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## A FIRST SURVEY OF METAZOAN PARASITES IN THE FISHES OF LAKE POWELL, UTAH

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**ABSTRACT.**—Lake Powell (Colorado River drainage, Utah and Arizona, USA) is an important and unique fishery comprising several nonnative fishes. There are no previous studies of the parasites of the fishes of Lake Powell. We provide a general survey of the metazoan parasites found in the numerically dominant fish species of the reservoir. We collected and surveyed for parasites in 236 fishes of 8 species. We found 832 parasites comprising 13 species. All of the parasite species we found are widespread throughout North America and other parts of the world. Six of the 13 species of parasites we found have been documented previously in the Colorado River system. In general, benthic-feeding fishes exhibited higher parasite richness and intensity compared to pelagic-feeding fishes. This study serves as a baseline for parasite community studies in Lake Powell and can provide a comparison for future studies.

**RESUMEN.**—El lago Powell (drenaje del río Colorado, Utah y Arizona, EE.UU.) es una pesquería única e importante compuesta por varios peces no nativos. No hay estudios previos sobre los parásitos de los peces del lago Powell. Proporcionamos un estudio general de los parásitos metazoarios encontrados en las especies de peces numéricamente dominantes de la reserva. Colectamos y estudiamos los parásitos de 236 peces de 8 especies. Encontramos 832 parásitos de 13 especies. Todos los parásitos que encontramos son especies que tienen una distribución amplia en toda América del Norte y en otras partes del mundo. Seis de las trece especies de parásitos han sido documentadas previamente en el sistema del Río Colorado. En general, los peces bentónicos exhibieron mayor riqueza e intensidad de parásitos en comparación con los peces pelágicos. Este trabajo sirve como base para el estudio de las comunidades de parásitos del lago Powell y puede proporcionar una comparación para estudios futuros.

Lake Powell was created in 1963 from the closing of the Glen Canyon Dam on the Colorado River (National Academy of Sciences 1991). It has become a renowned destination for recreational water sports and is considered an exceptional fishery. Nonnative sport fishes dominate the fish fauna of Lake Powell (Maxwell and Thoesen 1965). To effectively manage this system, it is essential to understand the ecological context and ecological interactions of the fishes. Parasites are an important, yet often overlooked, component of the ecology of fishes (Williams et al. 1992, Poulin 1999). Parasite surveys have been published for the Lower Colorado River (Linder et al. 2012) and Little Colorado River (Choudhury et al. 2004, Stone et al. 2007) below Lake Powell; however, there has been no parasite survey for the fishes of Lake Powell.

Parasite occurrence in a given community is determined by presence of appropriate hosts and opportunity for immigration and colonization. However, few consistent patterns exist for community assembly of parasites of

freshwater fishes, making it difficult to predict the expected assemblage of parasites in a wholly nonnative community of fishes, such as that found in Lake Powell (Price and Clancy 1983, Kennedy and Bush 1994, Poulin 1996, 1997). To document diversity and abundance of fish parasites in Lake Powell and the distribution of fish parasites among fish species, we surveyed for metazoan parasites in numerically dominant fish species in 2 separate years.

### METHODS

We surveyed for parasites in April of 2013 and 2015. In 2013, we sampled fishes near Wahweap Bay (36.99768, -111.477110; Lone Rock Canyon, Warm Creek Bay, Gunsight Bay, Antelope Creek), and in 2015 we sampled fishes near Bullfrog Bay (37.506614, -110.724569) and the surrounding canyons (Crystal Springs, Forgotten Canyon, and Moqui Canyon). During both sampling periods, we collected fish by using standard hook-and-line practices along with gill nets to supplement

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TABLE 1. Continued.

