DISTRIBUTION, ABUNDANCE, AND GENETIC POPULATION STRUCTURE OF WOOD RIVER SCULPIN, COTTUS LEIOPOMUS

Kevin A. Meyer^{1,3}, Daniel J. Schill¹, Matthew R. Campbell², and Christine C. Kozfkay²

ABSTRACT.-The Wood River sculpin Cottus leiopomus is endemic to the Wood River Basin in central Idaho and is a nongame species of concern because of its limited distribution. However, status and genetic population structure, 2 factors often central to the conservation and management of species of concern, have not been assessed for this species. We used backpack electrofishers to survey streams that were small enough (i.e., <10 m wide) to collect quantitative distribution and abundance data for Wood River sculpin, and we used mitochondrial DNA control region sequencing to investigate the distribution of genetic variation across the species' range. Of the 102 study sites surveyed, 20 sites (20%) had Wood River sculpin present, including 50%, 15%, and 0% of the sites predicted a priori to contain, possibly contain, or not contain the sculpin, respectively. Native redband trout Oncorhynchus mykiss gairdneri were present at 21 study sites, including 18 of the 20 sites that contained Wood River sculpin. Sixty-one study sites (60%) were dry or had too little water to contain any fish. We estimated that a minimum of 1.36 million Wood River sculpin (≥20 mm total length) currently reside in the basin. The presence of Wood River sculpin was positively associated with stream width:depth ratio and percent cobble/boulder substrate and negatively associated with stream gradient. Mitochondrial DNA haplotype differences were observed between and within the 3 major river subbasins supporting sculpin, with the most striking differences observed between populations in the Camas Creek subbasin and the other 2 subbasins, among which no haplotypes were shared, suggesting relatively long-term isolation. Our results suggest that the Wood River sculpin remains relatively widespread and abundant within its endemic range, despite obvious changes in historical stream connectivity caused by irrigation diversions and other chronic habitat alterations.

Key words: Cottidae, population status, occupancy, abundance, genetic population structure, conservation, endemic species.

The Wood River sculpin Cottus leiopomus is 1 of 8 sculpin species found in Idaho (Simpson and Wallace 1982). Although some of Idaho's sculpin species are widely distributed throughout the state, the Wood River sculpin is endemic only to the Wood River Basin, with 3 associated subbasins, the Big Wood River, the Little Wood River, and Camas Creek. The Wood River sculpin is a benthic species that inhabits flowing water, from small streams to large rivers, and is often found in sympatry with native redband trout Oncorhunchus mykiss gairdneri. The extent of this sympatry has not been investigated, but it likely occurs because of similar requirements of clean, cool water and coarse substrate (gravel and larger) for spawning and rearing of both species.

Wood River sculpin are recognized as a nongame species of concern by the Idaho Department of Fish and Game (IDFG 2006) because of their naturally restricted distribution and recent habitat fragmentation and alterations, the latter of which stem from irrigation diversions and reservoirs, floodplain encroachment, livestock grazing, and stream channelization. Despite the protected status of Wood River sculpin, little is known of the species' status or genetic population structure, 2 factors central to the conservation and management of native fish populations. In fact, to our knowledge, no publications exist on this species except a few unpublished agency reports. Although the Wood River sculpin is most likely a sedentary species like other stream-dwelling sculpin (Mc-Cleave 1964, Hill and Grossman 1987, Petty and Grossman 2004, Schmetterling and Adams 2004), the large number and size of irrigation withdrawals and diversions in the basin (Thurow 1988, Megargle 1999) could exacerbate the lack of population connectivity and threaten the existence of populations lower in the basin or at the lower end of streams.

In light of the scarcity of existing data for this species, the main objective of this study was to evaluate the current distribution and abundance of the Wood River sculpin throughout

¹Idaho Department of Fish and Game, 1414 E. Locust Lane, Nampa, ID 83709.

²Idaho Department of Fish and Game, 1800 Trout Road, Eagle, ID 83616. ³E-mail: kmeyer@idfg.idaho.gov



Fig. 1. Location of study sites where Wood River sculpin and redband trout were present or absent and where genetic samples were collected in 2003 in the Wood River Basin, Idaho. Numbers correspond to genetic sample locations in Table 5. Fifth-order and larger streams that were excluded from analyses (see methods) are shown as wider gray lines compared to other stream segments.

its historical range. To assess the extent of sympatry between Wood River sculpin and redband trout, we also collected distribution and abundance data on trout. We measured a variety of habitat conditions to assess what factors might influence the distribution and abundance of the Wood River sculpin across its range. Finally, we conducted a preliminary survey of the genetic structure of Wood River sculpin populations using mitochondrial DNA (mtDNA) data.

