

1 Introgressive Hybridization Between Native and Hatchery-origin Non-native Steelhead (*Oncorhynchus*
2 *mykiss*)

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7 Abstract

- 8 1. The artificial-propagation and release of individuals from non-local populations is a widespread
9 practice that can threaten the genetic integrity of native, locally adapted populations due to
10 domestication effects from the artificial rearing environments and inter-breeding with the local
11 populations. We examined introgressive hybridization in a threatened population of winter-run
12 steelhead (anadromous *Oncorhynchus mykiss*) in the Willamette Basin, Oregon. The non-local,
13 hatchery, summer-run steelhead is released annually into the basin as mitigation for the impact
14 of numerous dams.
- 15 2. Sixteen microsatellite loci were used to detect introgression in natural-origin adult steelhead
16 migrating into the basin prior to spawning during 2013 and 2014. Bayesian clustering analysis
17 (STRUCTURE) was used to identify the level of admixture in the population and assign individuals
18 to clusters.
- 19 3. The Bayesian clustering analysis indicated that there are most likely two populations (or
20 clusters) in the study area, a native, coastal, winter-run steelhead and a non-local, summer-run
21 steelhead which was derived from artificial crosses between summer-run coastal and interior
22 redband populations. Introgressive hybridization was detected in 26.4% of the natural-origin
23 adult steelhead. First generation (F_1) hybrids were estimated as 4.9 to 10.1% of the natural-
24 origin adult steelhead. Hybrids backcrossed to the native, coastal, winter-run steelhead were
25 nine times more numerous than backcrosses to the hatchery, summer-run steelhead. Upstream
26 migration timing was significantly different between the native, winter steelhead and the F_1
27 hybrids.
- 28 4. Low numbers of summer steelhead and back-cross summer-run hybrids were identified in the
29 natural-origin population consistent with reduced fitness of hatchery summer steelhead in
30 natural environments. Conservation actions that protect native populations from hatchery fish
31 include altering stocking practices (such as integrated management or sterility), and protecting
32 remaining intact populations by designating genetic preserves and preventing the release of
33 hatchery-origin or hybrid steelhead into these areas.

34 Keywords: hatchery effects, hybridization, introgression, microsatellite, non-native species, steelhead,
35 *Oncorhynchus mykiss*

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39 1. Introduction

40 Introduction of non-native species is a widespread management action implemented to support
41 harvest in forestry, fisheries and wildlife (Laikre et al., 2010). The intentional introductions for the
42 purposes of harvest programs threaten native populations due to the high propagule pressure that is
43 typical in these applications (Lockwood, Cassey, & Blackburn, 2005; Simberloff, 2009). Throughout the
44 western United States, fisheries mitigation programs largely developed in the early and mid-1900's rely
45 heavily on artificial propagation to replace lost fisheries production (e.g. U. S. House Document 531,
46 1949; Naish et al., 2007 and citations therein). Most of these programs pre-date the U.S. Endangered
47 Species Act passed in 1973, and the subsequent listing of numerous salmonid species during the 1990s
48 in the Pacific Northwestern, United States. Fisheries management and mitigation programs are
49 challenged to balance these conflicting mandates, and may use non-local or non-native fish to satisfy
50 harvest goals. Non-local and non-native salmonids cause significant impacts to the native populations,
51 communities and ecosystems (Buoro, Olden, & Cucherousset, 2016). Conflict between harvest or
52 recreation management, and the conservation of native species (e.g. Shouse, 2003) can result in
53 hatcheries having undefined or conflicting goals and objectives (Naish et al., 2007). Therefore, the
54 evaluation of interactions between native and propagated populations is important to determine if
55 harvest and conservation programs are in conflict, and identify possible management strategies to
56 reduce or eliminate impacts to native species.

57 Non-local, domesticated, hatchery salmonids largely impact the native conspecifics through
58 individual-based performance, such as physiological and fitness impacts. (Buoro et al., 2016). Artificially-
59 reared non-native or non-local populations often have reduced fitness in the wild due to domestication
60 effects (Araki et al., 2008; Le Luyer et al., 2017). The demographic effect of continued introductions has
61 been shown to cause rapid extinction of the native species even with reduced fitness of the non-native
62 or non-local population (Huxel, 1999; McGinnity et al., 2003). Reduced fitness of hatchery fish in the
63 wild and loss of production in the native population is identified in several commonly reared salmonid
64 species including: coho salmon (*O. kisutch*, Theriault, Moyer, Jackson, Blouin, & Banks, 2011), steelhead
65 trout (*O. mykiss*, Leider, Hulett, Loch, & Chilcote, 1990; Araki, Cooper, & Blouin, 2007; Berntson,
66 Carmichael, Flesher, Ward, & Moran, 2011), Atlantic salmon (*Salmo salar*, Jonsson, Jonsson, & Hansen,
67 1991; McGinnity et al., 2003) and Chinook salmon (*O. tshawytscha*, Chilcote, Goodson, & Falcy, 2011).
68 Hatchery and wild crosses also have reduced fitness in the wild with the poorest performance from non-
69 local, hatchery sources (Araki, Berejikian, Ford, & Blouin, 2008).

70 Hatchery steelhead derived from an artificial mixture of non-local, summer-run subspecies are
71 currently stocked in the Willamette Basin, a tributary to the lower Columbia River, United States, where
72 the native winter-run steelhead population is protected under the Endangered Species Act. The stocking
73 program mitigates for impassable dams that block access for anadromous populations of steelhead and
74 alters the remaining accessible habitats. The hatchery population of summer-run steelhead supports a
75 local recreational fishery due to the timing of return migration (spring to fall) and prolonged freshwater
76 rearing prior to spawning (up to 12 months). The use of highly domesticated hatchery stocks, such as the
77 Skamania stock used in the Willamette Basin (see below), is thought to limit the risk of colonization and
78 hybridization due to low fitness in the wild (Naish et al., 2007). The objective of this study was to identify
79 the extent of genetic interactions between the non-local, hatchery, summer-run steelhead and the
80 native, natural-origin, winter-run steelhead by assessing introgression and run timing of hybrids. This
81 study focuses on the effects of introgression on run timing, the distinguishing phenotype between the
82 summer and winter run steelhead. However, genetic impacts can occur in other phenotypes that were
83 not measured, but can be meaningful to local adaptations and evolutionary potential.

84 1.1 Steelhead Taxonomy and the Willamette Basin

85 *O. mykiss* exhibits diverse life history strategies. Variation is expressed within and among
86 populations and includes multiple return times for adults during spawning migrations, varying periods of
87 freshwater and ocean residency, and plasticity of life history between generations (Behnke, 1992; Busby
88 et al., 1996). Two subspecies of *O. mykiss* are recognized in the Pacific Northwest: a coastal subspecies
89 (*O. m. irideus*, also commonly called rainbow trout) and an interior subspecies (*O. m. gairdneri*, also
90 commonly called redband trout) (Behnke, 1992). Life history expression includes adult rearing in
91 freshwater rivers (resident or fluvial) or ocean (anadromous) in both subspecies. Steelhead, the
92 anadromous form of *O. mykiss*, exhibit different run timing and maturation phenotypes. The winter-run
93 matures in the ocean and migrates upstream immediately prior to spawning. The summer-run matures
94 in freshwater after returning from ocean rearing resulting in a protracted period during freshwater
95 rearing (Prince et al., 2017). Both phenotypes spawn in the late winter/spring months (~January to May)
96 (Van Doornik et al., 2015). The coastal subspecies largely expresses the winter-run maturation
97 phenotype, but may also express an alternate summer-run phenotype in some locations, usually in
98 upper basin habitats. The interior, redband trout only expresses a summer-run, maturation phenotype
99 (Behnke, 1992; Busby et al., 1996).

