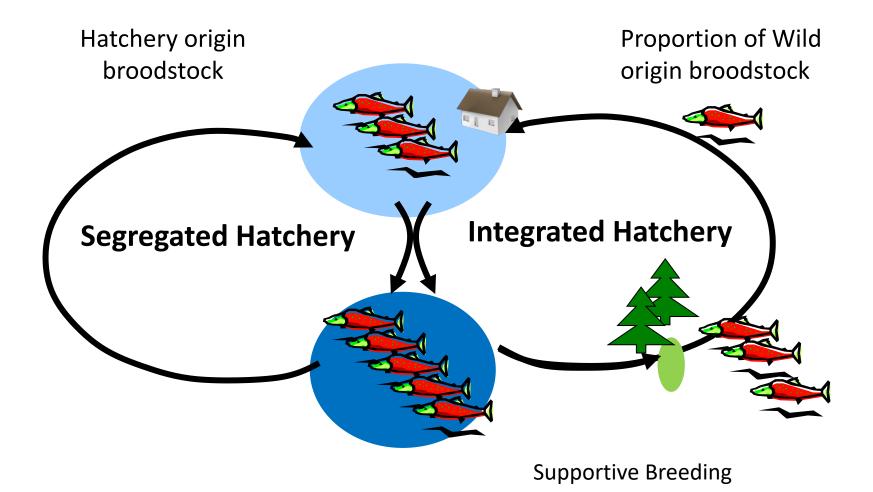
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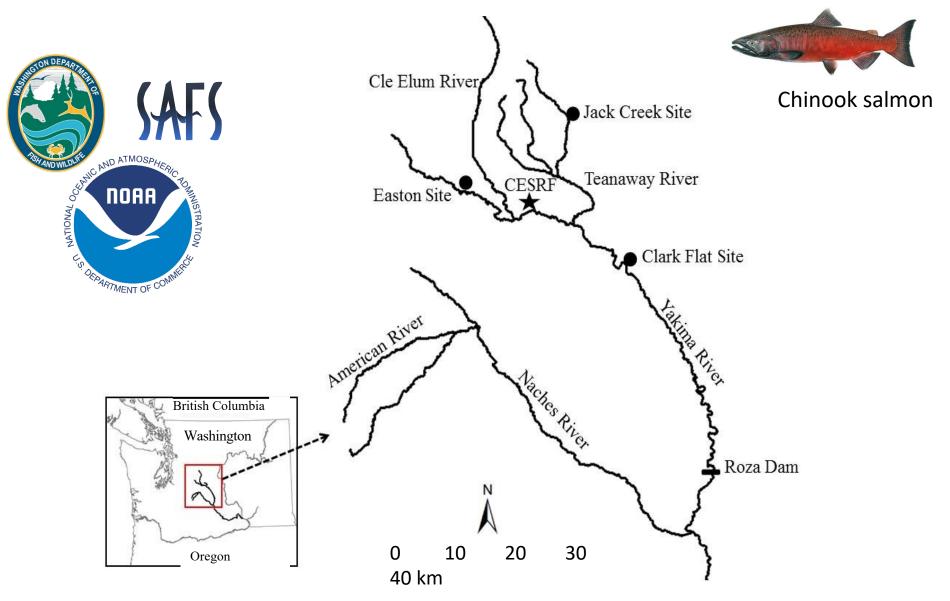