Fish species	EFN	cestode						copepod		trematode monogenean			nematode	
		<i>Bothriocephalus claviceps</i>	<i>Carypophyllaeus fimbriiceps</i>	<i>Carypophyllaeus laticeps</i>	<i>Corallotaenia</i> sp.	<i>Proteocephalus ambloplitis</i>	<i>Proteocephalus</i> sp.	<i>Ergasilus</i> sp.	<i>Digenea</i> sp.	<i>Dactylogyrus</i> <i>extensus</i>	<i>Camallanus oxyccephalus</i>	<i>Hysterothylacium</i> sp.		
Striped bass, <i>Morone saxatilis</i>	25	1	1	1	1	5	4	8	1	1	7	4		
		1	167	243	1	39	22	20	11	2	46	7		
		0.04	0.08	0.01	0.01	0.06	0.04	0.09	0.01	0.01	0.08	0.04		
		1	24	243	1	8	6	3	11	2	7	2		
		0.04	1.9	2.7	0.0	0.4	0.2	0.2	0.1	0.0	0.5	0.1		
TOTAL	89	1	167	243	1	39	22	20	11	2	46	7		
		0.01	0.08	0.01	0.01	0.06	0.04	0.09	0.01	0.01	0.08	0.04		
		1	24	243	1	8	6	3	11	2	7	2		
		0.01	1.9	2.7	0.0	0.4	0.2	0.2	0.1	0.0	0.5	0.1		

sample sizes. We obtained samples of 10 fish species as noted in Table 1.

We caught and dissected fishes on-site to ensure the collection of live parasites. We used standard techniques as described by Pritchard and Kruse (1982). We employed stereomicroscopes and examined all internal organs, gills, and body surfaces for metazoan parasites. We temporarily preserved cestodes, monogeneans, and digeneans found in the gastrointestinal tract and liver by placing individuals on a slide glass with a drop of alcohol-formalin-acetic acid (AFA), covering them with slide covers, and wrapping nylon twine around the whole slide. We then stored the slides in jars of 70% ethyl alcohol for transportation to the lab for staining and identification. Nematodes and copepods were preserved in individual test tubes filled with 70% ethyl alcohol.

We stained cestodes, monogeneans, and digeneans in the lab using Mayer's carmalum staining mixture (Pritchard and Kruse 1982) and then created permanent slides of each species using Canada balsam. We left nematodes and copepods whole and unstained in vials of 70% alcohol. We identified parasites to the genus level and to the species level when possible using taxonomic keys and descriptions in Hunter (1927), Schmidt (1970) Gardener and Schmidt (1986), Oros et al. (2010), Hoffman (1999), Fagerholm (1982), La Rue (1914), and Essex (1928).

We then calculated prevalence, mean intensity, and mean abundance for each parasite species (Bush et al. 1997). Prevalence was calculated as the number of fish of a given species infected by a parasite of a given species divided by the total number of fish examined of that species. Mean intensity was calculated as the number of total parasites found of a given species divided by the number of infected fish of a given species. Mean abundance was calculated as the number of total parasites of a given species divided by the total number of fish examined of a given species. Parasite richness is defined as the number of different species found in a single host or single fish species. We did not include fish species for which fewer than 10 individuals were examined (i.e., *Lepomis cyanellus* and *Dorosoma petenense*). Although our focus was on internal parasites, we did find one leach (Family Hirudinidae) attached to the pelvic fin of one channel catfish (*Ictalurus punctatus*). Voucher

specimens of all parasites were deposited in the H.W. Manter Laboratory of Parasitology at the University of Nebraska State Museum, University of Nebraska, Lincoln, Nebraska, USA (Appendix 1).

## RESULTS

### Fish

In 2013, we examined 87 fishes of 7 species, all nonnative (the term *nonnative* refers to species that are not natively found in the Colorado River system). Black crappie (*Pomoxis nigromaculatus*) and striped bass (*Morone saxatilis*) were the most numerous; each made up 28.7% of the total number of fishes sampled. Smallmouth bass (*Micropterus dolomieu*) exhibited the highest parasite richness, with 6 species of metazoan parasites found among sampled fish. Common carp (*Cyprinus carpio*) exhibited the highest average intensity of infection, with one individual infected with 243 parasites of one species (Table 1).

