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Fish communities respond to hydrology and elevation in restored western Kentucky wetlands

Lucas Zuklic

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Fish communities respond to hydrology and elevation in restored western Kentucky wetlands

A thesis
presented to
the faculty of the Department of Biological Sciences
Murray State University
Murray, Kentucky

In partial fulfillment
of the requirements for the degree:
Master of Science

By
Lucas Zuklic

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1
2
3
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5 research possible. I would first like to thank my advisor, Dr. Michael Flinn, for all his help,
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Chapter 1

Hydrologic conditions are the most important factors in determining fish communities found in restored western Kentucky wetlands

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163 Abstract

164 *The Wetlands Reserve Program (WRP) is designed to restore wetland ecological function and*
165 *wildlife benefits; however, these projects rarely monitor biological responses. The objectives of*
166 *our study were to (1) identify environmental factors that were the most important in determining*
167 *wetland fish community composition, (2) examine the influence of the environment on specific*
168 *groups of fish (large-river fishes and Kentucky Species of Greatest Conservation Need (KY-*
169 *SGCN) wetland fishes), and (3) compare fish community composition and diversity in wetlands*
170 *of different condition (e.g., degraded, restored, and reference). From April 2019 to August 2020,*
171 *monthly electrofishing surveys were performed in degraded (n=2), restored (n=8), and reference*
172 *(n=2) wetlands throughout western Kentucky, USA. Over 12,500 individual fish accounting for*
173 *53 species were collected. Non-metric multidimensional scaling was used to visualize fish*
174 *community composition and vector analysis was used to determine which environmental factors*
175 *most affected fish community composition. Our findings suggest that fish community composition*
176 *was most affected by environmental factors that were related to the influence of the Mississippi*
177 *River. Vector analysis identified that large-river fishes displayed a strong positive association*
178 *with wetlands that were more hydrologically influenced by the Mississippi River, whereas KY-*
179 *SGCN wetland fishes displayed a strong positive association with wetlands less hydrologically*
180 *influenced by the Mississippi River. ANOSIM and Hill diversity (Hill-Shannon; $q=1$) were used*
181 *to quantify differences in fish community composition and diversity among wetland condition.*
182 *Furthermore, our results indicate western Kentucky WRP restorations have not created unique*
183 *community composition but have rapidly fostered levels of fish diversity similar to reference*
184 *wetlands. Current and future WRP restoration managers may need to consider potential*
185 *tradeoffs between wetland wildlife communities and wetland function to best promote restoration*
186 *goals.*

187
188

188 Introduction

189 Wetlands are crucially important ecosystems for both humans and wildlife. Wetlands
190 provide humans with numerous ecosystem services that are valued at approximately 35 trillion
191 USD a year (Costanza et al. 2014). Wetlands are critical for wildlife because they are productive
192 and habitat-rich ecosystems that foster the existence of diverse assemblages of biota (Mitsch &
193 Gosselink 2015). Despite their value to humans and wildlife, wetlands have suffered large-scale
194 global losses (Dahl & Allord 1996; Mitsch & Gosselink 2015). Many of the United States' non-

195 coastal wetland resources exist as floodplain wetlands throughout the Mississippi River Alluvial
196 Valley (MAV). Once supporting nearly 10 million ha of bottomland hardwood forest, the MAV
197 has experienced radical alterations to its regional hydrology and large-scale land use conversion
198 to agriculture, which has led to a dramatic loss of wetlands, wildlife benefits, and ecosystem
199 function (Semlitsch 2000; Frederickson 2005; Rewa 2005; King et al. 2006; Faulkner et al. 2011;
200 Kleunder et al. 2015; USDA-NASS 2017).

201 In response to wetland loss throughout the United States, the Wetlands Reserve Program
202 (WRP), by 2013, had restored over 1 million ha of wetlands, of which approximately 250,000 ha
203 were located throughout the MAV (Natural Resources Conservation Service 2013a). WRP
204 focuses on using a combination of reforestation and hydrologic restoration techniques to foster
205 ecosystem processes that allow wetland restoration (King & Keeland 1999; Hayes & Egan 2004;
206 Rewa 2005). Special emphasis is placed on restoring hydrology. Hydrology has been found to be
207 the critical driver of wetland processes that, when restored, most quickly returns wetland
208 function and wildlife benefits (Bedford 1996; Brinson & Rheinhardt 1998; Rewa 2000; Zedler
209 2000; Haynes & Egan 2004; Rewa 2005; Brauman et al. 2007; Hunter et al. 2008; Faulkner et al.
210 2011; King & Keim 2019). Through wetland restoration, WRP aims to return lost wetland
211 function and wildlife benefits (Natural Resources Conservation Service 2013a).

212 To assess wetland restoration success, WRP restoration practitioners employ a variety of
213 post-restoration monitoring studies (see Osmond et al. 2012). Typically, these studies focus
214 directly on the response of wetland function, water quality, or use vegetation response as a proxy
215 for wetland function (Rewa 2005; King et al. 2006; Faulkner et al. 2011). Post-restoration
216 monitoring efforts, however, are sometimes misleading as not all restorations are monitored and
217 restoration success criteria are not always clearly defined (Zedler 2000; Stanturf et al. 2001;

218 Ruiz-Jaen & Aide 2005; USDA OIG 2008; Moreno-Mateos et al. 2012). Additional problems
219 may arise if only wetland function is considered because wetland function and wildlife usage
220 have not always been found to be maximized at the same wetland (Zedler 2000). Even though
221 one goal of WRP is to return lost wildlife benefits, relatively few studies exist that quantify the
222 response of wildlife communities (Rewa 2005). Considering the goals of the WRP, wildlife
223 community responses should also be measured post-restoration.

224 The utility of evaluating post-restoration success using biological indicator species such
225 as Species of Greatest Conservation Need (SGCN) have been published (Benson et al. 2018).
226 Biological indicator species provide information on the functionality of a particular ecosystem,
227 as species within an assemblage vary in their environmental requirements and their sensitivity to
228 disturbance. To our knowledge, a paucity of studies exist that utilize fish community response as
229 a measurement of WRP post-restoration success (but see Leao 2005; Rewa 2005; Benson et al.
230 2018). The use of fish community response to WRP restorations throughout the MAV is well-
231 founded because hydrology is a major determinant of wetland function and fish community
232 structure. MAV wetland function and fishes depend on a natural flood-pulse cycle where water,
233 sediments, and nutrients are supplied to wetlands through seasonal connections to the floodplain
234 (Junk et al. 1989; Faulkner & Patrick 1992). For both large-river and SGCN wetland fishes,
235 seasonal connections to floodplain wetlands determines the diversity, community composition,
236 and overall success of these fishes (Welcomme 1985; Junk et al. 1989; Caldwell et al. 2011;
237 Zeug et al. 2005) by providing abundant energy resources (e.g., zooplankton) and granting
238 suitable spawning habitat (Welcomme 1985; Winemiller and Rose 1992, King & Keeland 1999;
239 Baber et al. 2002; Kluender et al. 2015). Hydrology and water quality in MAV floodplain
240 wetlands, however, can differ greatly among seasons as wetlands are more influenced by the

241 river channel in the winter and spring and more influenced by local climatic events during the
242 summer and fall (Junk et al. 1989; Mitsch & Gosselink 2015). Individual species of fish display
243 different responses to specific wetland environmental conditions (Jester et al. 1992; Miranda &
244 Lucas 2005; Dembkowski & Miranda 2012). For example, large-river fishes depend on seasonal
245 access to floodplain wetlands during flooding events while SGCN wetland fishes utilize the
246 floodplain's shallow depths, sluggish flow, dense vegetation, and soft substrate year-round
247 (Welcomme 1985; Junk et al. 1989; Petts 1989; Aarts et al. 2004; Hohausova et al. 2010;
248 Beesley et al. 2014; Kluender et al. 2015; Eisenhour et al. 2018; Simpson et al. 2021). Differing
249 responses of fishes, therefore, underline the importance to quantify the response of different
250 groups of taxa (Benson et al. 2018).