Hatchery Reform

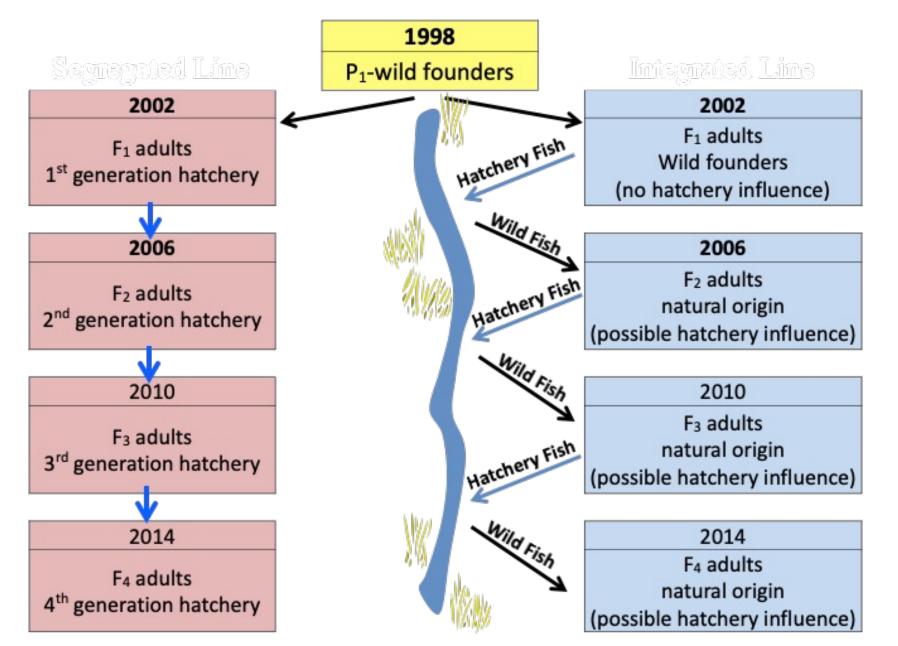


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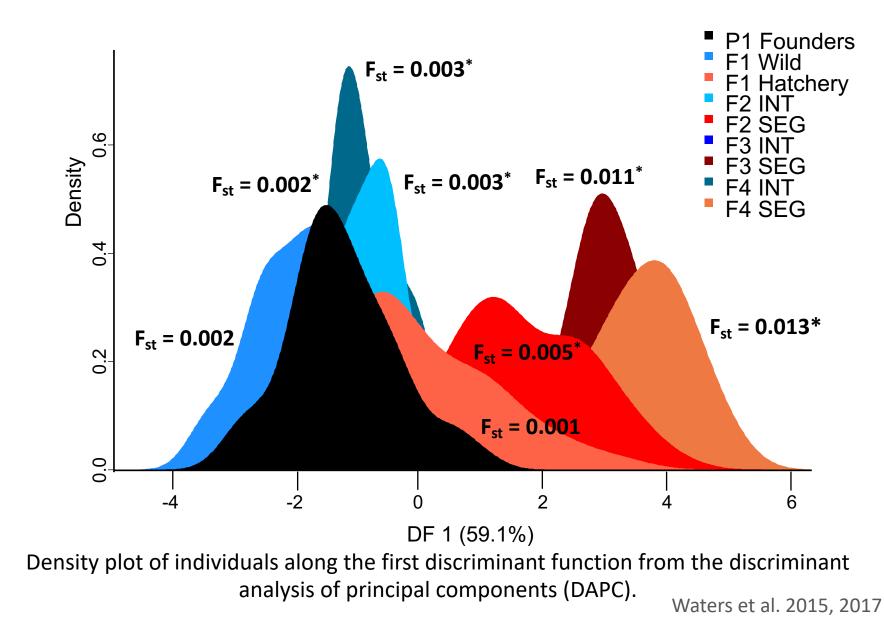
Experimental System