100 The native coastal steelhead in the upper portion of the Willamette Basin in western Oregon are
101 a distinct population listed as threatened under the Endangered Species Act in 1999 (NMFS, 2006). The

102 native steelhead upstream from Willamette Falls had a unique migration timing that is later than other
103 coastal steelhead populations thought to coincide with seasonal stream flow conditions that allowed
104 passage at Willamette Falls. Willamette Falls is a natural, 12-m high, horseshoe-shaped waterfall located
105 at river km 42 that was altered with navigation locks in 1873 and hydropower facilities in 1889 (OHS,
106 2018; PGE, 2018). A fish ladder was installed in 1885 to assist passage of salmon (PGE, 2018), prior to
107 this ladder, only the later-migrating native steelhead were able to ascend the falls.

108 The Willamette River Basin is 29,727 km² located in northwest Oregon between the Cascade and
109 the Coast Mountain ranges. The Willamette River is 301 km long and flows from south to north into the
110 Columbia River. Most of the urban population in the state resides in the basin including the cities of
111 Portland, Salem, Corvallis and Eugene (Fig. 1). Hydropower and flood control development in the basin
112 occurred during the 1960s and currently 13 high-head dams block access to 41% of native steelhead
113 habitat (Sheer and Steele, 2006). The accessible, lower elevation areas of the basin are impacted by
114 habitat alteration, urbanization and associated flood control, dam operations and associated altered
115 hydrology, climate trends, and landscape effects (see NMFS, 2016 for more detail).

116 1.2 Stocking History in the Willamette Basin

117 Stocking the non-local, summer-run (*Skamania* stock) steelhead began in 1966. Recently, the
118 hatchery program released an average of 572,411 juvenile steelhead annually between 2012 and 2017
119 at 6 sites in the study area (ODFW stocking records). Another non-native, hatchery-origin coastal
120 steelhead was artificially-propagated and introduced in the basin upstream from Willamette Falls
121 (commonly called early winter-run), and naturalized populations are thought to migrate into the
122 Willamette Basin during the fall months prior to spawning during the winter and early spring months
123 (~October to February 15) (Van Doornik et al., 2015). Introductions of this hatchery-origin, “early
124 winter” population (non-native coastal) were discontinued in the Basin upstream of Willamette Falls
125 during the 1990s, whereas the introductions of the *Skamania* summer-run hatchery steelhead are
126 ongoing (Van Doornik et al., 2015).

127 Juvenile hatchery, summer steelhead are marked by removing the adipose fin and these marked
128 adults are collected for broodstock at hatchery facilities; adults with intact adipose fins are naturally-
129 produced steelhead that can include: offspring of hatchery, summer-run steelhead that spawned in
130 rivers, native winter steelhead, or hybrids. Hatchery-origin, adult, summer-run steelhead out-number
131 the natural-origin steelhead 3:1 during the last 10 years (Fig. 2). Overlap among the migration and
132 spawning timing of these hatchery and native steelhead creates opportunity for spatial and temporal
133 overlap during spawning (Van Doornik et al., 2015).

134 2 Methods

135 The collection of steelhead was performed to capture a representative sample of the natural-
136 origin (adipose fin present) steelhead population passing Willamette Falls. Some hatchery-origin,
137 summer-run steelhead were analyzed only to provide a reference for the population clustering, and
138 provide an estimate of classification error from known (adipose fin removed) individuals. Sampling of
139 hatchery, summer steelhead ceased in late June due to high summertime water temperatures and
140 compliance with sampling permits, and did not resume later in the run after summertime temperatures
141 declined. Therefore, the hatchery, summer steelhead sample only captures the first half of the run
142 during 2013.

143 2.1 Fish Collections

144 Upstream migrating adult steelhead were captured at the adult fish trap located at the
145 Willamette Falls fish ladder from January 22, 2013 to June 26, 2013 and November 7, 2013 to June 4,
146 2014. Steelhead were tagged and sampled in approximate proportion to the number passing the fish
147 facility in each year. Sampled individuals represent an estimated 3.4% (2013) and 4.0% (2014) of the
148 natural-origin steelhead (adipose fin present) passing Willamette Falls.

149 Steelhead were anesthetized with AQUI-S 20E (AquaTactics, Kirkland, WA) and a radio-tag
150 implanted following the methods in Caudill et al. (2014). Total length and weight were measured and
151 fish were checked for marks or tags, sex, and evidence of injuries or disease. Hatchery-origin, summer-
152 run steelhead were identified by the removed adipose fin. Tissue was sampled as a caudal (tail) fin
153 punch and dried on paper. Telemetry records from fixed receiver sites (Fig. 1) and mobile tracking were
154 used to determine upstream extent of migration and infer spawning location using methods described
155 in parallel studies of Chinook salmon (Caudill et al., 2014; Keefer et al., 2017). Additionally, radio-tags
156 provided information on the extent of upstream migration and potential spawning location, as well as
157 the proportion of the individuals passing Willamette Falls that subsequently migrated downstream and
158 out of the study area. All collection, handling and tagging was performed using approved animal care
159 protocols and appropriate scientific collection permits.

160 2.2 Laboratory Methods

161 Sixteen microsatellite markers were used to genotype individuals. Thirteen of these markers
162 were standardized across the Columbia River basin and are summarized by Stephenson et al. (2009).
163 Additional primer sets analyzed were *Omy105*, *Omy2* (Heath, Pollard, & Herbinger, 2001) and *Omy77*
164 (Morris, Richard, & Wright, 1996). The DNA was isolated from fin clips using Qiagen DNEasy Blood and
165 Tissue™ extraction kits (Qiagen, Inc., Germantown, MD) according to manufacturer's protocols. Sixteen

166 microsatellite loci were amplified by PCR in three multiplex reactions using Qiagen Multiplex PCR Master
167 Mix in 96-well plates on C1000 Touch and S1000 thermal cyclers (Bio-Rad, Hercules, CA). PCR details are
168 provided in the Supporting Information, Section S.1. The PCR products were run on an Applied
169 Biosystems 3130xl Genetic Analyzer. Forward primers were fluorescently labeled (6-FAM from
170 Integrated DNA Technologies, Inc., Coralville, IA; VIC, PET, NED from Applied Biosystems, Inc.). PCR
171 product (1 μ l) was added to 10 μ l formamide and 0.15 μ l of LIZ500 size standard. Peaks were scored
172 using Genemapper software 5 (Applied Biosystems). All samples were amplified two or three times to
173 monitor for genotyping errors. Peaks were also visually checked for conformity to expected profiles.
174 Laboratory error rates were estimated at 1%.