251 Due to the suitability of utilizing fishes as biological indicators and to address knowledge
252 gaps related to fish usage of WRP restored floodplain wetlands in the MAV, our study examined
253 fish response to environmental conditions, i.e., hydrology, water quality, and biota in restored
254 wetlands. The objectives of this research were to (1) determine the relationships between fish
255 community composition and environmental conditions in western Kentucky wetlands, to (2)
256 determine the relationships between specific groups of fish (large-river fishes and Kentucky
257 SGCN (KY-SGCN) wetland fishes) and environmental conditions, and to (3) examine if
258 differences in wetland condition (e.g., degraded, restored, and reference) led to differences in
259 fish community structure. It was thought that wetland hydrology would be most important in
260 determining fish community composition, that the relative abundance of large-river fishes and
261 KY-SGCN wetland fishes would be determined by differences in hydrology, that fish
262 community composition in WRP restored wetlands would differ from those in reference and
263 degraded wetlands, and that WRP fish diversity would be intermediate between that of degraded

264 and reference wetlands. Hopefully, the fish/environment relationships found in our wetlands can
265 be used to inform future wetland restoration management decisions and ensure the most effective
266 management decisions (Merkle et al. 2019).

267 **Methods**

268 Study Area

269 Our study was conducted in the Mississippi Alluvial Plain and Mississippi Valley Loess
270 Plain ecoregions of western Kentucky (Omerink 1987). Historically, wetland resources in this
271 region were characterized by bottomland hardwood forests and stream floodplains that
272 experienced dramatic but predictable hydroperiods seasonally influenced by the upper
273 Mississippi River watershed (King et al. 2006; Mitsch & Gosselink 2015). A substantial amount
274 of wetland resources in this region still exist and are located along three major tributaries
275 (Mayfield Creek, Obion Creek, and Bayou du Chien Creek) that drain approximately 2,800 km²
276 into the Mississippi River. These wetlands are highly fragmented and suffer from degradation
277 from intensive agriculture and regional hydrologic modifications (Frederickson 2005, King et al.
278 2006). Current land use of this region of western Kentucky is dominated by cultivated crops
279 (64%) and forested floodplain wetlands (25%) (Dewitz 2019). Despite these anthropogenic
280 influences, the region retains some features of a large-river floodplain; for example, seasonally
281 high discharges reconnect the floodplain in the winter and spring (Mitsch & Gosselink 2015).

282 Wetland Selection

283 Eight wetlands restored by WRP in western Kentucky (Figure 1) were sampled. Restored
284 wetlands ranged in size from one to 20 ha and in age since hydrologic restoration from one to 13
285 years. WRP employed a variety of engineering techniques to restore local hydrology on the
286 selected wetlands (Personal communication, NRCS). Additionally, some restored wetlands were

287 planted with tree saplings to initiate reforestation and currently managed following Compatible
288 User Agreements, which include food plot planting, mowing, and water level management
289 (Personal communication, NRCS). Before wetland selection, pertinent WRP information from
290 the National Resources Conservation Service (NRCS) (including landowner contact, restoration
291 type, restoration age, restoration plans) was obtained. After obtaining restored wetland
292 information, the following criteria were used to select restored wetlands: (1) location on one of
293 the regional tributaries to the Mississippi River (Figure 1) and (2) similarity of hydrogeomorphic
294 wetland class (riverine following Brinson et al. 1993), hydroperiod (semi-permanent to
295 permanent following Cowardin et al. 1995) (Figure 2), and size (1-20 ha).

296 Poor condition (degraded wetlands; n=2) and good condition (reference wetlands; n=2)
297 wetlands were also sampled to compare with WRP restored wetlands (Figure 1). Degraded
298 wetlands were once natural wetlands that have experienced dramatic local hydrologic alterations
299 for agricultural purposes, but still exhibit some wetland characteristics. Both of our degraded
300 wetlands were positioned in active agricultural fields. Reference wetlands have not been
301 subjected to local hydrologic alterations, but ultimately still exist within a highly altered
302 landscape. One of our reference sites was a forested wetland located on a KY Wildlife
303 Management Area; the other site was a bottomland hardwood swamp positioned on an upstream
304 portion of one of our WRP easements.

305 Fish sampling

306
307 From April 2019 to August 2020, monthly electrofishing surveys were conducted at each
308 wetland. A 24-volt battery powered Smith-Root LR-24 backpack electrofisher was used to
309 collect fish from accessible wetland shoreline. Each site was sampled for 600-750 seconds at
310 400-500 volts and at 30 Hz on a 25% duty cycle. Collected fishes were measured, identified to

311 species using Etnier & Starnes (1993) and Pflieger (1997), enumerated, and then returned.
312 Specimens that could not be identified in the field were anesthetized with clove oil, preserved in
313 a 10% formalin solution, and later identified. For each sample, the abundance of each fish
314 species was divided by the effort (minutes) to determine catch per unit effort (CPUE)
315 (individuals per minute). CPUE was then averaged across all monthly sampling events to yield a
316 single value representing the CPUE of a specific fish species for each wetland.

317 Environmental influences on fishes

318 Twenty-eight environmental predictor variables (see below for explanations; Table 1)
319 were quantified to examine the influence on fish community composition and the relative
320 abundance of large-river and KY-SGCN wetland fishes. Large-river fishes were classified using
321 species descriptions from Etnier and Starnes (1993) and consisted of *Cycleptus elongatus* (blue
322 sucker), *Ictalurus punctatus* (channel catfish), *Aplodinotus grunniens* (freshwater drum),
323 *Dorosoma cepedianum* (gizzard shad), *Ictiobus bubalus* (smallmouth buffalo), *Sander*
324 *canadensis* (sauger). KY-SGCN wetland fishes were classified using Kentucky's Comprehensive
325 Wildlife Conservation Strategy report (2013) and consisted of *Umbra limi* (central mudminnow),
326 *Hybognathus hayi* (cypress minnow), *Lepomis marginatus* (dollar sunfish), *Fundulus chrysotus*
327 (golden topminnow), *Erimyzon sucetta* (lake chubsucker), *Lepomis miniatus* (redspotted
328 sunfish), and *Notropis maculatus* (taillight shiner).

329 Hydrology

330
331 Depth was recorded in each wetland from March 2019 to August 2020 using water level
332 loggers (HOBO® U20-001-04, Onset Computer Corporation). One logger was deployed in the
333 deepest wadeable location of each wetland. Depth was recorded every 15 minutes. Water level
334 data was averaged per day and daily depths were then used to calculate mean depth (m),

335 hydroperiod (days), and 13 indicators of hydrologic alteration (IHA) following Richter et al.
336 (1996) (Table 1). The percent of time each wetland experienced connectivity to its associated
337 stream was estimated by determining the stream depth at which each wetland was connected to
338 the stream and then by calculating the percent of time the stream was greater than that depth. For
339 wetlands on upstream portions of tributaries, stream depth was determined by using the United
340 States Geological Survey stream gauge 07024000 (USGS 2019-2020) on the Bayou du Chien
341 and for wetlands located on the Mississippi River floodplain stream depth was determined by
342 using the United States Geological Survey stream gauge 07022000 (USGS 2019-2020) on the
343 Mississippi River.

344 Hydrologic connectivity

345
346 Four landscape variables were calculated to indirectly quantify hydrologic connectivity
347 between wetlands and other permanent bodies of water. Distance to Main Channel was
348 quantified by measuring the straight-line distance from the center of each wetland to the main
349 channel of the nearest major stream in ArcGIS Pro (Version 2.7, Esri Inc.). Topography was
350 calculated by using 10-m U.S. Geological Survey (2017) 1/3 arc-second Digital Elevation
351 Models (DEMs) to find the mean slope inside a 1-km buffer around each wetland. The “elevation
352 profile” tool in USGS Stream Stats was used to delineate each wetland’s boundary and assign
353 elevation (m) values for 50 different locations within each wetland. Elevation was calculated by
354 taking the median of each wetland’s 50 elevation values. Waterway Distance to the Mississippi
355 River was calculated by using the “flow-path” tool in USGS Stream Stats to find the shortest
356 waterway distance (km) from each wetland to the main channel of the Mississippi River.
357 Waterway distance was considered as a proxy for the hydrologic influence the Mississippi River
358 exerted on each wetland; greater values imply lower hydrologic influence.

359 Water quality

360

361 Changes in water temperature (°C) and dissolved oxygen (DO) (mg/L) were recorded in

362 each wetland from March 2019 to August 2020 using multi-parameter sondes (YSI® EXO2,

363 Xylem Incorporated). One sonde was suspended in each wetland approximately midway in the

364 water column in the deepest wadeable location. Sondes recorded data at 15-minute intervals.

365 Measurements were averaged per day and then used to calculate minimum temperature,

366 maximum temperature, mean temperature, mean DO, and minimum DO for each wetland.