In 2015, we examined 149 fishes of 8 species, all nonnative. Gizzard shad (*Dorosoma cepedianum*) made up 26.2% of the total sample, with smallmouth bass, striped bass, and walleye (*Sander vitreus*) each making up another 18.1% of the total sample. Channel catfish exhibited the highest parasite richness, with 4 species of parasite found among the sampled fish. Common carp exhibited the highest average intensity of infection, with one individual infected with 57 parasites of one species (Table 2).

Combining both sampling years, we collected and examined 236 individual fish of 8 nonnative species which encompassed all of the numerically dominant fish species in the reservoir. In general, benthic- and nearshore-feeding fishes exhibited higher parasite richness and intensity compared to pelagic-feeding fishes. Striped bass made up 22% of the total sampled fish. Channel catfish exhibited the highest prevalence of parasites. Common carp exhibited the highest average intensity of infection. Striped bass exhibited the lowest prevalence of parasites. Walleye and striped bass exhibited the lowest average intensity of infection. Parasite richness was highest in smallmouth bass, which exhibited 6 different species of metazoan parasites, followed by largemouth bass (*Micropterus salmoides*), black crappie, channel catfish, and common carp,

each infected by 4 parasite species. The lowest parasite richness was found in striped bass and walleye, each infected by 1 species, followed by gizzard shad infected by 2 species (Table 3).

### Parasites

In 2013, we found 559 individual parasites comprising 11 species. These included cestodes, copepods, trematodes, monogeneans, and nematodes. Six species of cestode made up 84.6% of the total number sampled in 2013 (Table 1). *Ergasilus* sp. exhibited the highest prevalence and *Caryophyllaeus laticeps* exhibited the highest mean abundance (Table 1).

In 2015, we found 273 individual parasites comprising 4 species of cestode and 1 nematode. Cestodes made up 89.4% of the total parasites examined. *Proteocephalus ambloplitis* exhibited the highest prevalence. *Corallobothrium fimbriatum* and *Khawia* sp. exhibited the highest mean abundances (Table 2).

For both sampling years combined, we found a total of 832 individual parasites in adult, larva, and metacercaria stages comprising 13 different species (Table 3). *Proteocephalus ambloplitis* was the most prevalent parasite, infecting 26 individuals among 5 species of fish. *Caryophyllaeus laticeps*, a cestode, had the highest mean intensity; however, all 243 individuals were found within a single common carp. Five species of parasites were found in only one species of fish (Table 3).

## DISCUSSION

We found substantially different numbers and types of parasites in the 2 different locations and years. In the 2013 sample, we found 166 *Caryophyllaeus fimbriceps* infecting 6 of the common carp sampled (Table 1). In the 2015 sample we did not find any *C. fimbriceps*; however, we only collected 3 common carp that year compared to the 11 collected in 2013 (Table 2). Similarly, we found 72 *Corallobothrium fimbriatum* infecting 8 of the channel catfish sampled in 2015, but none of the channel catfish collected in 2013 exhibited infection from *C. fimbriatum*. There was a large discrepancy in channel catfish sample sizes between the 2 years: 16 in 2015 and only 3 in 2013. These and other similar results are likely due to sampling variation, especially for relatively rare parasite species. It could also be that there are spatial or temporal variations among

TABLE 2. Fish host species and summary statistics for parasite species from fish sampled near Bullfrog Marina in April 2015. EFN = examined fish number, IFN = infected fish number, TPN = total parasite number, P = prevalence, MI = mean intensity, MA = mean abundance.