Experimental lines

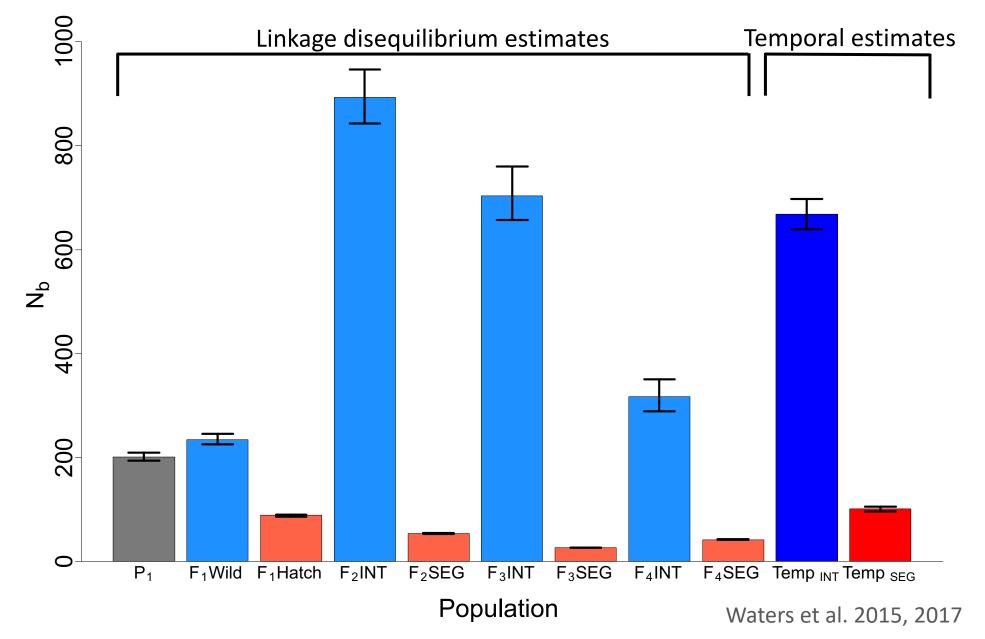


Temporal change in genetic variation



6

Effective number of breeders, N_b



7

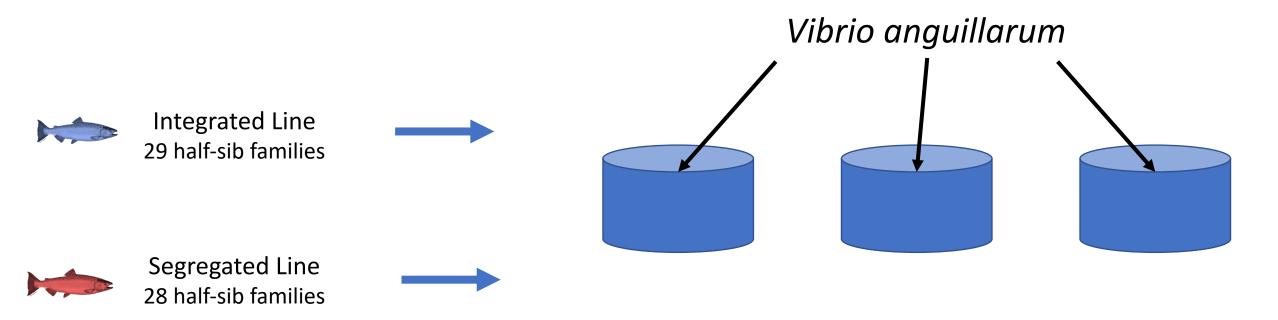
Study question

 How do different hatchery management practices impact disease response in propagated populations?

• Specifically, how does high (integrated) or low (segregated) gene flow impact disease response in hatchery fish?

CESRF represents an idealized, neutral system with best disease control practices

Experimental Design



1125 Fish (~375/tank)

Vibrio anguillarum

- Prevalent in Pacific Northwest
- Model bacterial pathogen
- Interested in generalized immune response to bacterial pathogens