367 Biotic variables

368

369 Zooplankton communities were sampled monthly from April 2019 through August 2020

370 at each wetland using a 9-cm diameter littoral sampling tube following Pennak (1962). Each

371 sample consisted of three replicates averaged together, one taken from open water, wetland edge,

372 and dense vegetation. Samples were poured into a volumetric container, volume (L) was

373 recorded, rinsed through a 43-µm sieve, and preserved in 4% buffered formalin solution. Later,

374 samples were subsampled to a maximum of 1/8 using a Folsom Plankton Splitter (WILDSCO,

375 Saginaw, MI) so that at least 50 zooplankters were found per subsample. Cladocerans and

376 copepods were enumerated but rotifers were excluded. After scaling back up to 100% from the

377 subsampled fraction, density was calculated by dividing the abundance of each sample by the

378 original volume of water.

379 Aquatic vegetation was sampled monthly at each site between April 2020 and August

380 2020, which corresponded with the region's growing season, using a 1m-by-1m quadrat. Each

381 sample consisted of nine replicates averaged together: three each taken from open water, wetland

382 edge, and dense vegetation. Percent cover of three aquatic vegetation groups (aquatic emergent,

383 aquatic submergent, and aquatic floating) was estimated using six cover categories (1 = 0-10%,

384 2= 11-20%, 3 = 21-40%, 4 = 41-60%, 5= 61-80%, 6 = 81-100%). The midpoint of the cover
385 categories was used for percent cover calculations.

386 The Kentucky Wetland Rapid Assessment Method (KY-WRAM) (Kentucky Division of
387 Water, 2016) was used to assess general wetland integrity and function. The KY-WRAM is used
388 to evaluate overall wetland function and ecological integrity and was developed for use by
389 regulatory agencies in wetland permitting decisions. The KY-WRAM assumes that wetlands
390 with high ecological integrity also have high wetland function. The KY-WRAM metrics capture
391 a full range of potential disturbances to wetland integrity and fall into six categories: size and
392 distribution; buffer and intensity of surrounding land uses; hydrology; habitat alteration and
393 habitat structure development; special wetlands; and vegetation, interspersions, and habitat
394 features. Each category is subdivided into additional metrics. The special wetlands category was
395 omitted because we were unable to gain access to the required information (all wetlands received
396 a score of 0 for this category). The final KY-WRAM score is the sum of all the metric scores and
397 ranges from 0 (very poor condition) to 90 (reference condition). Out of the metrics, 10 were
398 evaluated using ArcGIS Pro® software (Version 2.7, Esri Inc.) and high resolution orthoimages
399 (USDA 2018), and 10 were assessed in the field during a one-hour site visit per wetland during
400 July - September 2020.

401 Statistical Analyses

402
403 Statistical analyses were performed using R statistical software (version 4.0.5) (R core
404 team 2021) and the ‘Vegan’ software package (Oskansen et al. 2013). Nonmetric
405 multidimensional scaling (NMDS) was used to visualize variation in fish community
406 composition among wetland condition. Experimental units in ordinations were defined as the
407 average CPUE of each fish species present at each wetland in multivariate space. No fish species

408 were removed prior to ordination because we were interested in rare and uncommon taxa.
409 Ordinations were constructed with Bray–Curtis distances, run for the maximum number of
410 iterations (300), and chosen for minimum stress values. Dimensional solutions, stepping down
411 from six to one, were tested and determined by the use and examination of individual scree plots
412 (McCune & Grace 2002).

413 Analysis of similarity (ANOSIM) was used to test for differences in fish community
414 composition among *a priori* groups of wetland condition. The Gower distance was used in
415 ANOSIM and run for 9999 maximum iterations generating *r*-values between -1 and 1. Positive
416 values indicate differences among groups and significance was tested at $\alpha = 0.05$.

417 The influence of environmental predictor variables (Table 1) on fish community
418 composition, large-river fishes, and KY-SGCN wetland fishes was examined using vector fitting
419 analysis with the function ‘ENVFIT’ on NMDS ordinations. Each variable’s association with
420 each experimental unit was indicated by the vector’s direction while the strength of each
421 variable’s association was indicated by the vector’s length (McCune & Grace 2002). Vector
422 significance ($\alpha = 0.05$) was estimated using 999 random permutations of the data (Faith & Norris
423 1989). For ease of interpretation, the hydrologic variables 1-Day Maximum (m), 7-Day
424 Maximum (m), 30-Day Maximum (m), and 90-Day Maximum were grouped into one vector
425 labeled High Water Magnitude on ordination plots because of high correlation (overlapping
426 angle and vector magnitude) among variables (Flinn et al. 2008).

427 Vectors representing the relative abundance of large-river and KY-SGCN fishes were
428 also placed onto NMDS ordinations. Vectors representing these groups of fishes were intended
429 to help explain their association with the environment and were not used to predict fish
430 community composition.

431 Diversity

432

433 Hill diversity was calculated for each wetland based on condition. Hill diversity was
434 chosen due to its numerous advantages over other diversity indices (see Chao et al. 2014) and
435 was calculated according to Hsieh et al. (2016), and Chiu & Chao (2014). See Jost (2006) for a
436 more thorough review of the benefits of Hill diversity in relation to other indices or Roswell et
437 al. (2021) for a current consensus within the ecological community.

438 Before calculating Hill diversity, samples were standardized by 95% coverage to account
439 for uneven sampling effort (Chao & Jost 2012; Chao et al. 2014; Roswell et al. 2021; R package
440 iNEXT). Coverage is a relatively new method of sample standardization in ecology that
441 measures sample completeness and accounts for the abundance of species in the sampled
442 community. Coverage estimates the proportion of individuals in the community that belong to
443 species present in a sample (Roswell et al. 2021). For example, achieving coverage of 95%
444 means that 5% of individuals in the community were not sampled.

445 Hill diversity varies based on the choice of exponent used. Ecologists most commonly
446 use $q = 0$ (species richness), $q = 1$ (Hill-Shannon diversity), and $q = 2$ (Hill-Simpson diversity).
447 Hill-Shannon diversity ($q = 1$) was used because it results in all individuals being considered
448 equally as it counts species proportionately to their abundance or incidence (Chao et al. 2014).
449 Hill-Shannon diversity was calculated for each wetland using the iNEXT package which
450 provides functions to compute the most widely used Hill numbers ($q = 1$, $q = 2$, $q = 3$) for
451 individual-based abundance data or sampling-unit based incidence data. Incidence data was used
452 because it suitably represents timed surveys, e.g., timed electrofishing surveys, and because
453 Colwell et al. (2012), Chao et al. (2014), and Chao & Colwell (2017) demonstrated that
454 incidence data allows for biological inference just as powerful as abundance-based approaches.

455 Our input data for the iNEXT package consisted of species-specific incidence data from each
456 sample from each wetland which was categorized by each sample's wetland condition.

457 To describe patterns in Hill-Shannon diversity, sample and coverage-based rarefaction
458 and extrapolation curves were generated using the "estimate d" function (R package, iNEXT) to
459 determine how diversity increases with increasing sampling effort and completeness. Rarefaction
460 and extrapolation of Hill-Shannon diversity were conducted according to Hsieh et al. (2016) and
461 further discussed by Colwell et al. (2012), Chao & Jost (2012), and Chao et al. (2014). Sample-
462 based curves evaluated the number of individuals in a sample by plotting diversity estimates in
463 relation to the number of sampling units. Coverage-based curves were plotted against rarefied
464 sample completeness to illustrate diversity estimates in relation to sample coverage. All
465 extrapolation curves were plotted using a doubling in sample size and 500 bootstrap replicates
466 were used to estimate 95% confidence intervals. Confidence intervals, a known alternative to
467 standard statistical testing (Magurran 1988; Colwell, Mao, & Chang 2004), were used to
468 determine if differences between wetland condition were statistically significant.
469 Nonoverlapping 95% confidence intervals, associated with rarefied or extrapolated curves,
470 indicate possible significant differences at $\alpha = 0.05$ (Chao & Jost 2012; Chao et al. 2014).

471 **Results**

472

473 Fish Sampling

474

475 All wetlands had fish present. 12,518 fish from 17 families, 37 genera, and 53 species
476 were collected across all wetlands. The mean CPUE (individuals per minute) in restored
477 wetlands was 10.95 ± 1.91 (SE), 7.41 ± 1.94 (SE) in degraded wetlands, and 4.33 ± 0.42 (SE) in
478 reference wetlands. Degraded wetlands had 22 of the 53 recorded species, 47 species were found
479 in restored wetlands, and 38 species were found in reference wetlands. Golden topminnow was

480 only found in degraded wetlands, blacktail shiner (*Cyprinella venusta*), blue sucker, channel
481 catfish, common carp (*Cyprinus carpio*), freshwater drum, gizzard shad, grass carp
482 (*Ctenopharyngodon idella*), logperch (*Percina caprodes*), silver carp (*Hypophthalmichthys*
483 *molitrix*), sauger, and yellow bass (*Morone mississippiensis*) were only found in restored
484 wetlands, and bullhead minnow (*Pimephales vigilax*), pugnose minnow (*Opsopoeodus emiliae*),
485 red shiner (*Cyprinella lutrensis*), and taillight shiner were only found in reference wetlands.
486 Overall community composition of all wetlands was dominated by Centrarchidae (sunfishes)
487 (54% of all individuals collected) and Poeciliidae (livebearers) (18%). Across all wetlands, the
488 relative abundance of large-river fishes was 1.7% of all individuals collected whereas the relative
489 abundance of KY-SGCN wetland fish was 3.7%.

