Spatial and Temporal Dynamics of Spawning between Native Westslope Cutthroat Trout, Introduced Rainbow Trout, and their Hybrids, with Implications for Hybridization and Loss of Adaptation

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Introductions of Exotic Species













Hybridization

- Loss of locally adapted gene complexes and ecological adaptations in native populations
- Threatens the persistence of many rare and endangered species













Cutthroat trout

- 14 subspecies
- Historic distribution includes Pacific NW, Great Basin, Rocky Mountains, Southwest

Westslope cutthroat trout

- One of four major subspecies
- Distribution includes the Columbia Fraser, Missouri, and Hudson Bay drainages of the US and Canada



New RBT introgression (8/14) Static RBT introgression (2/14) Static no RBT introgression (4/14)

Ν

2002

Hitt 2002

What factors influence successful invasion of hybrids?



Time



Upper Flathead River Drainage









A native species stronghold





Population Declines







Columbia Falls

Kalispell

Upper Flathead River System

ALK &

Objectives

- I. Identify RBT and hybrid source populations in the upper Flathead River system;
- 2. Compare the timing, location and movements of spawning by RBT, WCT, and their hybrids; and
- 3. Describe the patterns of spatial and temporal overlap among parental taxa and hybrid types.



Objective: Compare spawning dynamics

 Ho_1 : There are no differences in the timing and location of spawning among WCT, RBT and hybrid trout.

Predictions:

- Hybrid and RBT spawn during increasing and peak flows; WCT during peak and declining flows; Hybrids have a longer spawning period
- Hybrid and RBT spawn in tributaries lower in the drainage as compared to WCT







Tagging section

and and

Telemetry Methods



Timing and location of spawning

Spawning and movement variables:

- Beginning migration date
- Beginning migration temperature (°C)
- Beginning migration flow (m³/s)
- Spawning date
- Spawning temperature
- Spawning flow
- Migration period (d)
- Migration distance (km)







Genetic analyses



2000-2007 • 27 WCT • 11 WCT-hybrids • 36 RBT-hybrids • 51 RBT

N = 125 fish

Number of fish

Statistical Analyses

- MANOVA (genotype, sex, and year with interaction terms)
- MANOVA (genotype)
- Tukey's tests for post-hoc comparisons
- *P* < 0.05 alpha level

Genotype









Results

- 29 spawning tributaries were used
- 98% moved upstream
- No evidence of mainstem or sidechannel spawning





Results

- MANOVA (genotype, year, sex, interactions)
- No differences due to sex and interaction terms
- RBT and RBT-hybrids spawned significantly later and at higher flows in 2002, likely due to a protracted and higher spring runoff
- MANOVA (genotype): Wilks' λ = 0.164, F = 13.804, P
 < 0.0001
- Significant differences for five of the eight spawning and movement variables



Spawning Date



Migration period



Spawning temperature



Migration distance



Genotype

Timing of Spawning





Discharge (m³/s)

Timing of Spawning



≥USGS

Timing of Spawning





Discharge (m³/s)

WCT

 Spawned in the middle to
 upper portions
 of the drainage





RBT and Hybrids







2 F_1 hybrids RBT 2 Likely produced by migration from Abbot Creek

(Boyer et al. 2008)

Conclusions

Spatial and temporal overlap is occurring in the lower drainage, but streams in the middle and upper drainage still provide reproductive segregation



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Genomic extinction is imminent if hybrid populations with high amounts of RBT admixture are not reduced or eliminated.