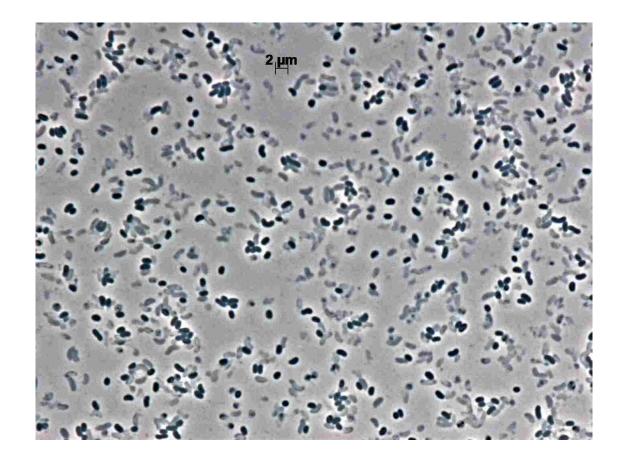
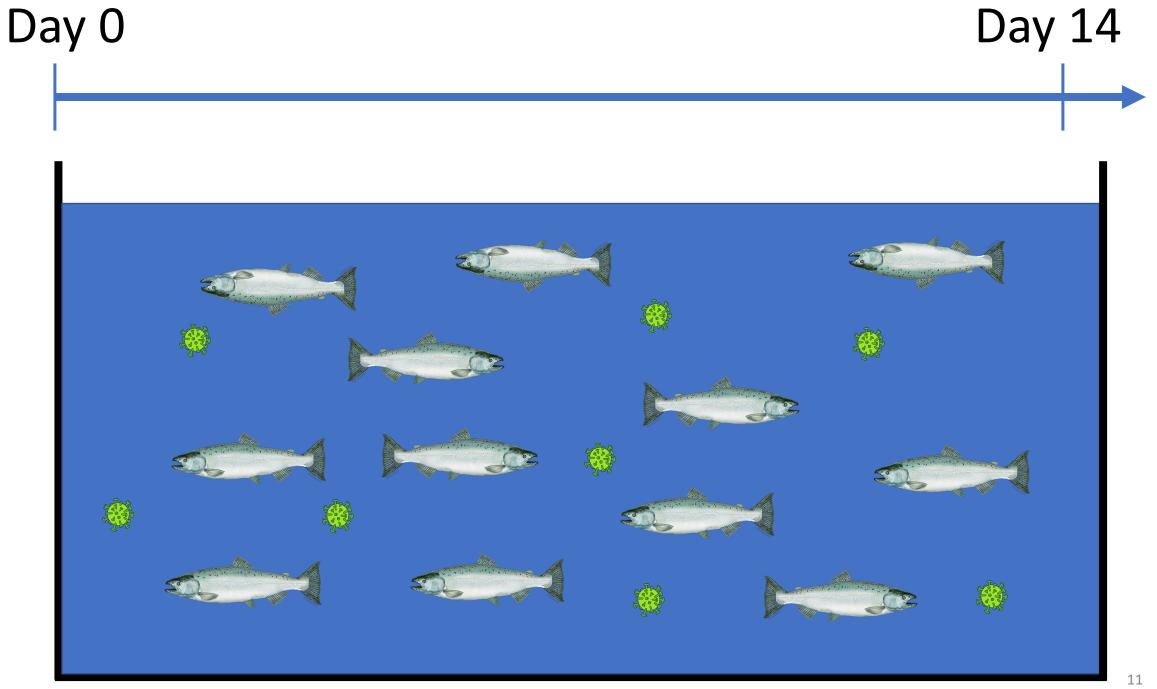
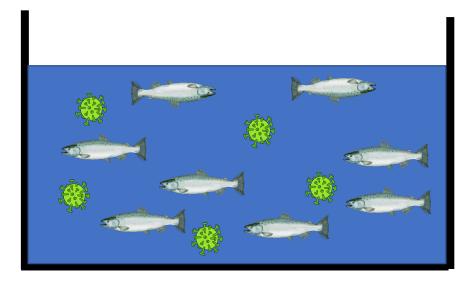
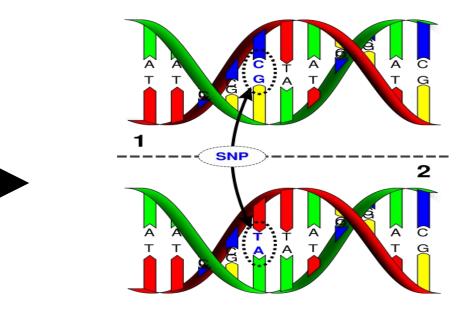