Parasite category		cestode					nematode	
Fish species	EFN		<i>Corallobothrium fimbriatum</i>	<i>Corallotaenia</i> sp.	<i>Khawia</i> sp.	<i>Proteocephalus ambloplitis</i>	<i>Hysterothylacium</i> sp.	
			<b>BENTHIC FISHES</b>					
Channel catfish, <i>Ictalurus punctatus</i>	16	IFN	8	3	1	7		
		TPN	72	46	23	13		
		P	0.5	0.19	0.06	0.44		
		MI	9	15	23	2		
		MA	4.5	2.9	14	0.8		
Common carp, <i>Cyprinus carpio</i>	3	IFN			1			
		TPN			57			
		P			0.33			
		MI			57			
		MA			19			
Largemouth bass, <i>Micropterus salmoides</i>	5	IFN				2	1	
		TPN				10	4	
		P				0.4	0.2	
		MI				5	4	
		MA				2	0.8	
Smallmouth bass, <i>Micropterus dolomieu</i>	27	IFN				11	11	
		TPN				18	25	
		P				0.41	0.41	
		MI				1.6	2.3	
		MA				0.7	0.9	
Walleye, <i>Sander vitreus</i>	27	IFN						
		TPN						
		P						
		MI						
		MA						
<b>PELAGIC FISHES</b>								
Black crappie, <i>Pomoxis nigromaculatus</i>	5	IFN						
		TPN						
		P						
		MI						
		MA						
Gizzard shad, <i>Dorosoma cepedianum</i>	39	IFN		1		1		
		TPN		2		3		
		P		0.03		0.03		
		MI		2		3		
		MA		0.05		0.08		
Striped bass, <i>Morone saxatilis</i>	27	IFN						
		TPN						
		P						
		MI						
		MA						
TOTAL	151	IFN	8	4	2	21	12	
		TPN	72	48	80	44	29	
		P	0.05	0.03	0.01	0.14	0.08	
		MI	9.0	12.0	40.0	2	2.4	
		MA	0.5	0.3	0.5	0.3	0.2	







parasite communities throughout the lake that account for the differences between the 2 sampling years. Two out of the 5 walleye sampled in 2013 were infected with *Hysterothylacium* sp., but there were no infections of any species of metazoan parasite in the 27 walleye sampled in 2015. It is possible that parasite numbers of *Hysterothylacium* were greater in 2013 compared to 2015, or it could be that *Hysterothylacium* does not occur uplake near Bullfrog. This phenomenon was also seen with *Camallanus oxycephalus*, which was present in one year's sample and completely absent in the other. Temporal variation and seasonal patterns of infection have been observed among other parasites in the Colorado River (Choudhury et al. 2004), which could explain these differences, although further sampling is needed to verify this hypothesis. Based on the differences we saw between the 2 sampling years, there is a high likelihood that other undetected parasites exist in the system at low levels. This study serves as a baseline for future comparison.

In contrast to the spatial and temporal variation observed, one consistent pattern between years was the overall lack of parasite infections in striped bass. Compared to our other species samples, we had a relatively large sample size of striped bass—25 in 2013 and 27 in 2015. Out of the 52 total fish, we found only one infected striped bass, which exhibited only one *C. fimbriceps* individual (Table 3). Though it is possible that we missed some of the rarer parasite species or sampled only healthy fish due to sampling variation, our data suggest that striped bass are generally parasite-free.

Parasite surveys conducted downstream of Lake Powell provide comparison to the patterns we observed. Surveys below the Glen Canyon Dam report *Dactylogyrus extensus* infecting common carp, *Corallobothrium fimbriatum* infecting channel catfish and striped bass, and a species of *Ergasilus* infecting channel catfish (Choudhury et al. 2004, Linder et al. 2012). Though we also found *D. extensus* infecting common carp, we did not find *Ergasilus* sp. infecting any channel catfish, nor did we find *C. fimbriatum* infecting striped bass. Other major differences between our survey and the surveys completed on the Lower Colorado River include the species of fish that were sampled. We only surveyed nonnative species, of which Choudhury et al. (2004) only sampled 2 (common carp and channel catfish) and

Linder et al. (2012) only sampled 3 (channel catfish, common carp, and striped bass). Linder et al. (2012) found a species of *Caryophyllidea* infecting a native cyprinid, flannelmouth sucker (*Catostomus latipinnis*), whereas we found *Caryophyllaeus laticeps* infecting a non-native cyprinid, common carp.