175 2.3 Statistical Analysis

176 The characterization of the natural-origin run is based on the genetic analysis of 'wild' (adipose
177 fin present) steelhead (n=161 in 2013 and n=206 in 2014). A sample of known hatchery-origin summer
178 steelhead (n=89) were included in the analysis during the 2013 spawning year to provide a genetic
179 reference for this cluster (representing only the first half of this run). Data from the natural-origin run
180 were first tested for allele frequency differences across spawning years for the natural-origin steelhead
181 using GENEPOP version 4.2 (Raymond & Rousset, 1995). All multiple comparisons were adjusted with a
182 Bonferroni correction (Rice, 1989). No significant differences in allele frequency exact tests were
183 identified so data were grouped across years for the remainder of the analyses.

184 STRUCTURE version 2.3.4 (Prichard, Stephens, & Donnelly, 2000) was used to identify the
185 appropriate K-value for the data set and to identify admixture among these populations in individual
186 samples. STRUCTURE was run using the admixture model without location prior with 10,000 iterations
187 for burn-in followed by 100,000 iterations. Ten independent runs were performed for K ranging from 1
188 to 8. The optimal number of populations (K) was chosen according to Prichard, Wen, & Falush (2010)
189 and Evanno, Regnaut, & Goudet (2005). A K-means test was performed using GENODIVE (Meirmans,
190 2013) to confirm the number of clusters identified using STRUCTURE.

191 Individual admixture values (Q-value) from the STRUCTURE output were averaged from the 3
192 runs with the lowest log likelihood, which indicates the best fit between the data set and estimated
193 parameters. Hybridized (admixed) individuals were identified from the averaged Q-values at two levels
194 of cluster membership (<0.8 and <0.9) to either parental cluster (native winter or hatchery summer)
195 with hybrids assigning between 0.2 and 0.8 or 0.1 and 0.9, respectively. Individual steelhead were
196 identified based on the Q-values as native, hatchery or hybrid, based on the most supported number of
197 clusters. F_{ST} and allele frequency exact tests were performed on the final population groups using

198 GENEPOP version 4.2 (Raymond & Rousset, 1995). The proportion of unmarked hatchery, summer
199 steelhead was calculated as the number of hatchery steelhead out of the total number of the natural-
200 origin steelhead. The hybridization rate was estimated as the proportion of the natural-origin first
201 generation (F_1) hybrids in our sample. Due to a wide range of Q-values documented for F_1 hybrids using
202 STRUCTURE (Bohling, Adams, & Waits, 2012), a strict criterion (Q-values from 0.4 to 0.6) and a relaxed
203 criterion (Q-values from 0.3 to 0.7) were evaluated to estimate hybridization in the sample. These Q-
204 values were selected based on STRUCTURE assignments compared to pedigree data that estimated
205 individual F_1 hybrids with Q-values from 23 to 73% (Bohling et al., 2012).

206 Simulated data were used to identify uncertainty in individual hybrid identifications, potential
207 bias in the population-level estimate of introgression, and to inform error associated with
208 misidentifications related to the different Q-value thresholds (Q=0.8 versus 0.9). Error for the individual
209 assignments was evaluated using HYBRIDLAB version 1.1 (Nielsen, Bach, & Kotlicki, 2006) to simulate F_1
210 and backcross hybrids from the data set. The input data for the simulated hybrids were derived from 70
211 individuals from each subspecies collected during 2013 with Q-values > 0.9 for native, winter steelhead
212 and <0.1 for hatchery, summer steelhead, with no missing data. Seventy F_1 hybrids and back-crosses to
213 each parental species were simulated from the parental data. The input parental data and output hybrid
214 data were run in STRUCTURE with K=2 and all settings and procedures were similar to those described
215 above. The assignments of the simulated individuals were compared to the true (known) genotypes.

216 A principal coordinate analysis on genetic distances was performed using GENALEX version 6.5
217 (Peakall & Smouse 2006, 2012) for individuals in the identified groups (native winter, hybrid and
218 hatchery summer, Q=0.9). A Chi squared test was used to test for differences from expected counts of
219 native winter, hybrid, and hatchery, summer steelhead (Q=0.9). Differences in the timing of upstream
220 migration based on date of capture between the native winter and hatchery summer steelhead and
221 hybrids were tested using a Kruskal Wallis test in R version 3.4.1 (R Core Team, 2017) followed by a
222 Dunn test for multiple comparisons using the FSA package (Ogle, 2018). Date was converted to the
223 number of days after November 1 when the fish was captured at the collection facility.

224 3 Results

225 STRUCTURE output identified the optimal number of clusters as two (Fig. S1), and the K-means
226 analysis supported two clusters in the data based on the maximum pseudo-F value (Meirmans, 2012;
227 2013). The genetic diversity measures (heterozygosity, allelic richness, etc.) between these two groups
228 are summarized in Table S1. The potential for additional clusters in the data set were explored due to
229 previous research in the vicinity of the study area identifying three clusters in the study area (hatchery

230 summer steelhead, east side and west side tributaries) (Van Doornik et al., 2015). The results
231 considering three populations are included in the Supporting Information (Section S.2). Briefly, the
232 source of a third cluster is undefined, and only a small portion of the natural-origin adult steelhead
233 collected at Willamette Falls (~7% reduced to 5% after removing individuals that subsequently migrated
234 downstream prior to spawning; Fig. S2, Section S.3). Steelhead in this cluster had an earlier passage
235 timing at Willamette Falls than the native (late) winter-run steelhead (Fig. S3), and different F_{ST} and
236 allele frequencies than the summer and winter groups in the data (Table S2). However, PCoA of genetic
237 distances did not indicate clear clustering among the different genotypes (Fig. S4). This additional cluster
238 was not related to adult tributary migration (Fig. S5) or life history. Hereafter, the results are presented
239 in terms of two steelhead populations, a native winter and a hatchery summer.

240 Hatchery-origin summer-run steelhead clustered together in the STRUCTURE analysis mostly
241 following the hatchery identification with adipose fin removed (Fig. 3). Three hatchery-origin steelhead
242 did not assign to the hatchery steelhead cluster (3.4% misassignment rate) based on these known
243 samples. One of these hatchery-origin steelhead had a Q-value equal to 0.74 which would classify as
244 hybrid based on the Q-value, but did predominantly assign to the correct cluster. One steelhead
245 assigned as a hybrid with equal admixture among the populations, and one steelhead assigned to the
246 native steelhead group with a population assignment of 0.94. Similarly, a small portion of natural-origin
247 steelhead assigned to the summer-run, hatchery group (3.5% at $Q=0.9$ and 4% at $Q=0.8$). Individual Q-
248 values were similar across the three independent runs examined (average standard deviation <0.005)
249 with hybrids having higher average standard deviation than the parental populations (Table 1).

250 The estimate of introgressive hybridization ranged from 19 to 26.4% of the natural-origin
251 steelhead at $Q=0.8$ and 0.9 , respectively. Hybridization rate, estimated as the number of F_1 hybrids,
252 ranged from 4.9% (Q-values 0.4-0.6) to 10.1% (Q-values 0.3-0.7). The counts of individual native winter
253 and natural-origin summer steelhead and hybrids were not significantly different across the two years
254 ($p>0.10$). Counts of the natural-origin (hatchery-lineage) summer, native winter and hybrid steelhead
255 were significantly different from the expected counts (χ^2 test, $p<0.0001$), where native, winter steelhead
256 were over-represented and hatchery summer steelhead were under-represented in the natural-origin
257 steelhead (Fig. S6). When we examined the putative back-cross individuals (Q-values 0.7-0.9 and 0.1 to
258 0.3 using the Q threshold of 0.9 for cluster assignment), there were nine times as many hybrids back-
259 crossed to the native winter steelhead than the non-native hatchery summer steelhead in the sample.