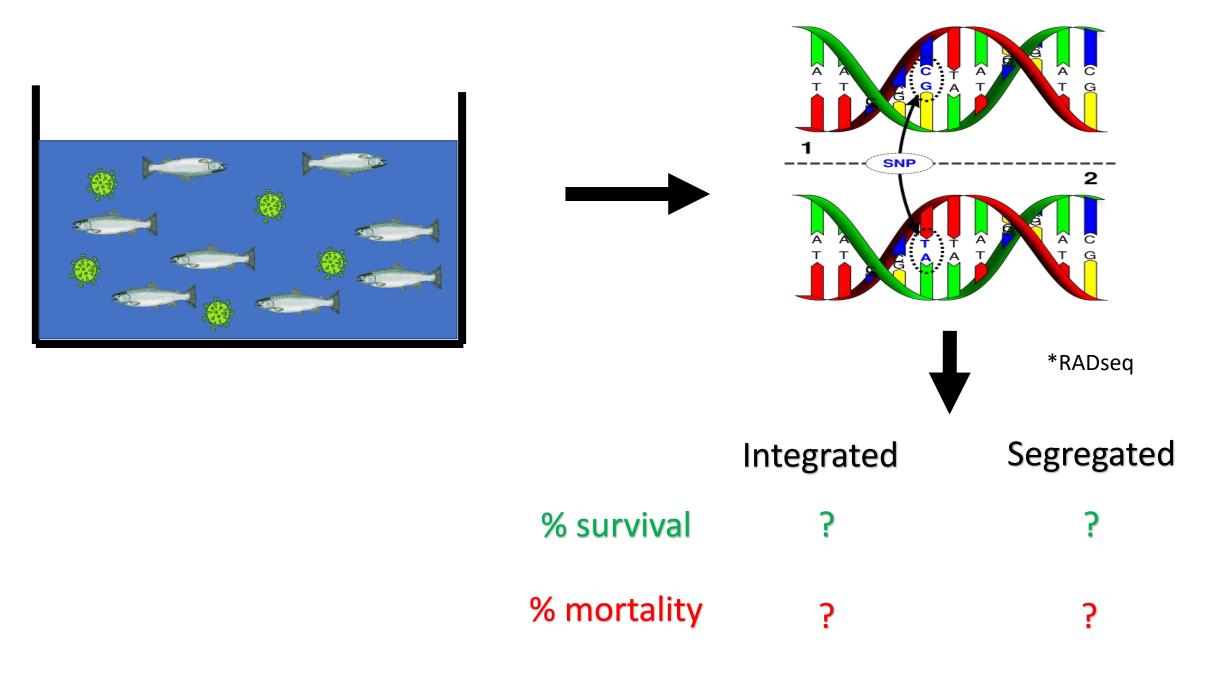
Image by Leibniz Institute DSMZ



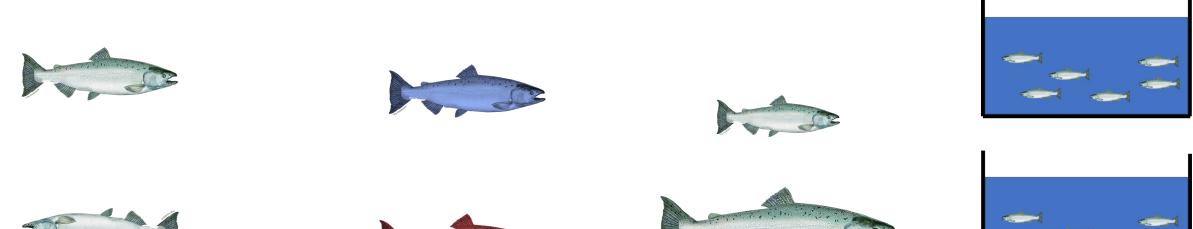




*RADseq



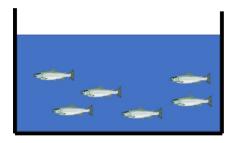
Mortality ~ Line + Length + Tank





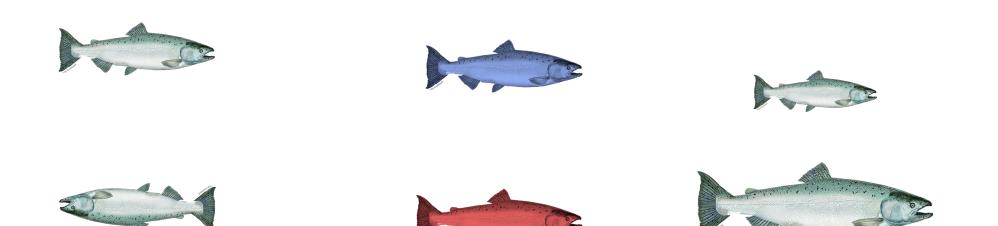