One of the major differences in the findings of our survey and previous surveys downstream was the absence of Asian tapeworm (*Bothriocephalus acheilognathi*) in Lake Powell. There is ample evidence of Asian tapeworm infecting native and nonnative cyprinids in the Lower Colorado River (Choudhury et al. 2004, Stone et al. 2007, Linder et al. 2012) and throughout waters across the American Southwest (Archdeacon et al. 2010). Asian tapeworm was introduced to North America through imported grass carp (*Ctenopharyngodon idella*) and is most commonly found infecting cyprinids (Diaz-Castaneda et al. 1995). In April 2013 and 2015, there was no evidence of Asian tapeworm in our samples. We may not have found any evidence of Asian tapeworm because of our relatively small sample size of cyprinids (14 common carp). However, Asian tapeworm has been recorded infecting a variety of fishes other than cyprinids in the Lower Colorado River (i.e., Ictaluridae, Fundulidae, Salmonidae; Choudhury et al. 2004), which suggests that if Asian tapeworm is present in Lake Powell, it is likely at low levels.

Although parasites have been shown to cause changes in fish behavior and to affect fishes negatively overall (Barber et al. 2000, Barber 2007, Horton and Okamura 2001), we found no cases where parasites appeared to affect fish health negatively. The majority of sampled fishes were infected with few parasites. Even in cases of severely infected fishes, like the common carp sampled in 2013 that was infected with 243 *C. laticeps* (Table 1), the relative volumes of parasites compared to the volumes of the relatively large-bodied fishes they infected were not substantial. As described in other parasite surveys, the collection methods we employed likely selected for fitter individuals that are less parasitized. Individuals that are highly parasitized are less likely to be actively moving and feeding, or are doing so at a decreased rate. If parasites are more prevalent among younger age classes, sampling methods similar to ours that target older and larger fishes are more likely to miss

individuals that are more highly infected (Linder et al. 2012).

Parasites observed in the fishes of Lake Powell exhibit similar life histories with slight variations in host specialization. The majority of observed parasites are cestodes that are transmitted via invertebrates to their definitive hosts. *Caryophyllaeus laticeps* and *Caryophyllaeus fimbriceps* are multihost, trophically transmitted cestodes that infect tubificid worms as larvae. When a benthic-foraging fish feeds on the worm, the fish becomes the definitive host for the parasite and the place where the parasite will reach maturity and reproduce, expelling eggs into the digestive tract and out into the water column (Kulakovskaya 1962, Bauer et al. 1969, Anderson 1976, Lumsden et al. 1982). These 2 parasites are found throughout Europe and North America (Hunter 1927, Chubb et al. 1982), and generally infect cyprinids. *Khawia* spp., *Bothriocephalus claviceps*, *Corallotaenia* sp., *Corallobothrium fimbriatum*, and *Proteocephalus ambloplitis* are cestodes with multiple hosts that have a strategy similar to other cestodes of releasing eggs into the water column. However, these species specialize in exploiting copepods as intermediate hosts, as opposed to annelids (Bangham 1925, 1928, Essex 1927, Hunter 1928, 1930, Hunter and Hunter 1929, Larsh 1941, Befus and Freeman 1973, Stromberg and Crites 1974, 1975, Brandt et al. 1981, Scholz 1997). All of these cestodes have been reported in multiple locations throughout North America, and they appear to be generalist parasites in habitat and in host fish species (Hare 1943, Bangham and Vernard 1946, Morrison 1957, Wilson 1957, Harms 1960, Anthony 1963, Becker 1967, Spall 1968, Becker and Houghton 1969, Woods 1971, Rubertone and Hall 1975, Baker and Crites 1976, Amin 1978, 1991, Sutherland and Holloway 1979, McReynolds and Webster 1980, Williams and Sutherland 1981, Hoffnagle et al. 1990, McDonald and Margolis 1995, Amin and Minckley 1996, Hoffman 1999, Scholz et al. 2001, Szmygiel and Reyda 2010, McAllister and Bursey 2011, Rosas-Valdez and Perez-Ponce de Leon 2011, Scholz et al. 2011). *Digenea* sp. (metacercaria) are multihost trematodes that are transmitted to their definitive host through mollusc intermediate hosts (Ginetsinskaya 1988). *Digenea* have previously been described in *Micropterus salmoides* in the United States (Becker and