490 Environmental influences on fish community composition

491 Final NMDS solutions consisted of two dimensions with a low final stress value (0.102)
492 and high interpretability compared to alternative solutions (Figure 3). Eight environmental
493 variables were found to be significant in determining fish community composition and were
494 overlaid as vectors onto NMDS ordination (Figure 3, Table 2). Environmental vectors displayed
495 a strong horizontal gradient: increasing Waterway Distance to the Mississippi River and
496 Zooplankton Density were associated with each other and were negatively associated with
497 increasing High Water Magnitude, Duration of Connectivity, and Low Water Duration. The
498 horizontal gradient, at least in part, helped explain the fish community composition of many of
499 our wetlands based on condition. Environmental vectors did not describe a vertical gradient well
500 and fish community composition of wetland condition, in many cases, had at least some vertical
501 aspect to it. Rise Count was an exception to this pattern as it lied in between vertical and
502 horizontal gradients: increasing Rise Count was more positively associated with increasing

503 Zooplankton Density and Waterway Distance to the Mississippi River and more negatively
504 associated with High Water Magnitude, Duration of Connectivity, and Low Water Duration.
505 Positive and negative associations existed between Rise Count and fish community composition
506 in many of our wetlands based on condition.

507 Environmental influences on river and KY-SGCN wetland fishes

508
509 The environmental variables High Water Magnitude, Duration of Connectivity, and Low
510 Water Duration had strong positive association with the relative abundance of large-river fishes
511 (Figure 3). Rise Count had strong positive association with the relative abundance of KY-SGCN
512 wetland fishes (Figure 3). Zooplankton Density and Waterway Distance to the Mississippi River
513 had at least some positive association with the relative abundance of KY-SGCN wetland fishes
514 (Figure 3). The vectors representing relative abundance of large-river fishes and relative
515 abundance of KY-SGCN wetland fishes had opposing relationships to one another (Figure 3).

516 Influence of wetland condition on fish community composition

517
518 NMDS ordination based on electrofishing CPUE data revealed little separation of fish
519 community composition by wetland condition (Figure 3). Results of ANOSIM analysis
520 confirmed that degraded, restored, and reference wetland fish community composition were not
521 significantly different ($r = -0.182$, $p = 0.732$).

522 Influence of wetland condition on fish diversity

523
524 Hill-Shannon diversity in restored wetlands was 28.6, 95% CI [27.7, 29.9] which was not
525 significantly different from Hill-Shannon diversity in reference wetlands 26.6, 95 % CI [24.9,
526 28.7] (Figures 4, 5). Hill-Shannon diversity in restored and reference wetlands were significantly
527 greater than Hill-Shannon diversity in degraded wetlands 18.1, 95% CI [17.4, 19.6] (Figures 4,
528 5). The estimated curve patterns of Hill-Shannon diversity accumulation per sampling unit for

529 degraded, restored, and reference wetlands were approaching asymptotic (plateau), suggesting
530 that the sampling strategy was sufficient in revealing true patterns of fish diversity associated
531 with these three wetland types. Coverage-based rarefaction and extrapolation further indicated
532 that sample completeness was consistent across wetland type as coverage values were all greater
533 than 95% (Figure 5).

534 *Discussion*

535
536 Our results indicated that hydrologic influence from the Mississippi River was more
537 important in determining fish community composition than wetland condition as differences in
538 hydrologic influence from the Mississippi River determined the relative abundances of large-
539 river and KY-SGCN wetland fishes. Hydrologic influence from the Mississippi River led to
540 differences in hydrology between wetlands with short and long waterway distances to the river.
541 Wetlands with short waterway distances were strongly influenced by the river as they
542 experienced direct lateral connectivity with the Mississippi River during its high magnitude
543 depth long-lasting spring flood pulse but seldom reconnected with the river after the spring.
544 Wetlands with long waterway distances were less influenced by the river as they did not
545 experience direct lateral connectivity with the Mississippi River and, therefore, experienced
546 lower magnitude depth shorter-lasting flooding and were more likely to flood after the spring
547 pulse. Even though restored wetlands did not have unique community composition, restored
548 wetlands had levels of diversity greater than those of degraded wetlands and was comparable to
549 reference wetlands. With these results our study has demonstrated that wetland restoration may
550 promote fish diversity and hydrologic influence is an important factor to consider regarding
551 specific groups of fishes.

552

553

554 *Influence of the Mississippi River*

555 Our study suggests that each wetland's waterway distance to the Mississippi River
556 determined the influence of the river's hydrology: differences in influence created contrasting
557 hydrologic conditions in short and long waterway distance wetlands (see above) (Figure 3).
558 Contrasting differences in hydrology most likely determined the relative abundance of large-
559 river and KY-SGCN wetland fishes and led to the contrasting associations between the two fish
560 groups. The hydrologic influence from the Mississippi River in wetlands with short waterway
561 distances was exacerbated due to the exceptional winter and spring flooding of 2019 and 2020.
562 According to the National Weather Service's (NWS) river gage at Cairo, IL, the Mississippi
563 River stayed above flood stage (40 ft) for 146 consecutive days from February 8th, 2019 to July
564 13th, 2019 and, during this time, it reached its third greatest height ever recorded (56.5 ft).
565 Although not as extreme as 2019, the Mississippi River still experienced exceptional flooding in
566 2020 as the NWS Cairo, IL gage reported 96 consecutive days above flood stage from February
567 8th, 2020 to May 12th, 2020, during which, the river reached its 21st greatest height ever recorded
568 (52.6 ft). Mississippi River flood events, like these, may become more likely as winter and
569 spring precipitation throughout the upper Mississippi River watershed is expected to increase 20-
570 30 % by the year 2100 (Wuebbles & Hayhoe 2004). The association between zooplankton
571 density and waterway distance to the Mississippi River (Figure 3) help support this claim, as
572 previous studies have found that greater riverine influence decrease water residence times and
573 maximize dilution effects leading to lessened zooplankton densities (Pace et al. 1992; Bozelli et
574 al. 2015; Godfrey et al. 2020).

575 Influence of the Mississippi River may have also been responsible for the long periods of
576 low water in wetlands with short waterway distances to the river (Figure 3). Watershed size

577 determines how local water levels respond to precipitation (USGS Watershed Science School
578 2019). Precipitation in a small watershed can drastically increase stream levels and induce
579 overbank flooding. The opposite is true of the Mississippi River as local water levels are
580 determined by precipitation that has occurred upstream and throughout its entire watershed.
581 Once water levels dropped in the Mississippi River after spring flooding, local precipitation was
582 unlikely to increase water levels enough to reconnect short waterway distance wetlands to the
583 river. Conversely, precipitation after spring flooding allowed wetlands further from the
584 Mississippi River to periodically receive stream inputs and maintain water levels throughout the
585 year. Water levels of the Mississippi River may have also affected the floodplain's water table
586 and affected drying. Flood-stage river levels in high order streams have been found to impede
587 floodplain water drainage (Berkowitz et al. 2020). After river levels dropped, the floodplain's
588 water table may have begun to rapidly drain leading to eventual drying in wetlands with short
589 waterway distances.