METHODS

Study Area

The Wood River is a tributary of the Snake River in central Idaho, which is composed of 3 major subbasins: the Big Wood River, the Little Wood River, and Camas Creek. Historical processes of both volcanism and glaciation potentially contributed to the isolation of fish populations in the Wood River Basin. First, the basin is not directly connected with the Snake River because a 20-m waterfall known as Malad Gorge Falls is located about 5 km from the mouth and has created a long-term (i.e., tens of thousands of years) barrier to fish migration (Fig. 1). In addition, the Big Wood River was historically connected to the Little Wood River much farther up in the drainage through the present day Silver Creek Valley (Jankovsky-Jones 1997). It has only been since the late Quaternary Period, following a series of lava flows, dam creation, lake formation, and subsequent glacial deposition and dam breaching, that the Big Wood River changed to its present course and connection with the Camas Creek subbasin (Jankovsky-Jones 1997). Two dams with no fish passage further isolate fish within the basin (Fig. 1): Magic Reservoir in the Big Wood River subbasin (built in 1910) and Little Wood Reservoir in the Little Wood River subbasin (built in 1939).

The basin contains over 6000 km of stream covering 7778 km² of semiarid valleys and mountainous headwaters that range from 837 m to over 3600 m in elevation. Vegetation includes western spruce-fir and pine forests at upper elevations and sagebrush steppe at low elevations. Precipitation is mostly in the form of winter snowpack and ranges from 18 cm in the lower valleys to 64 cm in the mountains. Discharge is driven by snowmelt and peaks between April and June, but it is modified by numerous irrigation diversions in mainstem and tributary streams. Native species in the Wood River Basin include Wood River sculpin, redband trout, mountain whitefish Prosopium williamsoni, bridgelip sucker Catostomus columbianus, largescale sucker Catostomus macrocheilus, Utah chub Gila atraria, longnose dace Rhinichthys cataractae, speckled dace Rhinichthys asculus, and redside shiner Richardsonius balteatus. Although Paiute sculpin Cottus *beldingii* have been reported from the lower Little Wood River (Merkley and Griffith 1993), their distribution is sparse, and it is unclear if they are native to the basin. Our samples did not contain any Paiute sculpin, which can be differentiated from Wood River sculpin by the presence of preopercular spines (Simpson and Wallace 1982). Introduced species that have established self-sustaining populations include brook trout Salvelinus fontinalis and brown trout Salmo trutta.

Study Site Selection

Data collection occurred from July to October 2003. Spatially balanced randomly selected study sites were generated with the help of the Environmental Protection Agency's Environmental Monitoring and Assessment Program (Fig. 1). The technique maps 2-dimensional space (in our study, a 1:100,000 scale hydrography layer) into 1-dimensional space with defined, ordered spatial addresses and uses restricted randomization to randomly order the spaces. Systematic sampling of the randomly ordered spaces results in a spatially balanced sample (Stevens and Olsen 2004). The study design and sampling frame was stratified in 2 ways to reduce overall variation in the abundance data and make estimates more accurate. First, all stream reaches were categorized *a priori* as either (1) likely to contain, (2) not likely to contain, or (3) possibly containing Wood River sculpin. This was based on professional knowledge of the geomorphology and anthropogenic alterations in the basin and on the suspected distribution of Wood River sculpin in the basin. Stream order (Strahler 1964) was used as a 2nd stratification.

Fish and Habitat Sampling

At each study site, fish were captured using electrofishing gear. Depending on stream width, crews of 2–4 people performed multiple-pass electrofishing removals (2–4 passes) using 1 or 2 gas-powered backpack electrofishing units. Pulsed DC was used, with pulses of 3–5 ms, 500–900 volts, and 30–60 Hz. Sampling occurred during low to moderate flow conditions (i.e., after spring runoff and before the onset of winter) to facilitate effective fish capture and standardize sampling conditions. Block nets were placed at the lower and upper ends of the reach to prevent fish from moving out of the sampling reach during electrofishing.

All fish were retained, identified, and enumerated, but in our study streams, nongame species other than Wood River sculpin (i.e., longnose dace, speckled dace, redside shiner, and bridgelip sucker) were caught at fewer than 7 sites and thus were encountered too infrequently to include in our analyses. Consequently, only Wood River sculpin and trout data are reported herein. Fish were measured to the nearest millimeter (total length = TL) and gram, and released. Study sites were typically (91% of the time) between 80 and 120 m in length (depending on habitat types, riparian vegetation, and our ability to place block nets) and averaged 96 m (range 45–130 m).

Maximum-likelihood abundance and variance estimates for Wood River sculpin and trout were calculated for each study site with the MicroFish software package (Van Deventer and Platts 1989). Wood River sculpin abundance estimates could not be obtained from large (i.e., >10 m in width; 5th-order streams and larger) mainstem reaches of the Wood River Basin. These reaches totaled 497 km in stream length (8% of all stream kilometers) and were excluded from our analyses, despite the fact that sculpin and trout were often present in these reaches. Consequently, this study focuses only on the small (<10 m in width) tributary streams and upper ends of the mainstem rivers.