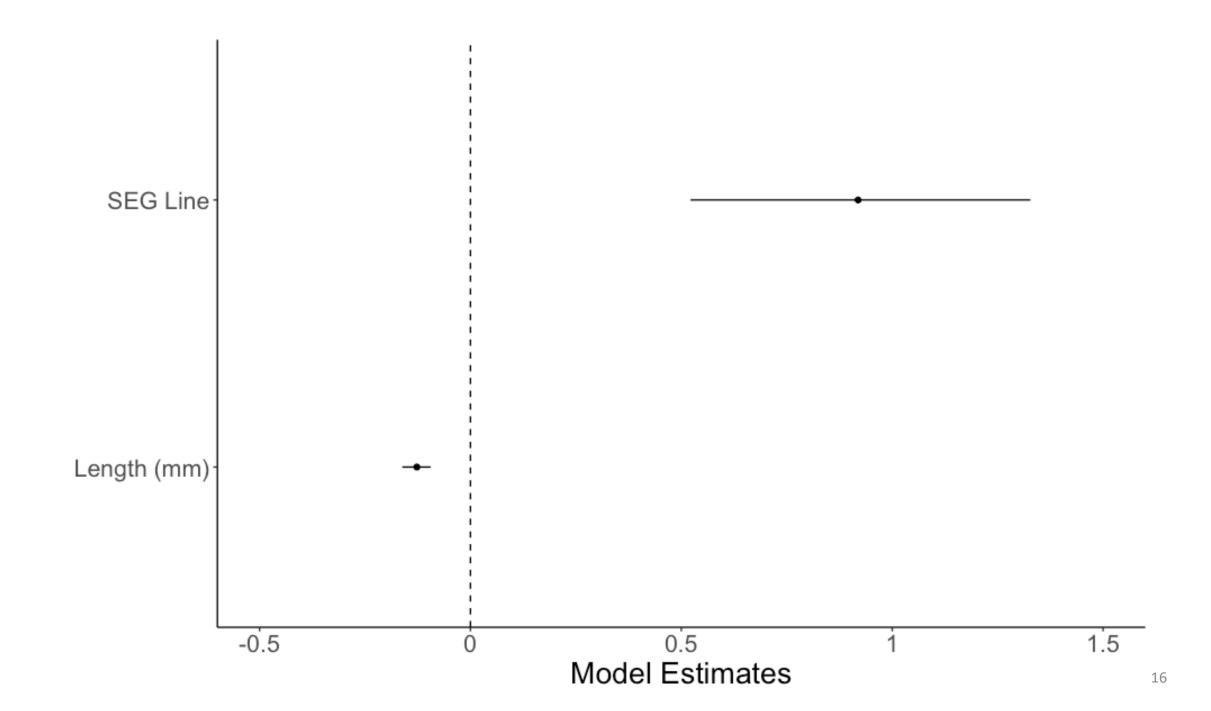


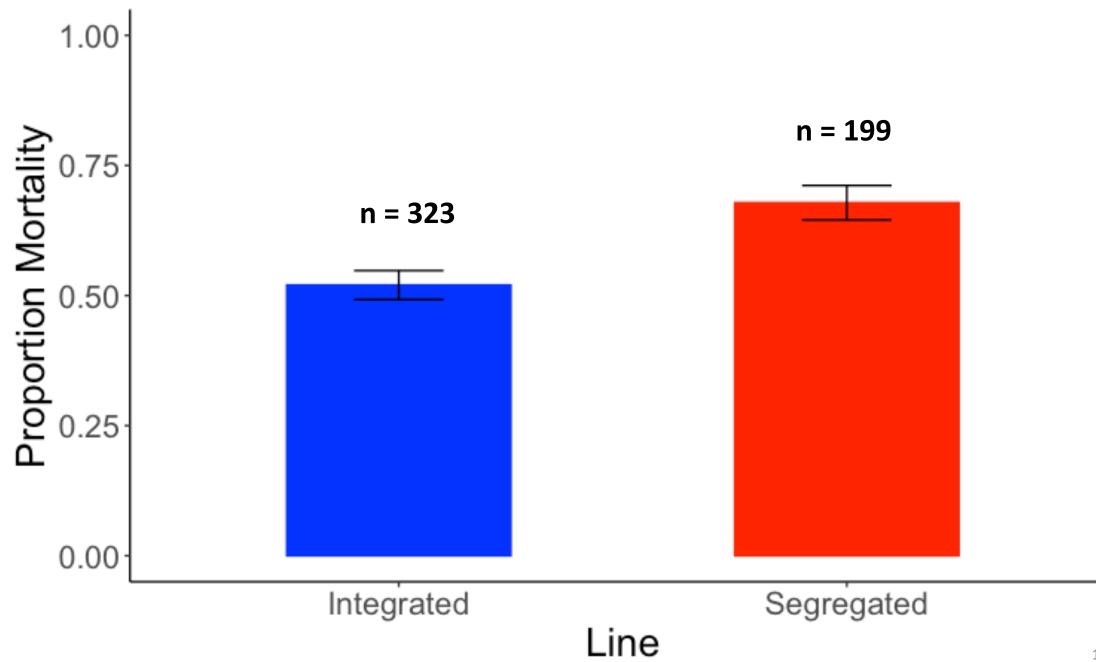
*Family included as random effect

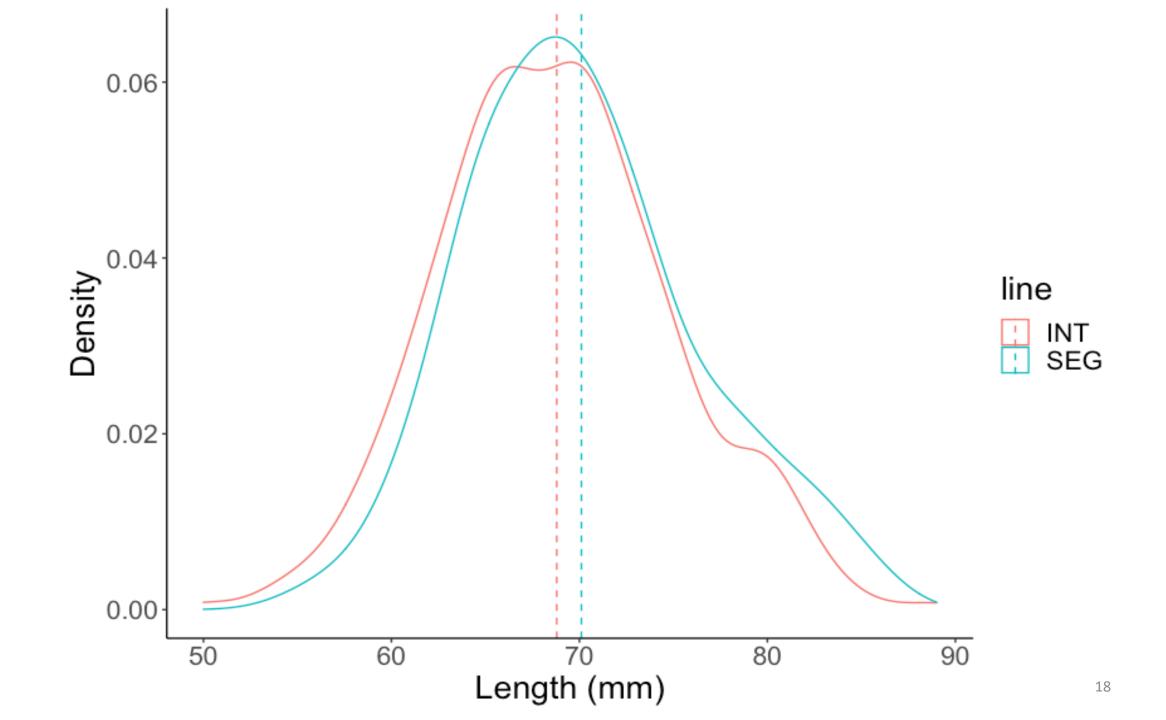
Mortality ~ Line + Length

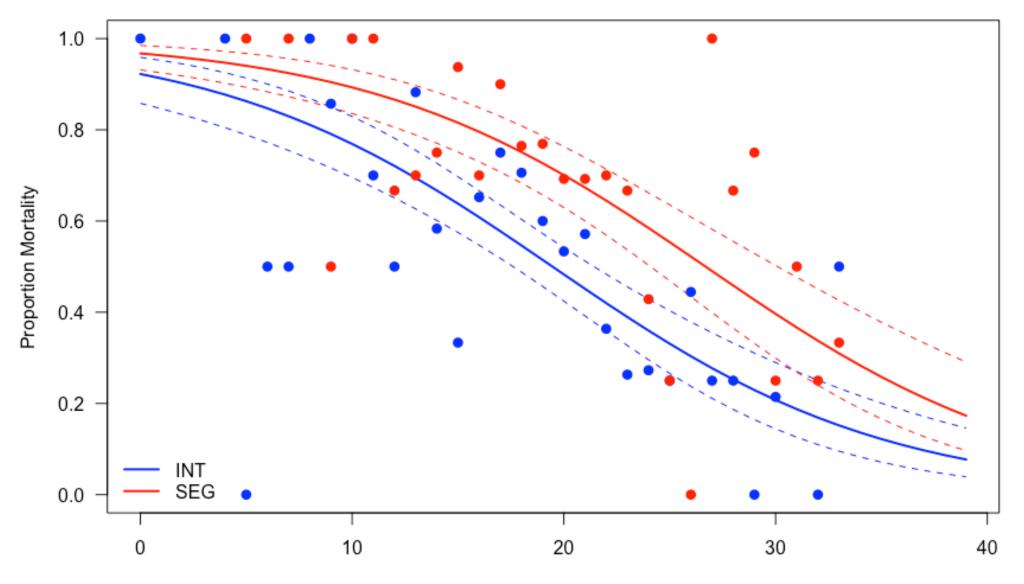