590 *Environmental influences on large-river and KY-SGCN wetland fishes*

591 Depth is a well-studied environmental factor that is important in structuring fish
592 communities (Rodriguez & Lewis Jr. 1997). Generally, when increased, depth has been found to
593 be beneficial for fishes as it provides habitat heterogeneity, environmental stability, and refugia
594 from poor water quality (Zeug et al. 2005; Shoup & Wahl 2009; Miranda 2011; Dembkowski &
595 Miranda 2012). For example, deeper depths increase habitat suitability for large river fishes (i.e.,
596 access to the floodplain) and serve as a trigger for reproduction (Welcomme 1985; Copp &
597 Penaz 1988; Junk et al. 1989; Winemiller & Rose 1992; Beesley et al. 2014, Kluender et al.
598 2015). Benefits provided to fishes by increasing depth may reach a threshold, however, and
599 depths exceeding that threshold may become detrimental to some fishes. High magnitude depth

600 can act as a measure of environmental stress for wetland fishes by decreasing floodplain habitat
601 suitability, i.e., creating lotic conditions or greatly inundating shallow littoral areas (Resh et al.
602 1988; Richter et al. 1996). When decreased, depth can limit habitat heterogeneity (Dembkowski
603 & Miranda 2012), can impose foraging limitations on fishes (Thomasz et al. 1997), increase
604 chance of predation, lead to poor water quality (Zeug et al. 2005), and increase likelihood of
605 drying, all of which may lead to depauperate fish assemblages or cause die-offs (Zeug et al.
606 2005; Shoup & Wahl 2009; Miranda 2011; Dembkowski & Miranda 2012).

607 Our study observed contrasting associations that existed between different groups of
608 fishes and wetland depth (Figure 3). Large-river fishes benefited from high magnitude depths as
609 it granted floodplain access. KY-SGCN wetland fishes, however, were likely negatively affected
610 by high magnitude depths because they require shallow littoral areas (Simpson et al. 2021) that
611 also coincide with predictable water levels and lentic conditions (Etnier & Starnes 1993;
612 Eisenhour et al. 2018). Relative abundance of KY-SGCN wetland fishes was negatively
613 associated with low water events (Figure 3), however, this association was probably due to the
614 influence of the Mississippi River, i.e., KY-SGCN wetland fish are less likely to utilize wetlands
615 strongly influenced by the Mississippi River. Even though large-river fishes were able to utilize
616 wetlands more influenced by the Mississippi River, prolonged low water events during the
617 summer of 2019 prevented year-round survival (e.g., samples were fishless until the spring of
618 2020) as fishes became trapped and were subjected to poor water quality, predation, and eventual
619 drying (Figure 6). Absence of prolonged low water and drying in wetlands less influenced by the
620 Mississippi River benefited KY-SGCN by providing year-round habitat.

621 Lateral connectivity is important for fishes as it is one of the most influential components
622 of floodplain ecosystem dynamics (Junk et al. 1989) that can affect water quality, primary

623 productivity (Knowlton & Jones 1997; Galat et al. 1998), fish metacommunity dispersal,
624 colonization, habitat utilization, and refugia from other adverse wetland conditions (Thomasz et
625 al. 1997; Baber et al. 2002; Miranda 2005; Zeug et al. 2005; Zeug & Winemiller 2008; Shoup &
626 Wahl 2009; Miyazono et al. 2010; Beesley et al. 2014). Conversely, a lack of lateral connectivity
627 may be detrimental to fishes as it can lead to stranding and mortality (Richter et al. 1996).

628 Lateral connectivity was present in wetlands with short and long waterway distances to the
629 Mississippi River. However, lateral connectivity experienced in our wetlands behaved differently
630 as lateral connectivity in wetlands with short waterway distances to the river were long-lasting
631 and seasonal while lateral connectivity between nearby streams and wetlands with long
632 waterway distances were shorter lasting, occurred more frequently, and occurred throughout the
633 year (i.e., Rise Count, Table 1; Figure 3). Greater relative abundance of large-river fishes in
634 wetlands highly influenced by the Mississippi River suggests that long-lasting lateral
635 connectivity with the river allowed these fishes to disperse from the river and utilize floodplain
636 habitat. Similarly, other studies have found that large-river fishes were the primary colonizers of
637 floodplain habitat next to large rivers that experienced direct lateral connectivity between the
638 river and floodplain (Miranda 2005; Zeug et al. 2005; Zeug & Winemiller 2008; Miyazono et al.
639 2010). Smaller frequent pulses resulting in year-round lateral connectivity most likely benefited
640 KY-SGCN wetland fishes as they are less tolerant of high magnitude flooding and lotic
641 conditions. Additionally, frequent pulses likely mitigated adverse water quality conditions
642 associated with the summer and fall which allowed year-round survival of fishes. KY-SGCN
643 wetland fishes require shallow, vegetated, lentic environments to complete their life cycles
644 (Etnier & Starnes 1993; Eisenhour et al. 2018) and source of lateral connectivity may be
645 meaningful. Habitat requirements for KY-SGCN wetland fishes were absent during periods of

646 lateral connectivity with the Mississippi River, which brought upon lotic conditions, colder water
647 temperatures, and exceptional depths. Additionally, a lack of vegetation was observed when
648 sampling after flooding, which may have been due to plants being inundated too long or from
649 scour.

650 *Influence of wetland condition on fish community composition*

651
652 The community composition of biota, including fishes, have been found to be more
653 similar during floods (Miranda 2005). Similarities in fish community composition occur because
654 annual seasonal floods homogenize floodplain environments by promoting lateral connectivity
655 which allows for the exchange of fish between the river and floodplain habitats (Hamilton &
656 Lewis 1990; Thomasz et al. 1997; Miranda 2005). Conversely, distinct fish community
657 composition is a common occurrence in wetlands with less lateral connectivity and is driven by
658 biotic interactions (e.g., predation and competition amongst fishes) and adverse water quality
659 (Gawlik et al 2002; Henning et al. 2007; Faulkner et al. 2011). Historic seasonal flooding (see
660 above) experienced in our wetlands likely drove similarities in community composition by
661 greatly promoting lateral connectivity, which allowed for greater dispersal of fishes. Sunfishes
662 and livebearers were dominant in our wetlands and these species are recognized as very common
663 floodplain dispersers capable of quickly colonizing recently flooded areas (Gkenas et al. 2011;
664 Alfermann & Miranda 2013). Many of our wetlands experienced long-lasting hydroperiods
665 which may have also been responsible for driving similarities in fish community composition as
666 these conditions have been found to ensure sunfish survival (Kushlan 1976; Hohausova et al.
667 2010; Alfermann & Miranda 2013) and possibly lead to competitive exclusion among other
668 species for food resources and prime available habitat (Carrara et al. 2012; De Bie et al. 2012).
669 The shared wetland geomorphic setting among our wetlands (i.e., riverine) may have created

670 similarities in fish communities as many our wetlands shared source pools of fish (e.g., Bayou du
671 Chien Creek, Mayfield Creek, Obion Creek, and the Mississippi River) during flooding.
672 Additionally, many wetlands were proximate to one another or located on the same easement
673 (Figure 1) allowing colonization from identical source pools of fish.

674 *Influence of wetland condition on fish diversity*

675 Even though fish community composition among wetland condition was similar, the
676 effect of wetland restoration and its influence on fish communities was evident during our study
677 as restored wetlands harbored levels of fish diversity comparable to reference wetlands and
678 greater than diversity in degraded wetlands (Figures 4, 5). Previous studies have also found
679 levels of diversity similar between reference and WRP restored wetlands (Juni & Berry 2001;
680 Benson et al. 2018). Diversity in our restored wetlands reached reference levels quickly as
681 wetlands were relatively young (i.e., 1-13 years). Similarly, Moreno-Mateos et al. (2012) found
682 that diversity in restored wetlands quickly reached reference wetland conditions (0-5 years) if the
683 wetland was in a warm climate and had a riverine geomorphic setting due to increased
684 biogeochemical functioning. The humid sub-tropical climate (i.e., hot summers and mild
685 winters) throughout our study area paired with the riverine geomorphic classification of our
686 restored wetlands probably influenced diversity in our wetlands. Restored wetlands' association
687 with two unique groups of fish (i.e., both large-river and KY-SGCN wetland fishes) (Figure 4)
688 may have also helped drive high levels of diversity in restored wetlands. Low levels of diversity
689 in degraded wetlands may have been due to lessened duration of lateral connectivity (Figure 3)
690 and/or unmitigated alterations to wetland hydrology (King et al. 2006).