Because electrofishing is known to be size selective (Sullivan 1956, Reynolds 1996), Wood River sculpin were separated into 2 length categories (<60 mm TL and \geq 60 mm TL), and trout were similarly separated into 2 length categories (<100 mm TL and \geq 100 mm TL). Abundance estimates were made separately for these size groups and summed for an overall estimate. Capture efficiencies for sculpin <20 mm and for trout <40 mm were especially low relative to larger fish; thus abundance values for sculpin <60 mm and trout <100 mm were probably underestimated (Peterson et al. 2004).

At each study site, we determined elevation (m) from U.S. Geological Survey 1:24,000 topographic maps using universal transverse mercator (UTM) coordinates obtained at the lower end of the reach that was electrofished. Stream order was determined from a 1:100,000 stream hydrology scale. Gradient (%) was determined using the software package All Topo Maps, version 2.1 for Windows (iGage Mapping Corporation, Salt Lake City, UT); the distance (m) between the 2 contour lines that bounded the study site was traced (average traced distance was about 1 km), and gradient was calculated as the elevational increment between those contours divided by the traced distance. We measured conductivity ($\mu S \cdot cm^{-1}$) with a calibrated handheld conductivity meter accurate to $\pm 2\%$. We noted land ownership as private or public.

Ten equally spaced transects were established within each study site, and from those transects, the remaining measurements were taken. Stream width (m) was calculated from the average of all transect readings. At each transect, depth was measured at one-fourth, one-half, and three-fourths of the distance across the channel, and the sum of the measurements was divided by 4 to account for zero depths at the stream margins for trapezoidalshaped channels (Platts et al. 1983, Arend 1999). Percent substrate composition was visually estimated as the percent of substrate within 1 m of each transect that was silt (<0.06 mm), sand (0.06–1.99 mm), gravel (2–63 mm), cobble (64–256 mm), boulder (257–4096 mm), or bedrock (>4097 mm). The percentage of unstable banks and the percentage of stream shading were also visually estimated at each transect. We averaged all habitat measurements across all transects for an overall mean for each study site.

Estimation of Fish Abundance

We estimated total Wood River sculpin and trout abundances separately for each subbasin using the stratified random sampling formulas from Scheaffer et al. (1996). As previously mentioned, estimates were stratified by stream order (1st through 4th) and by *a priori* distribution categorization for sculpin and trout (i.e., likely present, likely absent, and unknown); thus, there were 12 strata (L). We first summed the total length of stream within each stratum using the ArcView[®] geographic information system (GIS). The total number of sampling units in each stratum (N_i) was calculated by dividing the stream length by our typical sampling reach length (100 m). Abundance estimates were standardized to density per 100 linear meters of stream. We calculated a mean abundance (\overline{y}_i) within each stratum and an associated variance. For total population size (N_{census}) , we used the formula

$$(N_{census}) = \sum_{i=1}^{L} N_i \overline{y}_i$$

and for variance of N_{census} we used the formula

$$V(N_{census}) = \sum_{i=1}^{L} N_i^2 \left(\frac{N_i - n_i}{N_i}\right) \left(\frac{s_i^2}{n_i}\right)$$

where s_i^2 is the variance of the observations in stratum *i*, and n_i is the sample size within stratum *i*. From this we calculated 90% confidence intervals around the abundance estimates. All sample sites, including dry and fishless sites, were included in these estimates.

Fish-Habitat Relationships

We assessed whether stream characteristics that we measured were related to Wood River sculpin distribution and abundance. To do this we first removed all sites that were dry or had too little water to contain any fish. We then plotted all potential independent variables against presence-absence and abundance data in order to look for abnormalities and nonlinearity in the data. Both log and square-root transformations were needed to normalize the stream gradient data for comparison to presenceabsence and abundance data, respectively. Percent fine substrate could not be normalized with any data transformations and was therefore removed from further analyses. No other transformations were needed. Multicollinearity between independent variables was assessed with correlation analysis, but none was detected. Because width was used in the calculation of a standardized fish density metric (fish \cdot m⁻²), correlations between density and width or between density and width:depth ratio were not estimated.

To assess the relationship between stream attributes and the presence/absence of Wood River sculpin, we first calculated a mean and 95% confidence interval for stream conditions with and without Wood River sculpin, with statistically significant differences determined by nonoverlapping confidence intervals (Johnson 1999). We then related stream attributes to Wood River sculpin occupancy using logistic regression analyses, adding stream attributes to the model in an iterative process and using Akaike's Information Criteria (AIC) to assess the best model. AIC is an extension of the maximum-likelihood principle with a bias correction term that penalizes for added parameters in the model (Akaike 1973). Lower AIC values indicate better-fitting models. For comparison we also calculated an adjusted R^2 for discrete models (Nagelkerke 1991). To assess the relationship between stream attributes and density of Wood River sculpin, we used multiple regression analyses, adding stream attributes to the model in an iterative process and using R^2 to assess the best model.