260 The simulated data showed a wide range of individual Q-values for each genetic assignment
261 (native winter, hatchery summer and hybrids), but the average values across the samples were close to

262 the true genotypes (0.92 for native winter, 0.07 for hatchery summer, 0.49 for F_1 hybrids, 0.75 for F_1 x
263 native winter and 0.22 for F_1 x hatchery summer) (Figs. 4 and S7). Individual misassignments between
264 the native winter or hatchery summer steelhead and hybrid classifications generally ranged from 0 for F_1
265 hybrids using a $Q=0.9$ to 59% for back-crosses to hatchery summer steelhead using a $Q=0.8$ (Fig. 5). The
266 Q -value threshold of 0.9 generally provided lower error rates and the incorrect identifications were
267 nearly equal between the back-cross hybrids identified as parental type and the parental (hatchery or
268 native) groups identified as hybrids providing the most accurate estimate of introgression overall. A Q -
269 value threshold of 0.8 provided an unbalanced proportion of misassignments with fewer parentals
270 misassigned as hybrids, but high proportions of back-cross hybrids misassigned as parentals (>50%). In
271 terms of our F_1 hybrid criteria (relaxed versus strict), the strict criterion only correctly identified 40% of
272 the true F_1 hybrids and 11% of the back cross hybrids, whereas the relaxed criterion identified 73% of
273 the true F_1 hybrids and 31% of the back cross hybrids. Neither the strict nor relaxed criterion identified a
274 true parent as a hybrid.

275 Pairwise F_{ST} values were similar for the different Q -values, but were slightly greater for the Q -
276 value threshold of 0.9 (Table 2). Hybrids were more similar to the native group (F_{ST} 0.01, 0.02) than to
277 the hatchery summer group (F_{ST} 0.03). Only 9% of the variation of individual genetic distances was
278 explained by the first and second principal coordinate axes; however, the data show a clear gradation of
279 genetic distances among the groups with hybridized individuals intermediate between the native and
280 hatchery groups (Fig. 6).

281 The date of upstream migration was significantly different among the native winter, hatchery
282 summer and hybrid groups ($p<0.001$); however pairwise tests identified that the F_1 hybrids and native
283 winter steelhead were the only significantly different comparison ($p<0.001$). The median migration date
284 at Willamette Falls was earlier for all hybrid groups than the native winter-run steelhead (Figs. 7 and S8).

285 4 Discussion

286 Continued and frequent introduction of non-native or non-local populations results in high
287 propagule pressure that artificially increases the abundance and density of the non-local, hatchery
288 individuals (Lockwood et al., 2005; Simberloff, 2009), and counters the effects of reduced fitness of the
289 hatchery fish in natural environments. In the upper Willamette Basin, the hatchery summer steelhead in
290 the basin out-number the native winter steelhead preceding reproduction (Erdman, Caudill, Naughton,
291 & Jepson, in press). Introgression was detected in 26.4%, and F_1 hybrids were detected in 4.9 to 10.1%,
292 of the natural-origin samples. The simulation results support the estimated rates of introgression.
293 Hatchery summer steelhead were significantly under-represented in the natural-origin steelhead

294 samples indicating a lack of fitness in the natural environment. Individual putative back-cross hybrids to
295 the native winter steelhead were nine times more numerous than the hybrids back-crossed to the
296 hatchery summer steelhead. Hybrid steelhead had earlier median run timing than the native, winter-run
297 steelhead. Although the original (pre-stocking) genetic differentiation between these populations is
298 unknown, introgressive hybridization can alter the genetic integrity of the native population and lead to
299 replacement of the population by the non-native (Huxel, 1999; Ford, 2002). Local adaptations and
300 genetic diversity lost in these processes may not be recoverable in time frames relevant to the
301 management and preservation of natural resources (~100 years) which could reduce the evolutionary
302 potential of the population (Prince et al., 2017; Waples & Lindley, in press).

303 4.1 Accuracy of Hybridization and Introgression Estimates

304 A key question is whether admixture is under- or over-estimated at the population-level using
305 the STRUCTURE assignments. Estimation of introgression first depends on the ability to reliably detect
306 the clusters in the analysis which improves with greater genetic differences. Simulated data sets
307 estimate that STRUCTURE is about 97% accurate when F_{ST} is 0.05 (Latch, Dharmarajan, Glaubitz, &
308 Rhodes, 2006), a level of differentiation similar to the populations included in this study. Yet, at the
309 individual-level, hybrids have a wide range of admixture estimated by STRUCTURE, but the mean Q-
310 value of each type of hybrid was close to the expected value when tested with a known pedigree (also
311 shown in Bohling et al., 2012). Further, the user-defined K-value and Q-value thresholds will influence
312 the amount of introgression estimated using these methods (see Supporting Information, Section S.5 for
313 more discussion). In a simulated data set, using the Q-value of 0.8 resulted in 42% of backcrosses being
314 incorrectly assigned to the purebred group while using a Q-value of 0.9 reduced this error to 19% (Vaha
315 & Primmer, 2006), similar to our simulation results. Certainly, at the individual level and with low
316 genetic differentiation ($F_{ST} < 0.1$), there is error in the identification of individuals. Based on the findings
317 in Vaha & Primmer (2006) at an F_{ST} value of 0.06, we expect to correctly identify about 80% of the
318 individuals in this sample using 16 loci. The simulated hybrid data set confirms this 20% error in
319 assignment between parental groups and back-cross hybrids. Overall, these data and other studies
320 suggest high rates of correct assignment for F_1 hybrids and unbiased population-level rates.

321 4.2 Potential Factors Mediating Introgression from Hatchery Steelhead

322 Hatchery, summer steelhead have been stocked in the study area for about 10 generations, and
323 two explanations are possible for the observed hybridization pattern in the natural-origin population.
324 First, the genetic differentiation in the native winter and hatchery summer populations could be a result
325 of genetic mixing that is reducing genetic differentiation over time and moving the populations toward a

326 hybrid swarm. Briefly, the sample represents a “snapshot” during a temporal process. Under this
327 scenario, introgression is expanding toward an inevitable hybrid swarm, and delaying a management
328 action will result in the extinction of the native genotype. An alternative hypothesis suggests that the
329 level of introgression could be mediated by loss in fitness of hatchery summer steelhead and hybrids –
330 thereby resulting in a stable level of introgression (Chilcote et al., 2011). Epifanio & Philipp (2001) show
331 that introgression will asymptote at about 20% admixture in 10 generations when fitness of the hybrids
332 is low. This scenario requires very low relative fitness of the hybrids at 0.15. In their model, even a slight
333 increase of relative fitness to 0.20 resulted in complete introgressive hybridization within 11
334 generations. If introgression is stabilized at these levels, then it is possible that discontinuing the release
335 of artificially propagated summer-run steelhead which have very low fitness in the natural environment
336 could result in the decline of the summer-run steelhead in the population, and the potential negative
337 effects of introgression may be limited. Although assortative mating could be a mechanism that
338 maintains genetic differentiation providing another alternative hypothesis, Epifanio & Philipp (2001)
339 show that strong pre-mating isolation alone is insufficient to prevent a complete hybrid swarm if fitness
340 of the hybrids is at least 0.20 of the fitness of the native species or population. Even if considering a
341 conservative estimate of F_1 hybridization, the spatial and temporal overlap between the native winter
342 and hatchery summer steelhead is enough to cause introgression. Additionally, the timing of spawning
343 between the native winter and the hatchery summer steelhead overlaps (Van Doornik et al., 2015)
344 minimizing the potential for segregation in the natural environment.