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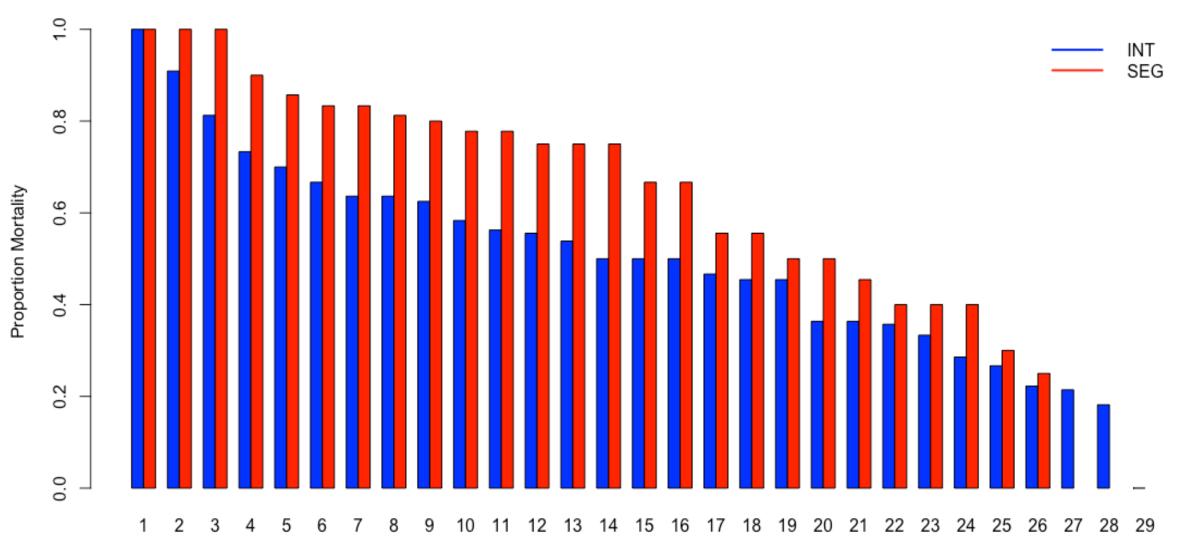








Body Length (mm) - 50 mm



Family Rank

Conclusions and Implications

- Managed gene flow (integrated) reduces overall susceptibility to V. anguillarum
- Disease resistance is variable by family
- Larger fish may be more resistance to disease

Future Direction

1. Incorporate models using "Days to Death" as response variable

2. Identify genetic markers associated with disease resistance