Genetic Population Structure

We chose mitochondrial DNA (mtDNA) analysis for this assessment because the high mutation rate, strict maternal inheritance, and low effective population size of mtDNA make it a useful tool for investigating the population structure of recently diverged or closely related taxonomic groups (Avise 1994). Additionally, the mtDNA control or "D-loop" gene region has already been successfully amplified and sequenced in several species of sculpin, and has demonstrated differences at the subbasin level between populations of fluvial sculpin *Cottus nozawae* (Yokoyama and Goto 2002) in Japan, as well as shorthead sculpin *Cottus confusus* and mottled sculpin *Cottus bairdi* in Idaho (IDFG, unpublished data).

Ten to 22 nonlethal fin clips from Wood River sculpin (total n = 187) were collected from 15 sample locations within the Wood River Basin (Table 5), but not necessarily the same locations as the fish abundance study sites (Fig. 1). Total genomic DNA was extracted from a 1×1 -mm piece of fin clip following methods described by Campbell (2000). DNA was resuspended in 100 µL TE. Approximately 450 bp of the mtDNA control region were amplified in a 20-uL reaction consisting of 1 uL DNA extract (approximately 2.5 ng $\cdot \mu L^{-1}$), 2.0 μL 10X buffer (Perkin Elmer), 2.0 μL MgCl₂, 1.6 µL bovine serum albumin, 1.0 µL of each primer (L/5'-TTC CAC CTC TAA CTC CCA AAG CTA G-3' [Lee et al. 1995] and R/5'-CCT GAA GTA GGA ACC AGA TG-3' [Meyer et al. 1990]), 1.6 µL 10.0 mM dNTPs (10mM each of dATP, dCTP, dGTP, and dTTP), 0.05 µL Perkin-Elmer Taq polymerase, and 9.3 µL dH₂O. Polymerase chain reaction conditions consisted of an initial denaturing cycle of 94 °C for 3 minutes, followed by 32 cycles of denaturation at 94 °C for 45 seconds, annealing at 52 °C for 45 seconds, and extension at 72 °C for 1 minute, with a final extension at 72 °C for 3 minutes.

Sequencing reactions were performed with a BigDye Terminator Cycle Sequencing Ready Reaction Kit (Applied Biosystems, version 3.1, http://www3.appliedbiosystems.com) using both the forward and reverse primers. Sequenced products were cleaned using Edge BioSystems gel filtration plates (Edge Biosystems, http: //www.edgebio.com) and were run out on a Prism 3730 DNA sequencer (Applied Biosystems). Sequences were edited using Sequencher (Gene Codes Corporation, version 4.1.2, http://www.genecodes.com), and the consensus sequences (394 bp) were aligned using the Clustal W program (Thompson et al. 1994) within the software program MEGA3 (Molecular Evolutionary Genetics Analysis, version 3.0; Kumar et al. 2004).

Differences in mtDNA haplotype frequency distributions among the 3 major subbasins were tested for significance using a Monte Carlo chi-square simulation (Roff and Bentzen 1989). Sequence divergence (p-distance) between haplotypes and a maximum-parsimony (MP) consensus tree depicting relationships between

Stream network and study sites	Big Wood River	Camas Creek	Little Wood River	Total
Total stream km in Wood River drainage	2847	1418	1738	6003
Stream km categorized <i>a priori</i> as "WRS present"	244	60	211	515
Stream km categorized <i>a priori</i> as "WRS unknown"	1455	470	746	2671
Stream km categorized a priori as "WRS absent"	1148	888	781	2817
Total number of sites sampled	52	15	35	102
Sites containing fish	19	4	9	32
Sites within "WRS present" streams	10	4	6	20
that contained WRS	7	2	1	10
Sites within "WRS unknown" streams	35	8	23	66
that contained WRS	5	0	5	10
Sites within "WRS absent" streams	7	3	6	16
that contained WRS	0	0	0	0
Sites containing redband trout	12	3	6	21
Dry or nearly dry sites	29	9	23	61

TABLE 1. Stream network (at the 1:100,000 scale) and distributional extent of Wood River sculpin (WRS) from study sites in 1st- to 4th-order streams sampled in 2003 within 3 subbasins of the Wood River, Idaho.