345 Few natural-origin, summer steelhead were detected in the population, and lower numbers of
346 putative hybrids back-crossed to the hatchery summer steelhead despite high abundances of hatchery
347 summer steelhead in the population during reproduction. Jones et al. (2015) found similar proportions
348 of hatchery, hybrid and wild steelhead in a stream in Southwest Washington (1% natural-origin,
349 hatchery-lineage steelhead and 29% hybrids), and conclude that the earlier spawning hatchery
350 steelhead were emerging during unfavorable conditions; however, the later emerging hatchery
351 steelhead were more likely to encounter better environmental conditions for early rearing resulting in
352 selection for this later migration timing that is more similar to the native population. The timing of adult
353 migration is a heritable trait in salmonids and has substantial fitness consequences if not synchronized
354 with local environmental conditions (Flagg, Waknitz, Maynard, Milner, & Mahnken, 1995; Quinn,
355 Peterson, Gallucci, Hershberger, & Brannon, 2002; McLean, Bentzen, & Quinn, 2005; Jones et al., 2015;
356 Hess, Zandt, Matala, & Narum, 2016).

357 Recent research indicates that modifications at the GREB1L gene are associated with premature
358 migration, and that premature migration (expressed in summer steelhead) is not masked in the
359 heterozygote (Prince et al., 2017). Yet, upstream migration after maturation is not a conduit for the
360 premature migration allele, and data patterns indicate selection against the intermediate phenotypes
361 (Prince et al., 2017). This interaction between run timing, maturation and spawning timing is likely a
362 complex expression of multiple gene complexes (Waples & Lindley, in press, and citations therein). The
363 earlier run timing expressed via the summer introgression in this data set suggests that the later
364 migrating hatchery summer steelhead (August to October) may have a greater likelihood to hybridize
365 with the native winter steelhead which spawn after the summer steelhead. An alternate explanation for
366 the observed influence of the later migrating hatchery summer steelhead in the hybridized individuals
367 could be less opportunity for removal from the population due to reduced fishing pressure or other
368 mortality during freshwater maturation.

369 4.3 Conservation and Management

370 Options for harvest-focused hatchery programs are to maintain programs with complete
371 segregation between the native, wild and hatchery populations, or to fully integrate the two populations
372 with the hatchery programs providing surplus production to meet harvest goals. Unfortunately, the
373 efficacy of these approaches has not been tested over the long term (Naish et al., 2007). Complete
374 segregation relies on the ability to collect all artificially propagated adults prior to spawning. Various
375 other management strategies to support segregation have been attempted to address threats from
376 hatchery fish, such as introducing sterile hatchery fish (Tiwarly, Kurbagaran, & Ray, 2004) where the
377 threat of introgression or inter-breeding is high (*e. g.*, Cotter et al., 2000), and the designation of genetic
378 preserves (also called gene banks) to protect and promote the recovery of native populations (*e.g.*
379 WDFW, 2015). Integration is also commonly implemented, but the rapid epigenetic effects creates
380 challenges to limit the transmission of genetic impacts to wild populations (Le Luyer et al., 2017).

381 Reduced fitness and misidentification are a concern when stocking hatchery fish without
382 complete segregation. Hybrids and natural-origin, hatchery individuals will be counted as native species
383 during abundance estimates and population demographic analyses. Currently, managers in the study
384 area are transplanting wild and presumably native winter steelhead returning to the basin into
385 inaccessible spawning areas upstream from dams, thereby introducing the hatchery-origin summer
386 steelhead genes into these areas. Although the majority (~70%) of the natural-origin returning adults to
387 the upper Willamette Basin are identified as native winter steelhead, the estimate of encounter rates
388 based on the observed proportions of F_1 hybrids in the data set is a concern. This likely underestimates

389 the effect of the hatchery summer steelhead due to the low expected fitness. For example, at a
390 hybridization rate of 5% and a relative fitness of 0.10, the hybrid encounter rates not expressed in the
391 observed data could be as high as 50%. If relative fitness is increased to 0.40, then hybrid encounter
392 rates would be about 12%.

393 Future research should examine the fitness consequences to the hybridized individuals and
394 identify the effects of the mitigation stocking program on the population demographic. Fitness can be
395 acquired from individual-based lifetime tagging studies and genotyping over time; however spawning
396 behavior, hybridization attempts and redd disturbance may be difficult to estimate due to high stream
397 flows, low water visibility and large habitats where steelhead occur in the study area. Importantly, long
398 term studies are needed to best identify strategies to prevent impacts to the native, wild populations.
399 Information on the migration and spawning behavior of the later migrating (passing Willamette Falls
400 between August 1 and October 31) hatchery summer run steelhead is lacking, and information on
401 whether these individuals are more likely to contribute to hybridization could assist in identifying
402 potential management actions to reduce effects in the basin. Additionally, fisheries management in the
403 local basin has included the practice of “re-cycling” hatchery, summer steelhead to maximize angler
404 opportunity (*i.e.* trapping, transporting and re-releasing adult hatchery steelhead downstream).
405 However, while this program boosts angler harvest by an estimated 15%, it also likely increases the
406 reproductive interactions between hatchery and wild steelhead by leaving many mature, hatchery
407 steelhead in-river (Erdman et al., in press).

408 Areas, like the Columbia River Basin, where existing harvest and conservation policies are
409 conflicting, demonstrate that clear guidance on priorities and updated policies that ensure and
410 encourage the conservation of native species and ecosystem function are needed to assist in the
411 complex processes implementing fishery programs under these laws. Yet, defining common policies has
412 proven challenging when management actions span multiple agencies with different underlying
413 objectives. Even within agencies, different departments may have their own competing objectives and
414 complex infra-structure. Areas currently developing new hydropower projects should strive to more
415 fully balance the conservation of native species and the replacement (or mitigation) for lost subsistence
416 or commercial fisheries, and create programs and policies that can evolve with scientific advances and
417 socio-political concerns. Areas where artificial propagation is planned or ongoing should implement
418 genetic monitoring practices to ensure the preservation and evolutionary potential of native genotypes
419 (Schwartz, Luikart & Waples, 2007; Bohling 2016), and promote adaptive management based on
420 scientific findings and best-available science.

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https://wdfw.wa.gov/conservation/fisheries/steelhead/gene_bank/puget_sound/summary_of_candidate_populations_7-1-15.pdf [26 March 2018].

Accepted version

655 Table 1. Average Q-value for individual assignments and standard deviation for hatchery and native
656 steelhead and hybrids in the upper Willamette River Basin. Results for the two tested Q-values (0.9 and
657 0.8) are presented.