Houghton 1969). *Camallanus oxycephalus*, a nematode, is a multihost parasite that uses copepods as an intermediate host. It is a generalist parasite and has been described throughout North America (Bangham 1941, Becker and Houghton 1969, Aho et al. 1991, McDonald and Margolis 1995, Hoffman 1999, cf. Zhang 2012). The life history of *Hysterothylacium* sp., another nematode, is similar to that of other species already noted, but the species is more of a generalist in terms of its intermediate host choice, using annelids, molluscs, echinoderms, crustaceans, and other fish species (Fagerholm 1982, Anderson 2000, Rohde 2005). With its broader intermediate host capabilities, *Hysterothylacium* sp. is also capable of infecting a broad range of top predatory fishes as definitive hosts and it occurs worldwide (Hare 1943, Meyers 1978, Fagerholm 1982). *Dactylogyrus extensus*, a monogenean, has a direct life cycle that does not require an intermediate host. It relies solely on a definitive fish host to complete its life cycle. Eggs are released directly into the water column where they hatch. Larvae then wait for a fish to inhale them where they attach themselves to the gill filaments of the host fish. After attachment on a suitable host, larvae mature and begin reproducing (Izyumova and Zelentsov 1969, Dawes 1977). *Dactylogyrus extensus* has been described only in common carp in North America (Fantham and Porter 1948, Haderlie 1953, Mizelle and Klucka 1953, Bangham 1955, Roberts 1957, Hoffman 1967). *Ergasilus* sp. is part of a small group of ectoparasitic copepods that also have direct life cycles. *Ergasilus* sp., often referred to as gill louse, is capable of prolonged periods of free swimming before attaching itself to the body or gills of the host fish (Freyer 1969, Abdelhalim et al. 1991). These parasites are non-host-specific, but have been described among centrarchids in North America and Great Britain (Roberts 1965, Freyer 1969).

Benthic-feeding fishes showed higher prevalence and intensity of parasite infections than pelagic-feeding species. On average, each species of benthic-feeding fish was infected by 3.8 species of parasites, whereas each species of pelagic-feeding fish was infected by only 2.3 species of parasites. Mean intensity of parasite infections was 21.1 for benthic-feeding fishes and 2.2 for pelagic-feeding fishes. These differences are consistent between years and

locations and are unlikely to be due to differences in sample sizes. Using our results as a baseline, future parasite surveys can compare parasite infections among fishes to gain insights into changes occurring among fish communities over time and throughout the lake.

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APPENDIX 1. Accession numbers for metazoan parasites collected from the fishes of Lake Powell, Utah, USA, and deposited at the University of Nebraska State Museum, Parasitology Collection.

Taxon	Accession number
<i>Bothriocephalus claviceps</i>	HWML 103056
<i>Camallanus oxycephalus</i>	HWML 99863
<i>Caryophyllaeus fimbriceps</i>	HWML 103054
<i>Caryophyllaeus laticeps</i>	HWML 103053
<i>Corallobothrium fimbriatum</i>	HWML 103058
<i>Corallotaenia</i>	HWML 103057
<i>Dactylogyrus extensus</i>	HWML 103061
<i>Digenea</i>	HWML 103060
<i>Ergasilus</i>	HWML 99865
<i>Hysterothylacium</i>	HWML 99864
<i>Khawia</i>	HWML 103055
<i>Proteocephalus</i>	HWML 103063
<i>Proteocephalus ambloplitis</i>	HWML 103059
<i>Proteocephalus ambloplitis</i>	HWML 103062