haplotypes were obtained from aligned sequences using MEGA3 (Kumar et al. 2004). The MP tree was obtained using the closeneighbor-interchange algorithm with search level 3, in which the initial trees were obtained with the random addition of sequences (100 replicates). All positions containing gaps and missing data were eliminated from the dataset (complete deletion option). To assess nodal strength, we performed 1000 bootstrap replicates. Since Paiute sculpin have been tentatively identified in the lower Little Wood River (Merkley and Griffith 1993) and resolved as a sister species to Wood River sculpin (Kinziger et al. 2005), we included a D-loop sequence from a Paiute sculpin (University of Alabama Ichthyological Collection [UAIC] 11528). We also included a D-loop sequence from a Wood River sculpin sample that had previously been sampled and sequenced (D.A. Neely unpublished data) from the Big Wood River and archived at the UAIC (11631). Two additional D-loop sequences from torrent sculpin *Cottus* rhotheus (UAIC 11670) and mottled sculpin Cottus hubbsi (UAIC 11633), collected in Idaho, were included for reference purposes (D.A. Neely, unpublished data).

RESULTS

Wood River sculpin distribution and abundance was determined at 102 study sites in 1st- to 4th-order streams throughout the Wood River Basin, 61 of which were dry or had too little water to sustain any fish (Table 1). Fiftytwo of the study sites were on private property (65% of which were dry or nearly dry), whereas 50 sites were on public land (52% of which were dry or nearly dry).

The *a priori* categorization of Wood River sculpin distribution was reasonably accurate, given that sculpin were present at 10 of 20 sites (50%) where we expected to find them, 0 of 16 sites where we expected them to be absent, and 10 of 66 sites (15%) where we were unsure if they would be present (Table 1). Redband trout were found at 18 of 20 sites (90%) where Wood River sculpin were found, and 3 of 82 sites (4%) where Wood River sculpin were not found.

Estimation of Fish Abundance

Estimated abundance of Wood River sculpin in Idaho, including only 1st- through 4th-order streams, was 1,356,600 ± 594,660 (90% confidence interval; Table 2). Forty-six percent of that abundance came from areas that were identified *a priori* as likely to contain Wood River sculpin, whereas 54% of that abundance was estimated to be in streams where we were unsure if Wood River sculpin would be present. Comparatively, we estimated that there were 175.630 ± 97.200 redband trout in the Wood River Basin in 1st- through 4th-order streams (Table 2). Among all study sites (including those that were dry or had no fish), average linear abundance and areal density of Wood River sculpin were 0.40 individuals · m⁻¹ and 0.09 individuals \cdot m⁻², respectively. At only those sites where sculpin were present, average linear abundance and areal density were 2.04 individuals \cdot m⁻¹ and 0.48 individuals \cdot m⁻², respectively.

Species	Size group	Occurrence category	N_{census}	90% confidence interval
Wood River sculpin	≥60 mm	Present	402,313	89,977-714,652
		Absent	0	
		Unknown	550,461	126,130-974,792
	<60 mm	Present	226,533	24,915-428,151
		Absent	0	
		Unknown	177,293	503-365,333
TOTAL		All	1,356,600	761,940-1,951,260
Redband trout	≥100 mm	Present	55,175	14,044-96,306
		Absent	0	
		Unknown	21,297	9427-33,167
	<100 mm	Present	18,602	54-42,982
		Absent	0	
		Unknown	80,556	156-164,346
TOTAL		All	175,630	78,430-272,830

TABLE 2. Estimated number of Wood River sculpin and redband trout present in 2003 in 1st- to 4th-order streams of the Wood River Basin, Idaho. Ninty-percent confidence intervals are given.

TABLE 3. Average stream conditions and associated confidence intervals (CIs) at sites with (n = 20) and without (n = 23) Wood River sculpin in the Wood River Basin, Idaho.

	With Wood I	River sculpin	Without Wood River sculpin		
Variable	Mean	95% CI	Mean	95% CI amplitude	
Stream order	2.7	0.4	2.1	0.6	
Elevation (m)	1944.9	89.0	2010.4	171.9	
Gradient (%)	2.5	0.8	6.6	3.3	
Conductivity ($\mu S \cdot cm^{-1}$)	220.0	55.0	207.0	55.0	
Width (m)	4.4	1.1	2.8	0.6	
Depth (m)	0.13	0.02	0.16	0.06	
W:D ratio	32.5	4.8	22.6	4.7	
Percent fine substrate	0.5	0.3	1.4	0.7	
Percent cobble/boulder substrate	3.6	0.7	2.5	0.9	
Percent shading	1.6	0.4	1.7	0.6	
Percent unstable banks	0.5	0.4	0.6	0.4	
Redband trout density (number \cdot m^{-1})	0.16	0.10	0.05	0.10	

Fish-Habitat Relationships

At sites that contained enough water for fish to be present (n = 43), there appeared to be relationships between certain stream conditions and the distribution of Wood River sculpin (Table 3). Most notably, the likelihood of Wood River sculpin presence increased in a stream as width: depth ratio and percent cobble/ boulder substrate increased and gradient decreased (Fig. 2). The inclusion of all 3 variables in a logistic regression model gave the best AIC score (46.4) and explained 52% of the total variation in Wood River sculpin occupancy (using the adjusted R^2 for discrete models). The next-best model was a 2-variable model that included gradient and cobble/boulder substrate (AIC = 49.5) and explained 41% of the variation in Wood River sculpin occupancy.