658

	Q=0.90		Q=0.80	
	Avg	SD	Avg	SD
Native, winter	0.96	0.00098	0.95	0.00132
Hatchery, summer	0.03	0.00069	0.04	0.00076
Hybrid	0.65	0.00375	0.56	0.00423

659

660

661 Table 2. F_{ST} and allele frequency tests among clusters identified by STRUCTURE at $K=2$. F_{ST} values using a
662 Q-value 0.8 are shown above the diagonal and Q-value 0.9 is shown below the diagonal. All pairwise
663 comparisons had significantly different allele frequencies.

664

	Native, winter	Hybrid	Hatchery, summer
Native, winter		0.015	0.057
Hybrid	0.013		0.022
Hatchery, summer	0.065	0.028	

665

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667

668 Figure Legends

669

670 Figure 1. Map of study area and locations of the downstream-most impassable dams (gray rectangles)
671 and stationary radio-telemetry receiver sites (blue circles) in the study area. Fish were captured and
672 tagged at Willamette Falls fish ladder.

673

674 Figure 2. Counts of hatchery summer and native winter steelhead at Willamette Falls by year.
675 Identification is based on adipose fin presence/absence. Data from Oregon Department of Fish and
676 Wildlife (ODFW 2017).

677

678 Figure 3. STRUCTURE plot K=2 for adult steelhead in the upper Willamette Basin. Individuals are
679 represented by one bar in the plot and samples are ordered by date captured during the upstream
680 migration (x-axis). The clusters group with native winter (green) and hatchery summer (red) steelhead.
681 Hybridized individuals show contributions from both winter and summer steelhead corresponding to the
682 Q-value shown on the y-axis.

683

684 Figure 4. STRUCTURE output for individual simulated genotypes (y-axis) versus true genotype (x-axis).
685 The simulated hybrids are shown on the x-axis according to the proportion of native winter steelhead
686 (0.5 for F_1 hybrids, 0.25 for F_1 x summer and 0.75 for F_1 x winter).

687

688 Figure 5. Percentage of individuals misassigned in each genotype category using a minimum Q-value of
689 0.9 and 0.8 for cluster assignment. Misassignment rate is the proportion of true (simulated) individuals
690 assigned as: known winter or summer as hybrid; known F_1 or back cross hybrids as winter or summer.

691

692 Figure 6. Principal coordinate analysis (PCoA) of individual genetic distances for steelhead in the upper
693 Willamette Basin (Q=0.9) for native winter (blue diamonds), hatchery summer (black triangles), and
694 hybrids (orange squares) identified by STRUCTURE output.

695

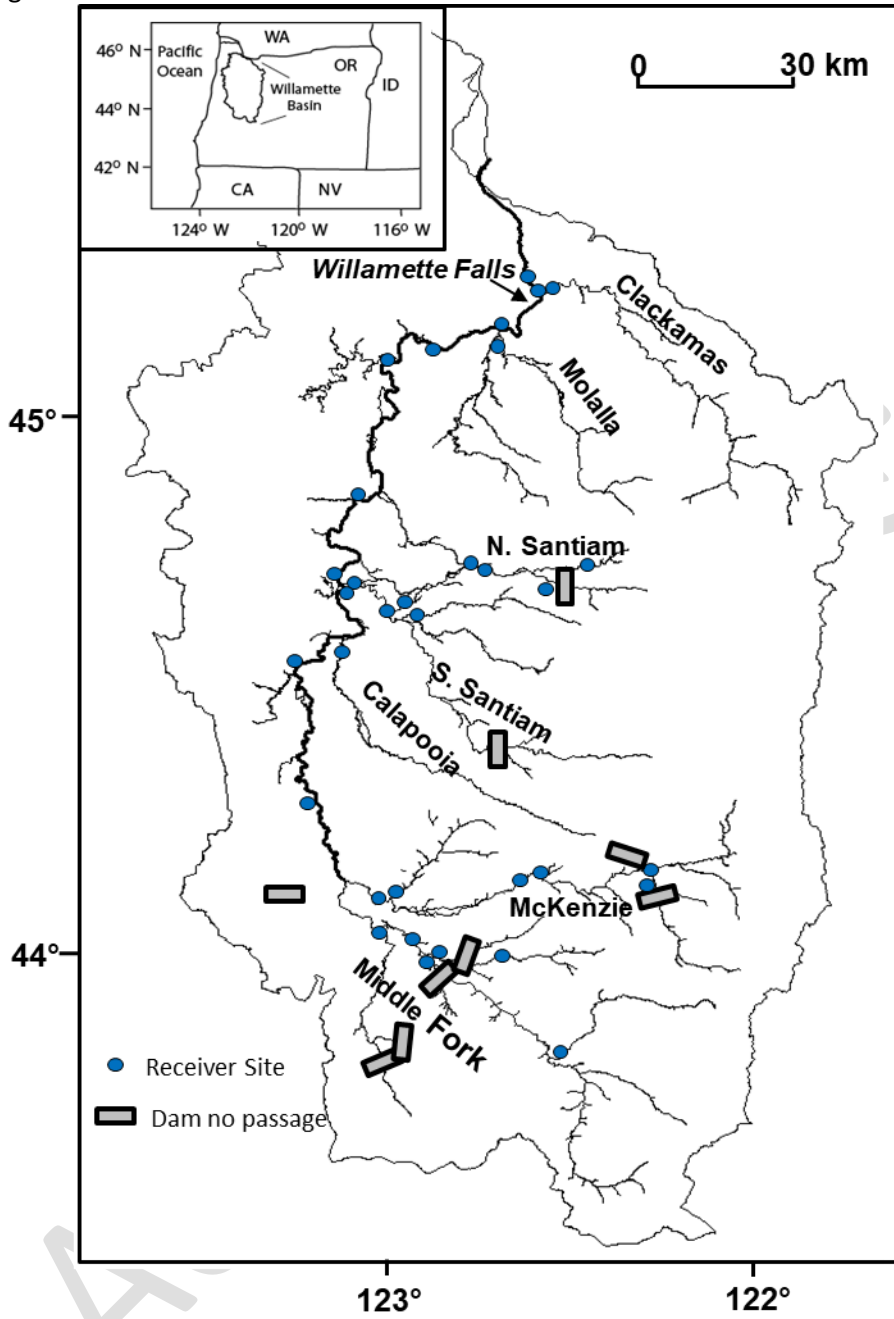
696 Figure 7. Migration timing (range and median) of natural-origin, steelhead by genotype (native, winter
697 steelhead, F_1 and back cross hybrids), and hatchery-origin, summer steelhead at Willamette Falls. The
698 native winter and hybrid run timing is based on the genotyped sample collected for this study. The
699 hatchery summer steelhead run timing is based on the run counts at Willamette Falls for the adult
700 returns for the 2014 spawning year (return date 1 November 2013 to 31 October 2014) (ODFW, 2017).

