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Smoltification in an Impounded, Adfluvial Redband Trout Population Upstream from an Impassable Dam: Does It Persist?

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ARTICLE

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FIGURE 1. Study area indicating location of Mann Creek, Weiser River drainage, southwestern Idaho, relative to Brownlee Dam (left panel), and the Oxbow, and Brownlee dams on the Snake River (right panel).

Thrower and Joyce 2004; Thrower et al. 2008). Examples of anadromy include (Haner et al. 1995), and increased Na^+ -ATPase activity in the gill lamellae of residual anadromous populations that produce anadromous offspring (Zaugg and McLain 1972). Thrower and Joyce (2004) documented an anadromous population that produced anadromous offspring. The enzyme assay that detects Na^+ -ATPase activity in the gill lamellae is one of the more quantitative measures of anadromy. Pascual et al. (2003) documented anadromy in a stock of rainbow trout that was thought to originate from a resident freshwater population. Genetic analyses also suggest that gene flow between sympatric resident and anadromous forms of rainbow trout is more common than previously suspected (Olsen et al. 2006; McPhee et al. 2007; Pearsons et al. 2007).

Numerous factors can determine whether a salmonid can complete an anadromous life cycle. One important life history factor is the ability of juveniles to undergo smoltification, the physiological and morphological adaptation to seawater in preparation for ocean life. Smoltification and migration in individual fish can be influenced by growth, size, water temperature, and photoperiod (Zaugg and Wagner 1973; Pereira and Adelman 1985; Hirata et al. 1988). Additionally, genetics (Nichols et al. 2008) and landscape features (Narum et al. 2008) may influence life history expression and whether fish undergo smoltification. Numerous physical and biological indicators of the complex process of smoltification have been reported, including decreased condition factor (Zaugg and McLain 1972; Beeman et al. 1995), a silver body coloring or high skin

METHODS

Sampling.—Juvenile redband trout migrants were captured with a rotary screw trap (Thedinga et al. 1994; Roper and Scarnecchia 2000) positioned 2 km above the reservoir from 18 March to 3 June 2009. Water temperature was recorded hourly at the screw trap location with a Hobo Pro version 2 water

temperature data logger. Stream discharge was measured periodically throughout the trapping season with a flowmeter to establish a relationship between discharge and staff-gauge readings from a gauge located near the trap.

Fork length (FL) for each fish captured was recorded to the nearest millimeter and weight (W) to the nearest 0.1 g. Condition factor (K) was calculated for each fish by means of the formula $K = W/FL^3$ (Westers 2001). Scale samples were collected from each fish and independently aged by two readers. An age was assigned to each fish by using a double-blind protocol described by Scarnecchia et al. (2006). Fish age was determined from each scale sample independently by two readers and when discrepancies occurred, samples were re-aged independently by the same two readers. If there were any remaining discrepancies, the two readers consulted with each other and identified a single age for the sample.

Juveniles were assigned one of three body coloration patterns described by Negus (2003): banded, intermediate, or silver (Figure 2). Banded fish retained all parr marks, and the body was dark, colorful, or both. Intermediate fish had no parr marks near when dis-

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hereafter referred to as DART. The theoretical 50% passage date for Mann Creek redband trout at LGR was calculated by adding the number of days estimated to travel from the Mann Creek trap to LGR to the 50% passage date at the trap in Mann Creek. Redband trout would have to travel approximately 424 km downstream to reach LGR. We used published travel rates for steelhead smolts (Giorgi et al. 1997; Moore et al. 2010) and mean travel times reported by DART from populations in the Snake River basin that travel approximately 400 km to estimate travel time from Mann Creek to LGR.

Analyses—Analysis of variance (ANOVA) (significant at $P = 0.05$) was used to compare mean ATPase activities between banded and nonbanded fish for three migration periods: early (19 March–10 April), middle (28 April–5 May), and late (21

May–4 June). In 2006, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2007, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2008, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2009, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2010, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2011, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2012, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2013, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2014, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2015, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2016, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2017, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2018, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2019, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2020, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2021, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2022, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2023, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2024, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods. In 2025, the difference in ATPase activity between banded and nonbanded fish was significant ($P = 0.0001$) for the early migration period, but not for the middle ($P = 0.10$) or late ($P = 0.10$) periods.

studies. Furthermore, Thrower and Hard (2009) found that migrant progeny of resident *O. mykiss* had much lower marine survival rates than migrant progeny of anadromous parents, and Pearse et al. (2009) found that relatively few resident *O. mykiss* successfully completed an anadromous life cycle. Although smoltification is a necessary process for salmonids to complete an anadromous life cycle, it is only one of many factors that determine whether a fish can successfully migrate to and from a marine environment.

In our study area, coastal-origin hatchery rainbow trout *O. mykiss* have been stocked extensively since 1967 (Kozfkay et al. 2009). These stockings may have led to introgression of native and hatchery fish. Genetic analyses indicated that fish sampled from Mann Creek Reservoir were introgressed (Matt Campbell, Idaho Department of Fish and Game, unpublished data), while fish sampled in the headwaters of Mann Creek did not show signs of introgression (Kozfkay et al. 2011). There is evidently a barrier that prevented hatchery fish from introgressing with the headwater populations. However, it is unknown whether the putative smolts in our study are derived from the introgressed population, the pure population, or both populations. Future studies should examine genetic differences between the banded and nonbanded fish in this study as they could provide important information regarding smoltification traits and capabilities in *O. mykiss*.

Our results suggest the possibility that in portions of the Snake River drainage where steelhead have been extirpated as a result of artificial barriers, remnant populations may retain the potential for anadromy if migratory paths were reconnected. To fully interpret the evidence presented here that an isolated redband trout population continues to undergo morphological, physiological, and behavioral changes associated with smoltification, future research should examine *O. mykiss* populations, especially those in more interior regions, that have been blocked from migration for much longer periods of time. Examining populations with a longer history of isolation from marine environments could be useful for understanding the persistence of smoltification capabilities as well as other co-occurring aspects of redband trout life history.

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