At sites that contained Wood River sculpin (n = 20), there was little correlation between any stream conditions and Wood River sculpin density (Table 4). Average depth was most strongly correlated to density, but inclusion of depth and stream order in a multiple regression model explained only 30% of the variation in Wood River sculpin density, whereas depth alone explained 28% of the variation in sculpin density. For redband trout, density was most positively correlated to percent stream shading and most negatively correlated to elevation (Table 4).

Genetic Population Structure

Seven haplotypes (unique DNA sequences) were observed in samples of Wood River sculpin (Table 5). Because previous sequencing in our lab of the D-loop mtDNA region in



Fig. 2. Observed frequency of occurrence (histograms) and probability of occurrence predicted from logistic regression models (dashed lines) relating Wood River sculpin to stream conditions in the Wood River Basin, Idaho. The centers of the histograms are the midpoints of the bins used in the frequency distributions. *P*-values are from the combined logistic regression model that included all 3 stream variables (see text).

other sculpin species has yielded 13 haplotypes, the new haplotypes were labeled Cottus_14 through Cottus_20 (GenBank accession EU369333–EU369352).

Statistically significant haplotype frequency differences were observed between the 3 subbasins (P < 0.005, $\chi^2 = 197$). The Cottus_14 haplotype was most common (66.3%), occurring in all sample locations within the Big Wood and Little Wood river subbasins (Table 5). This haplotype was fixed within 8 of 11 sample locations in the Big Wood and Little Wood river subbasins and was the dominant haplotype in the other 3 locations. The Cottus_15, 16, and 20 haplotypes were present in the Big Wood River subbasin and Cottus_19 in the Little Wood River subbasin, but all 4 were found at low frequencies and only at sin-

TABLE 4. Correlations (*r*) between stream attributes and density (fish \cdot m⁻²) of Wood River sculpin (WRS) and redband trout (RBT) at study sites surveyed in 2003 in the Wood River Basin, Idaho.

	Der	nsity
Stream attribute	WRS	RBT
Gradient (%)	-0.30	-0.12
Stream order	0.47	-0.11
Elevation (m)	-0.16	-0.35
Conductivity ($\mu S \cdot cm^{-1}$)	0.22	0.13
Depth (m)	0.53	0.07
Percent cobble/boulder substrate	0.23	-0.20
Percent unstable banks	-0.07	-0.08
Percent shading	0.04	0.43
Redband trout density	-0.01	_

gle locations. In contrast, Cottus_14 was not observed within any of the sampling locations within the Camas Creek subbasin, where instead the most common haplotype was Cottus_17 (63.2%). The Cottus_18 haplotype was also observed, but only in 2 sample locations in 1 stream. Sequence divergence between the 7 haplotypes observed within the Wood River Basin ranged from 0.3% to 1.0% (data not shown). Sequence divergence between these haplotypes and the reference Paiute sculpin sequence ranged from 5.9% to 6.4%.

All haplotypes observed in the Wood River Basin and the reference Wood River sculpin sequence (UAIC 11631) clustered together in the MP consensus tree (84% bootstrap support; Fig. 3). Within this group, there were no haplotypes that clustered together with greater than 50% bootstrap support. As expected, the reference Paiute sculpin sample (11528) grouped at a basal position to the Wood River sculpin samples (99% bootstrap support).

DISCUSSION

Our results suggest that Wood River sculpin are somewhat limited in distribution and reside in disconnected populations within their historical range in the Wood River Basin. In fact, Wood River sculpin were captured in <20% of the study sites and were estimated to occupy only 11% of the 1st- through 4th-order stream kilometers in the basin. Much of the reason for their meager distribution was that 60% of the sites we surveyed were dry or nearly dry during our sampling period. This arid condition is due in part to irrigation diversions over the last 150 years but also to natural hydrologic conditions common in the Wood River

TABLE 5. Sample number (corresponding to Fig. 1) and location, polymorphisms (associated with each haplotype),
haplotype, and haplotype frequency at genetic sample locations in the Camas Creek (CA), Big Wood River (BW), and	d
Little Wood River (LW) subbasins in the Wood River Basin, Idaho.	