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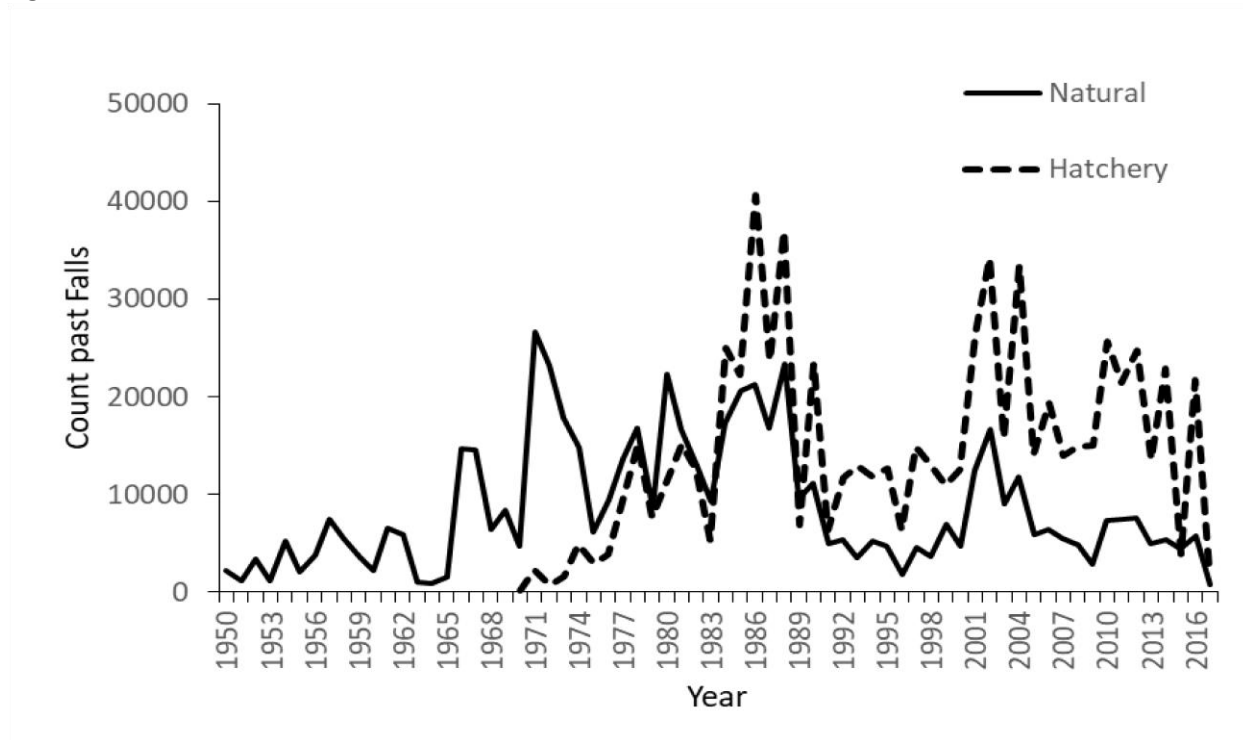
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704 Figure 1



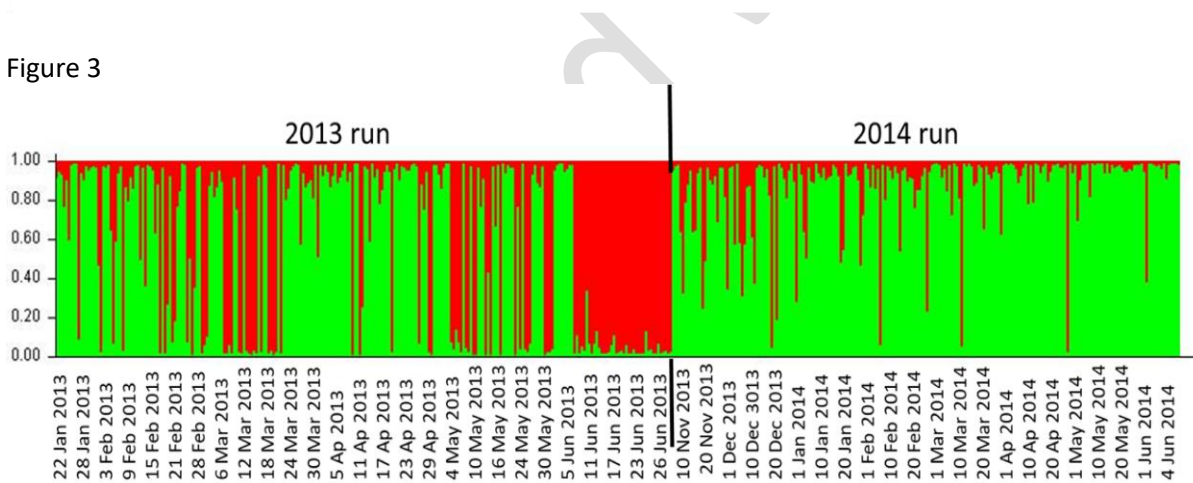
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708 Figure 2



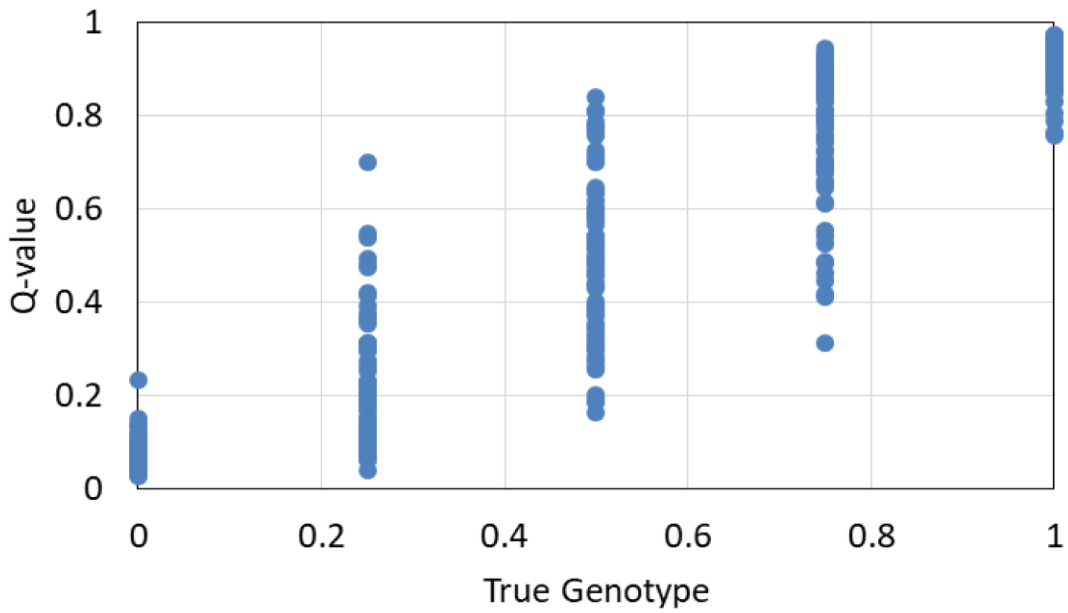
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Figure 3