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695 Management and conservation implications

696
697 The influence of hydrology on the relative abundances of large-river and KY-SGCN
698 wetland fishes was evident throughout our study. Therefore, there may be utility in using
699 hydrologic conditions as conservation management tools to influence community composition to
700 better promote the presence of specific fish taxa in restored wetlands. Ultimately, restoration
701 managers will have to consider tradeoffs in fish communities associated with the hydrologic
702 conditions they hope to achieve through restoration. For example, if restoration managers aim for
703 wetland hydrology to be influenced by the Mississippi River, large-river fishes may be promoted
704 but at the expense of wetland obligates. Promotion of large-river fishes may be important as
705 some species, like channel catfish and *Ictiobus* sp. (buffalo), have commercial and recreational
706 value. Riverine influence on wetland hydrology may, however, lead to a greater likelihood of
707 drying, presence of lotic conditions, or high magnitude depths. Drying may trap and cause
708 mortality amongst large-river fishes utilizing these wetlands. Large scale die-offs may also be
709 beneficial for other taxa such as wading birds, amphibians, and wetland plants. If restoration
710 managers desire wetland hydrology to be less influenced by the Mississippi River, wetland
711 obligates may be promoted but not necessarily large-river fishes. Promotion of wetland obligate
712 fishes is important because many of these species are KY-SGCN, which, although not
713 recreationally or commercially important, are of important conservation concern. Restoration of
714 wetlands less influenced by the Mississippi River may improve the conservation status of these
715 fishes by restoring environments that fulfill their specific habitat requirements (Eisenhour et al.
716 2018). Conservation of KY-SGCN wetland fishes is important because of their intrinsic value
717 and specific habitat requirements which may indicate proper wetland ecosystem functioning
718 (Benson et al. 2018; Simpson et al. 2021). Regardless of wetland restoration managers' goals,

719 future wetland restorations throughout the lower Ohio River tributary ichthyofaunal region of
720 Kentucky may benefit KY-SGCN wetland fishes as this area harbors high quality source pools of
721 KY-SGCN wetland fishes (Eisenhour et al. 2018; Personal communication, KDFWR).

722 Limitations to our study

723
724 Patterns in floodplain fish community composition and diversity often vary seasonally
725 because of hydrologic variation due to flooding and drying events (De Graaf 2003; Arrington &
726 Winemiller 2006; Tedesco et al. 2008). Our study did not quantify seasonal patterns of
727 community composition and diversity because we did not have adequate statistical inference due
728 to a lack of samples taken during the winter and spring at several wetlands because of seasonal
729 flooding. This lack of samples may have led to under representations in fish community
730 composition and diversity because these samples may have been the most robust as flooding
731 would have thoroughly mixed river and floodplain metacommunities. The methods used in our
732 study were limited to data collected by backpack electrofishing. Utilizing multiple sampling
733 gears may lead to better estimations of community composition and diversity. Further research in
734 western Kentucky, as well as throughout the MAV, will improve our understanding of how fish
735 respond to wetland restoration and to wetland environmental conditions. Further research should
736 include studies that examine the effects of different wetland restoration techniques on fish
737 communities and environmental conditions to inform managers which restoration techniques best
738 promote the desired fish communities. Additionally, future studies may consider choosing
739 degraded and reference sites that are also strongly influenced by the Mississippi River to
740 determine if wetland condition determines fish communities in wetlands with similar hydrology.

741
742
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744

745 Conclusions

746
 747 Wetland hydrology had the greatest influence on fish communities from the variables
 748 examined. Therefore, wetland restoration practitioners may consider focusing on specific aspects
 749 of wetland hydrology to promote desired communities or increase the presence of specific taxa.
 750 The influence of wetland restoration throughout the MAV and other large-river floodplain
 751 ecosystems may have region-wide implications on fish communities as our study found high
 752 levels of fish diversity in restored wetlands. Knowledge gaps associated with a lack of post-
 753 restoration monitoring, however, exist with large wetland restoration programs and, therefore,
 754 quantifying fish communities post-restoration provides wetland restoration programs with insight
 755 and direction for future restoration and management efforts. Undoubtedly, restoration
 756 practitioners will face and must consider tradeoffs associated with wetland restoration practices
 757 and, hopefully, these results better inform future recommendations and restoration projects.
 758 Regardless, the need remains for future studies that span across larger temporal and spatial scales
 759 to better understand how wetland restoration practices can influence the entire MAV regions.

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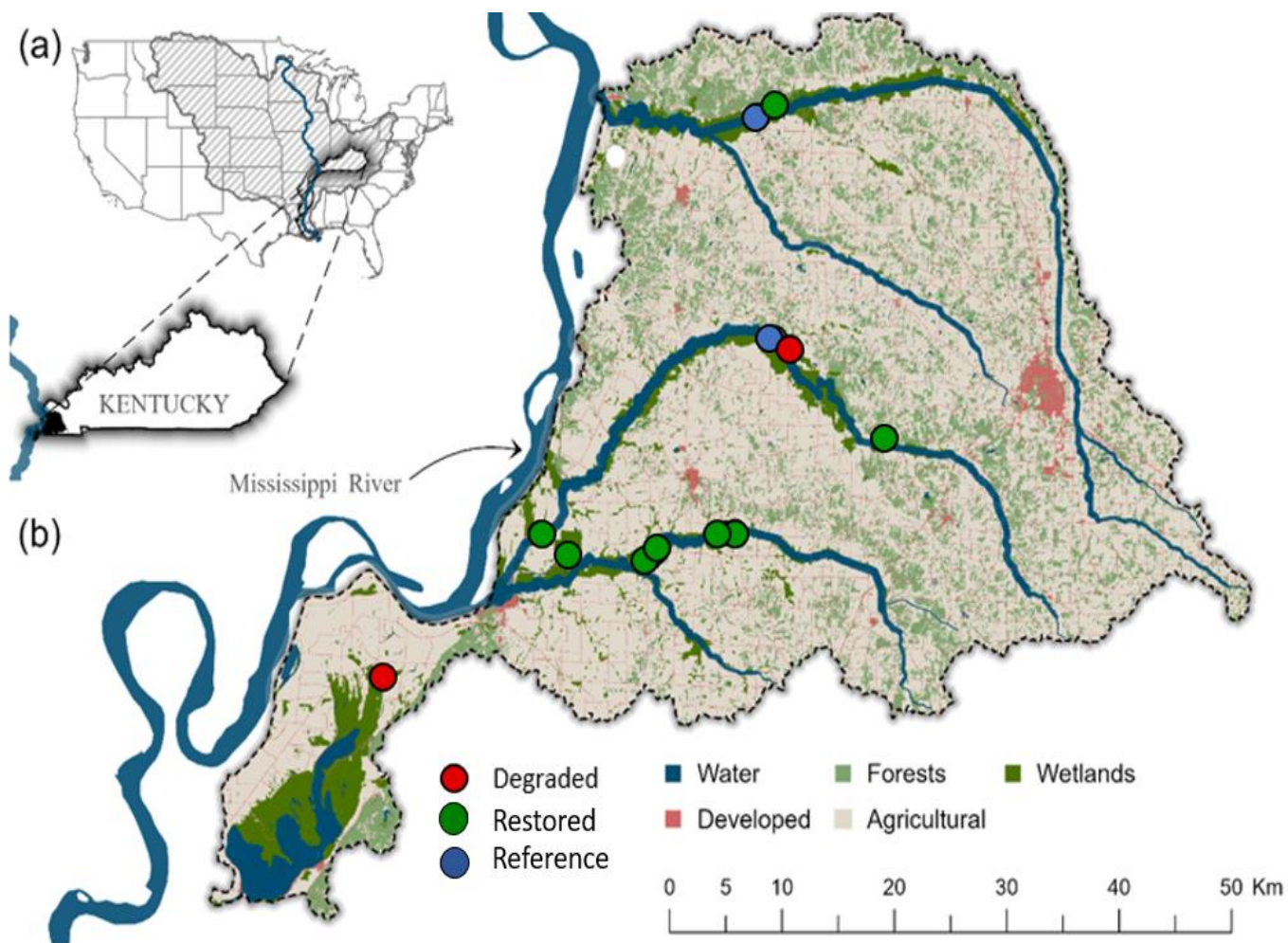
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Figures and Tables