				Haplotype frequency						
Polymorphisms 129 b.p. 185 b.p.		C G	T G	T G	C G	C G	C A	A G		
			220 b.p.	_	_	-	_	Α	_	_
			281 b.p.	С	Т	С	С	С	С	С
			290 b.p.	Т	С	С	С	С	Т	Т
Sar	nple n	umber and location		Cottus_14	Cottus_15	Cottus_16	Cottus_17	Cottus_18	Cottus_19	Cottus_20
1	CA	Soldier Creek					22			
2	\mathbf{CA}	Phillips Creek					13			
3	\mathbf{CA}	Willow Creek					1	11		
4	CA	Upper Willow Creek	ζ.					10		
5	BW	Big Wood River		12						
6	BW	Warm Springs Creek	c C	11						1
7	BW	Trail Creek		12						
8	BW	Silver Creek		10						
9	BW	Gladiator Creek		9	2	1				
10	LW	Little Wood River		14						
11	LW	Iron Mine Creek		10						
12	LW	Friedman Creek		10						
13	LW	Muldoon Creek		13					2	
14	LW	Little Copper Creek		10						
15	LW	Porcupine Creek		13						



Fig. 3. Maximum-parsimony consensus tree of observed Wood River sculpin mitochondrial DNA haplotypes and 3 reference sequences. Bootstrap values (see text) are shown when >50.

basin (Castelin and Chapman 1972). Thus distribution and connectivity of fish in this basin was probably always naturally restricted at some level (see also Meyer et al. 2006), and it is difficult to estimate how much historical 1st- through 4th-order stream habitat is no longer occupied. There are some museum records of Wood River sculpin that reveal historical occupancy, but those we are aware of that were collected from locations downstream of their current extent of occupancy are limited to the mainstems of the Big Wood and Little Wood rivers.

Regardless of the exact historical distribution of Wood River sculpin, we found during our surveys that where flowing water was present in the basin, Wood River sculpin were also typically present and usually were quite abundant. For instance, at the 32 sites that supported at least 1 species of fish, Wood River sculpin were present 63% of the time. Moreover, they were nearly as widely distributed as and far more abundant than redband trout, which are considered to be robust in this basin (Thurow 1988). Although Wood River sculpin were over 7 times more numerous than redband trout, at sites where both were present, Wood River sculpin biomass averaged only 23.5 kg \cdot ha⁻¹ compared to 208.3 kg \cdot ha⁻¹ for redband trout. Previous studies have shown similar relationships between trout and sculpin density and biomass (Neves and Pardue 1983, Erman 1986).

Despite the somewhat restricted distribution of Wood River sculpin in the basin, several factors suggest that this is not indicative of a population at risk. First, true distribution and abundance were unavoidably underestimated in our study because sampling efficiency constraints in large river segments (i.e., 5th-order and higher) precluded inclusion of these segments, though sculpin were captured in many of these reaches when the reaches were sampled independently for salmonid populations (K. Meyer unpublished data). Second, depletion electrofishing typically underestimates both fish distribution (Bayley and Peterson 2001, Reynolds et al. 2003) and abundance (Riley and Fausch 1992, Peterson et al. 2004), especially for small fish such as sculpin. Furthermore, because of their sedentary nature, stream-dwelling sculpin already tend to naturally be fragmented into disconnected populations (Hill and Grossman 1987, Goto 1998, Petty and Grossman 2004). Not only were Wood River sculpin more abundant than redband trout, but compared to estimates for Yellowstone cutthroat trout O. clarkii bouvieri, another species of conservation interest in southern Idaho, Wood River sculpin were 5.5 times more abundant per square kilometer (cf. Mever et al. 2006). Yellowstone cutthroat trout and Wood River sculpin have similar longevity, usually living to a maximum of 6 years of age, and both mature at between 2 and 4 years of age, but Wood River sculpin are far less fecund (see Meyer et al. 2003, 2008).

Wood River sculpin were almost always present when redband trout were present and almost always absent when redband trout were absent (Fig. 1), but it is unlikely that there was any causative effect between the species. In fact, at sites containing Wood River sculpin, their density and that of redband trout were not correlated (r = -0.01), and habitat variables that were correlated to fish density were different between species. The most likely explanation is that both species require similar basic habitat requirements such as cold water for metabolic function and coarse substrate for spawning and rearing. Our results suggest that, once these basic needs are met, additional factors affecting distribution and abundance differ between the 2 species, as for other sympatric populations of salmonids and sculpin (e.g., Prenda et al. 1997, Inoue and Nakano 2001). Competition between sculpin and trout sometimes occurs (Brocksen et al. 1968, Hesthagen and Heggenes 2003, Ruetz et al. 2003), but habitat selection shifts can minimize potential competition between sculpin and trout (Prenda et al. 1997).