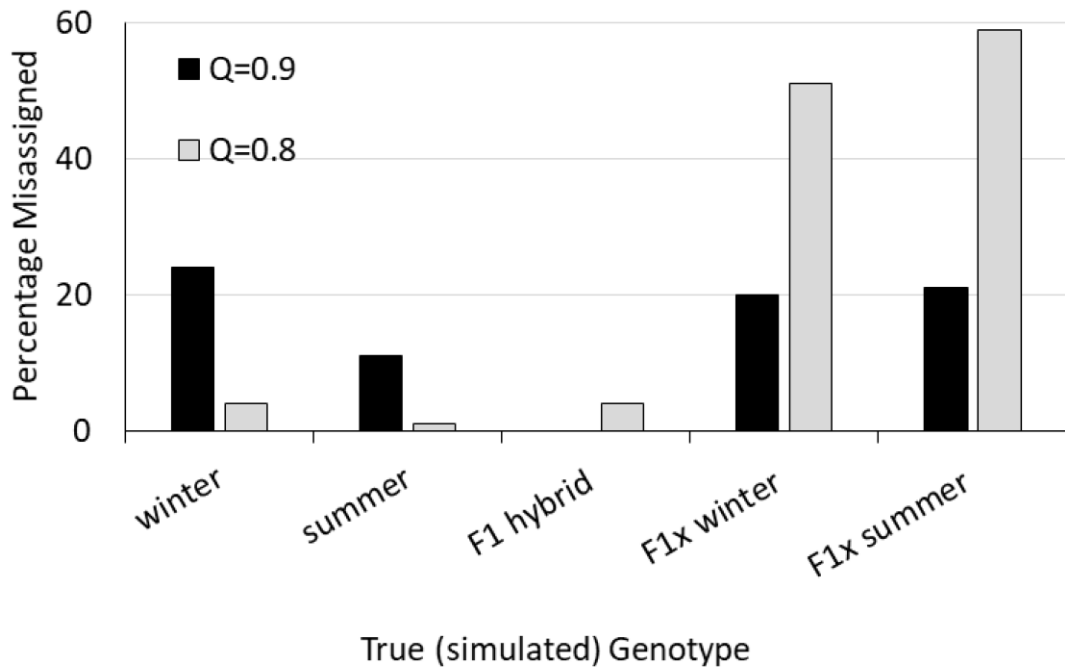


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715 Figure 4

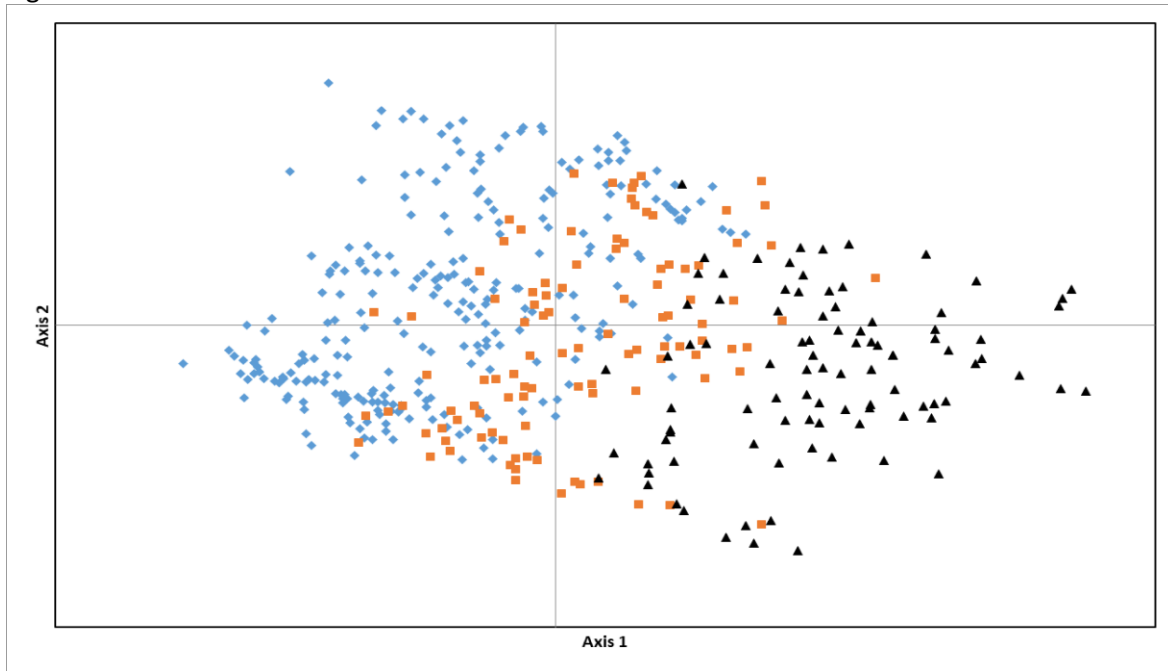


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718 Figure 5



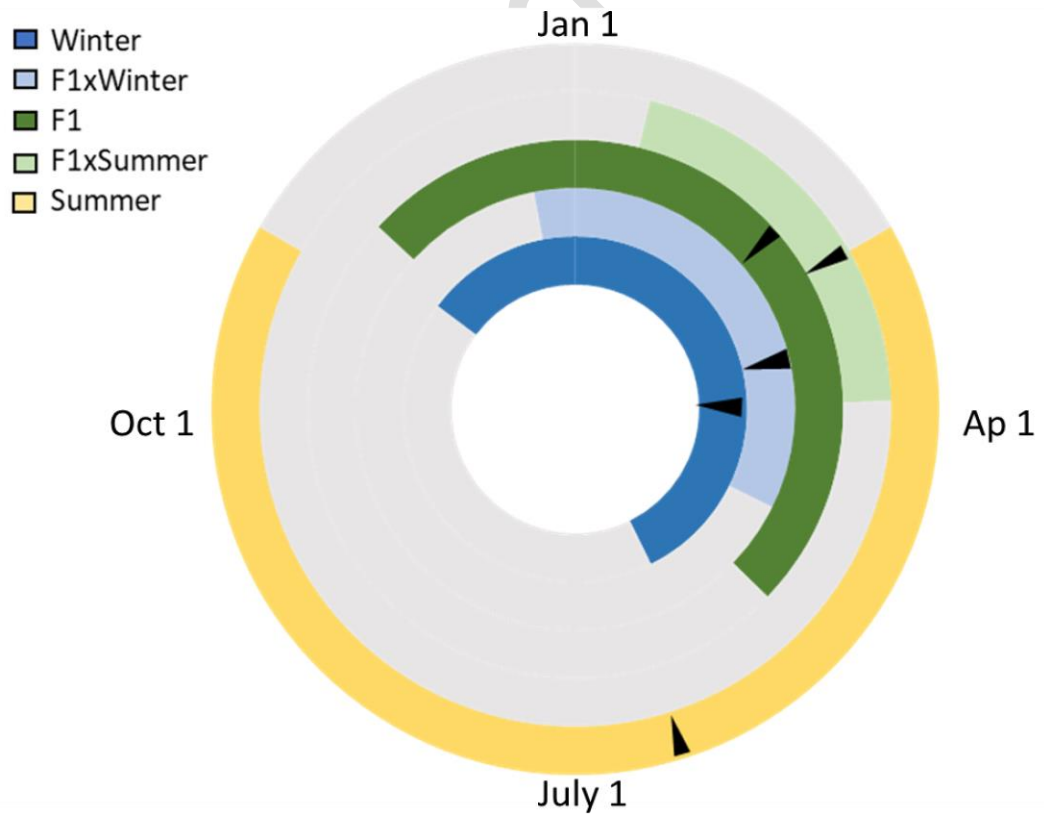
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722 Figure 6



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Figure 7



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