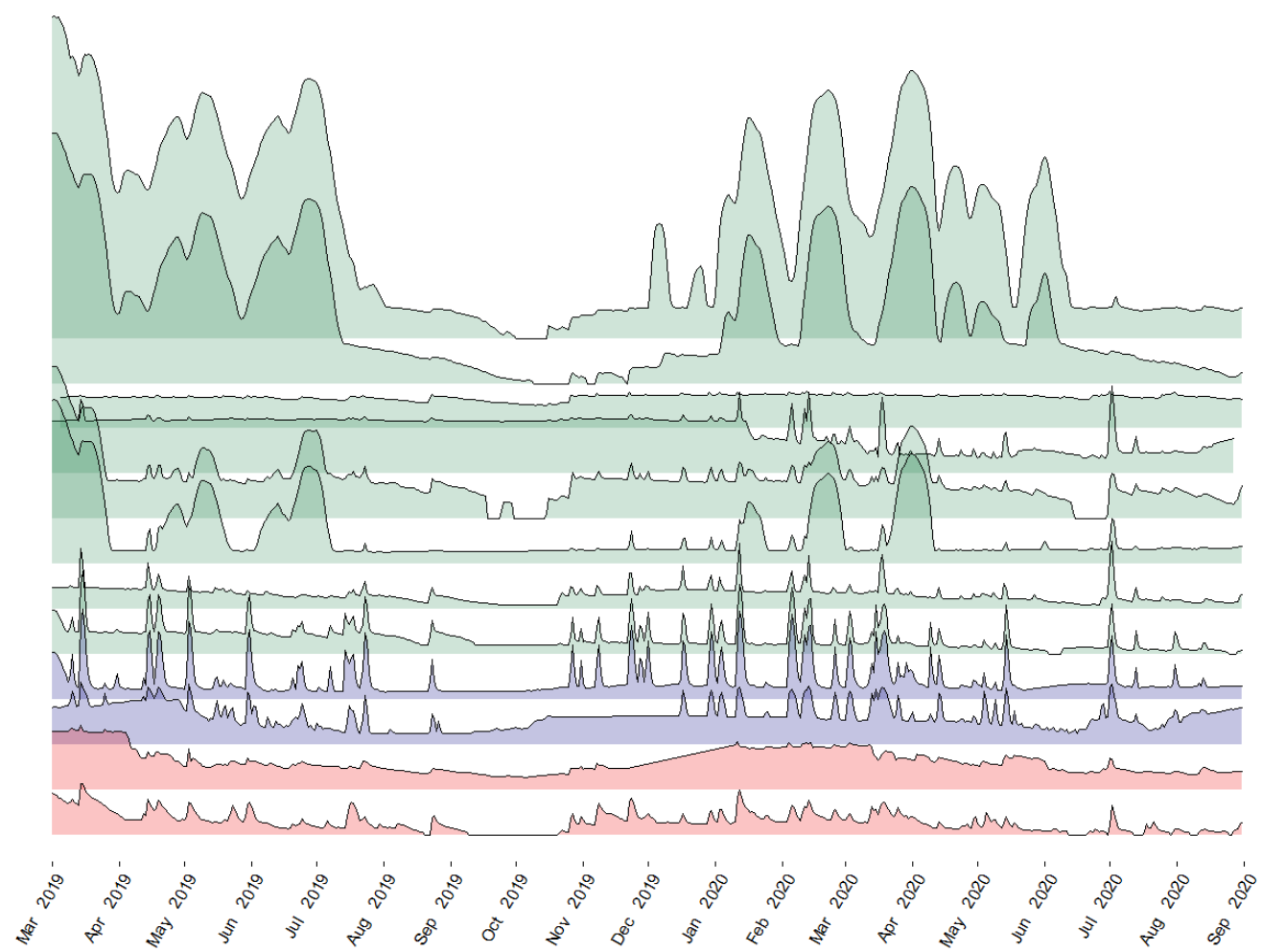


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980 **Figure 1.** (A) Location of study area in western Kentucky and part of the Mississippi River watershed, USA. (B)
 981 Twelve study wetlands include degraded (n=2), restored (n=8), and reference wetlands (n=2) (land use
 982 classifications from 2018 USDA National Imagery Program).
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Figure 2: Hydrographs of degraded (red, n=2), restored (green, n=8), and reference wetlands (blue, n=2) in western Kentucky, USA. Wetland depth was collected at 15 min intervals from March 2019 to September 2020.

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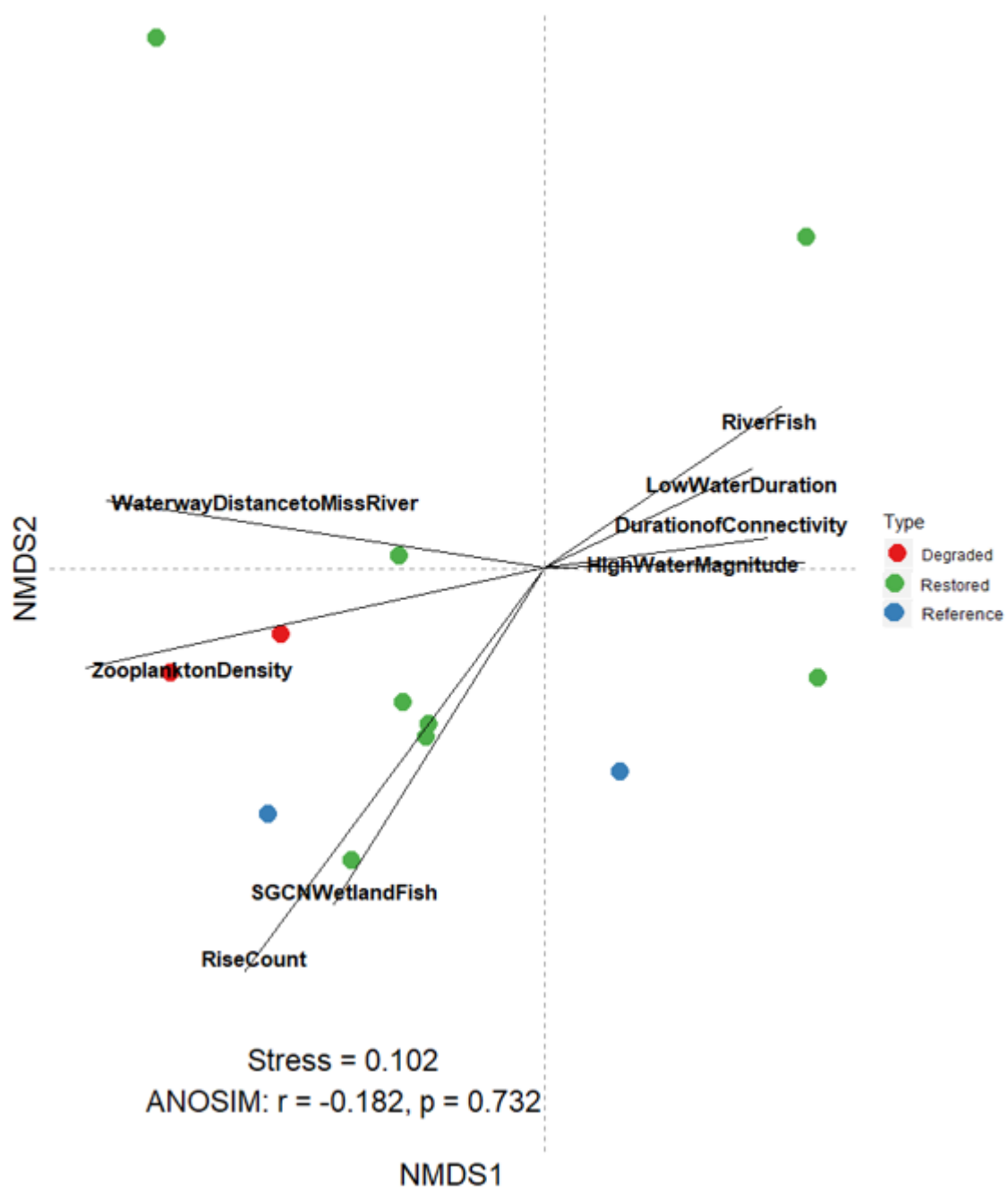
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1008 **Table 1:** Environmental metrics (28) considered for vector analysis in NMDS ordinations measured in wetlands
 1009 throughout western Kentucky, USA. Each metric was calculated using data collected over the entire sampling period
 1010 for each wetland in our study. Indicators of hydrologic alteration following Richter et al. (1996) are denoted with the
 1011 abbreviation 'IHA'.
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Metric	Definition	Measures
<u>Hydrologic metrics</u>		
Mean Depth (m)	The average wetland depth.	Magnitude
Hydroperiod	The number of days a wetland had water.	Duration
Duration of Connectivity	The percent of time a wetland exhibited lateral connectivity to its nearest stream (USGS-NWIS).	Connectivity
1-Day Maximum (m) (IHA)	The maximum single day depth recorded.	Magnitude/Duration
1-Day Minimum (m) (IHA)	The minimum single day depth recorded.	Magnitude/Duration
7-Day Maximum (m) (IHA)	The maximum 7-day rolling average recorded.	Magnitude/Duration
7-Day Minimum (m) (IHA)	The minimum 7-day rolling average recorded.	Magnitude/Duration
30-Day Maximum (m) (IHA)	The maximum 30-day rolling average recorded.	Magnitude/Duration
30-Day Minimum (m) (IHA)	The minimum 30-day rolling average recorded.	Magnitude/Duration
90-Day Maximum (m) (IHA)	The maximum 90-day rolling average recorded.	Magnitude/Duration
90-Day Minimum (m) (IHA)	The minimum 90-day rolling average recorded.	Magnitude/Duration
Minimum Date (day of the year) (IHA)	The date the lowest single recorded depth occurred.	Timing
Maximum date (day of the year) (IHA)	The date the greatest single recorded depth occurred.	Timing
Low Water Duration (days) (IHA)	The number of consecutive days depths stayed below the 25th percentile.	Duration
High Water Duration (days) (IHA)	The number of consecutive days depths stayed above the 75th percentile.	Duration
Rise Count (IHA)	The number of occurrences that wetland depth rose from the previous day.	Frequency/Rate of change
<u>Water quality metrics</u>		
Mean Dissolved Oxygen (DO) (mg/L)	The average DO measurement.	Magnitude
Minimum Dissolved Oxygen (DO) (mg/L)	The minimum single day DO recorded.	Magnitude
Mean Temperature (C°)	The average temperature recorded.	Magnitude
1-Day Maximum (C°)	The maximum single day temperature recorded.	Magnitude
1-Day Minimum (C°)	The minimum single day temperature recorded.	Magnitude