Few stream conditions we measured were strongly correlated to Wood River sculpin distribution or abundance. Nevertheless, considering their benthic nature and need for cobble/ boulder substrate for spawning and rearing, we were not surprised to find that Wood River sculpin were more likely to occur where gradient was low and cobble/boulder substrate was prevalent. Meyer et al. (2008) concluded that Wood River sculpin growth and survival also was inversely related to gradient. The fact that fish habitat models usually are region specific and difficult to transfer to areas outside the original study area (Fausch et al. 1988) is largely extraneous to this study, because Wood River sculpin exist only in the Wood River Basin. Nonetheless, our results should be viewed as introductory, and more research on habitat requirements and preferences for Wood River sculpin may further explain their current distribution and abundance. While numerous studies have identified specific habitat preferences for stream-dwelling sculpin (e.g., Brown 1991, van Snik Gray and Stauffer 1999, Inoue and Nakano 2001), others have concluded that microhabitat and macrohabitat conditions were of little importance and instead suggested that sculpin population dynamics were regulated most strongly by density-dependent processes such as competition for food and space (Petty and Grossman 1996, Grossman et al. 2006).

Results from this study indicate significant haplotype frequency differences both among and within the 3 major river subbasins supporting Wood River sculpin. The most striking difference was observed between the Camas Creek subbasin and the other 2 subbasins, in which no haplotypes were shared, although there was not clear phylogenetic support for this geographical subdivision. Significant genetic differences were also observed within the Camas Creek subbasin, with Willow Creek exhibiting a predominant haplotype not found in the other 2 sample locations. These results suggest that there is very limited gene flow between populations in the Camas Creek subbasin and populations in the Big Wood and Little Wood river subbasins. Similar conclusions can be drawn between Willow Creek and the other 2 sample locations within the Camas Creek subbasin.

Fewer genetic differences were observed between sample locations in the Big Wood and Little Wood river subbasins, with one haplotype dominant throughout these subbasins. Several additional haplotypes were also observed in low frequency within Gladiator Creek and Warm Springs Creek in the Big Wood River subbasin and in Muldoon Creek in the Little Wood River subbasin. While the majority of the sample locations in the Big Wood and Little Wood river subbasins shared the same haplotype, the lack of haplotype diversity within these subbasins prevented any conclusions from being drawn regarding current levels of gene flow among these populations. However, as previously mentioned, other species of sculpin have low dispersal distances (<50 m; Hill and Grossman 1987, Goto 1998, Petty and Grossman 2004) and exhibit restricted gene flow over short distances (Yokovama and Goto 2002, Lamphere 2005), and it is likely that gene flow between populations of Wood River Sculpin within subbasins is similarly low.

There is no evidence to suggest that the differences observed between the Camas Creek subbasin and the other 2 subbasins were the result of sampling separate species of sculpin. This has been a concern in other drainages in Idaho where some sculpin species have been very difficult to distinguish morphologically (D. Zaroban, Albertson College Museum Fish Curator, personal communication). Sequencing of the mtDNA D-loop region in fluvial sculpin in Japan indicated that intraspecific sequence divergence at the basin level ranged between 0.6% and 1.8% (Yokoyama and Goto 2002). In our study, pairwise sequence divergence estimates among the 7 haplotypes observed in the Wood River Basin were relatively low (0.3%-1.0%) and similar in range to what we have observed at the basin level in other species of sculpin in Idaho (M.R. Campbell unpublished data). In contrast, sequence divergence between the reference Paiute sculpin sequence and the Wood River sculpin haplotypes was much higher (5.9%-6.4%).

It is likely that a combination of other factors are responsible for the haplotype differences observed between the Camas Creek subbasin and the other 2 subbasins, including contemporary influences (dispersal behavior, population isolation as the result of anthropogenic influences, and population size) and historical processes. As mentioned previously, a review of the geological history of the Wood River Basin indicates that until the late Quaternary Period (1) the Camas Creek subbasin may have been isolated from both the Big Wood and Little Wood river subasins, and (2) the Big Wood and Little Wood river subbasins may have been directly connected. Both of these geological factors could have contributed to the fixed differences observed between populations in the Camas Creek subbasin and the other 2 subbasins.

In summary, the Wood River sculpin was the most abundant and one of the most widely distributed species of fish in 1st- to 4th-order streams in the Wood River Basin, being present in most locations (63%) where any species of fish were present. Our genetic results provide some evidence of restricted gene flow among populations of Wood River sculpin, which is consistent with the life history and low dispersal of many species of streamdwelling sculpin. Obviously, natural and anthropogenic changes (drought and global warming, habitat alterations, and irrigation diversions) can exacerbate population fragmentation and isolation. The long-term consequences of population isolation (demographic and genetic) for this species will probably depend mostly on population size, which appears to be large, although highly variable, in many streams throughout the species' range. The stability of population size over time should be monitored. Initially, we suggest that any conservation monitoring and management efforts focus on the Camas Creek subbasin, because estimated abundance was lowest in this subbasin and sampled populations from this subbasin contained mtDNA diversity not found in the other 2 subbasins.

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