Biotic metrics

Vegetation	The average aquatic vegetation scores.	Habitat
Zooplankton Density (individuals/L)	The average zooplankton density.	Food resources
KY –WRAM Score	The KY-WRAM score indication ecological integrity (0-100) (100 indicates highest quality).	Wetland quality
<u>Hydrologic connectivity</u>		
Distance to Main Channel (m)	The distance to the main channel of the nearest stream.	Connectivity
Topography	The mean slope inside a 1km buffer around wetland.	Connectivity
Waterway Distance to Mississippi River (km)	The shortest waterway distance from each wetland to the main channel of the Mississippi River (USGS Stream Stats).	Hydrologic influence of Mississippi River
Elevation (m)	The median of elevation (n=50) (USGS Stream Stats).	Connectivity



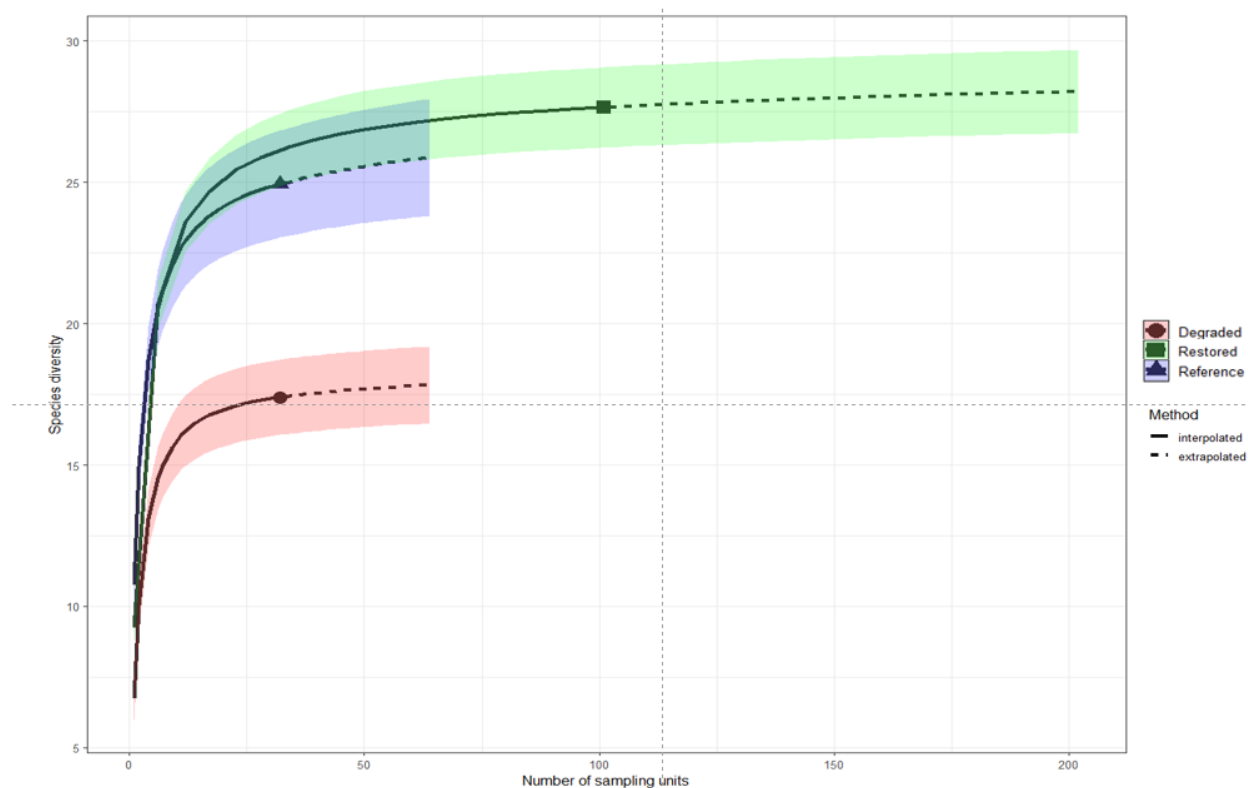
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Figure 3: NMDS ordination of fish community composition from wetlands in western Kentucky, USA. Ordination is based on per taxa CPUE from electrofishing that occurred monthly from April 2019 to August 2020. Symbol colors indicate level of wetland condition (degraded, restored, reference). All variables included in Table 1 were tested and only significant variables were placed onto ordination as vectors. Vectors representing the relative abundance of river fish and KY-SGCN wetland fish were also placed onto ordination. The vector High Water Magnitude is a combination of the metrics 1-Day Maximum (m), 7-Day Maximum (m), 30-Day Maximum (m), and 90-Day Maximum (m). See Table 2 for definitions of variables used as vectors.

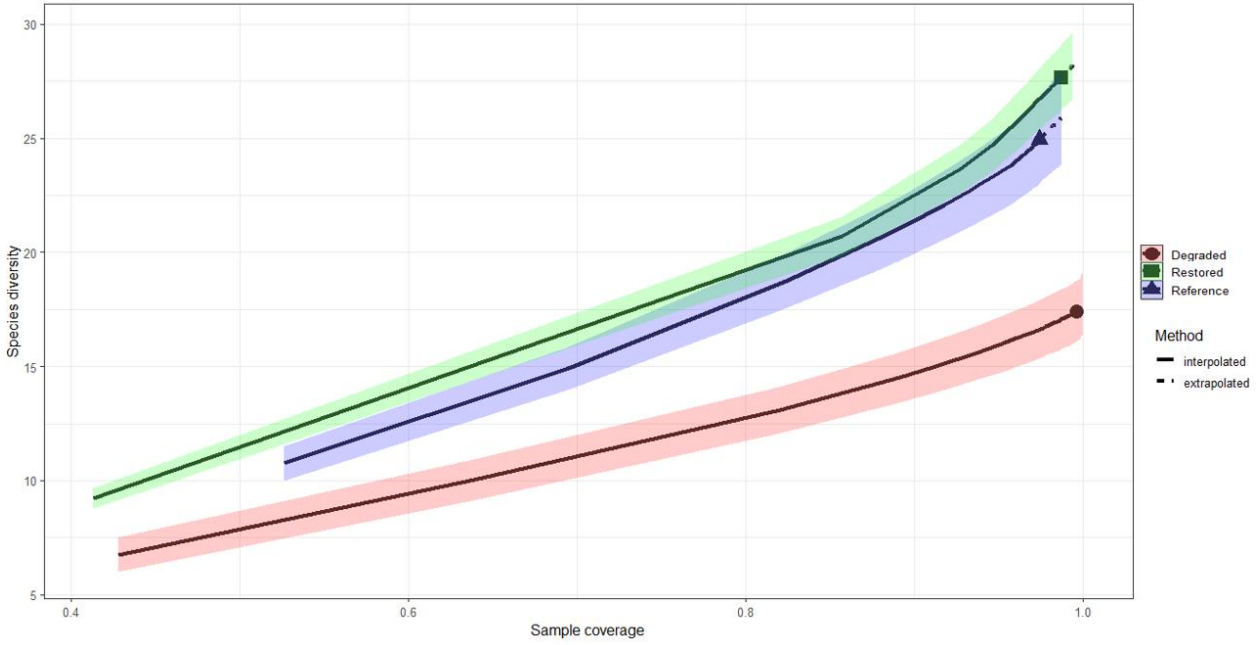
1026 **Table 2:** Correlation coefficient and p-values associated for vectors placed onto NMDS ordination that had
 1027 significant associations with fish community composition.
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Vector	Correlation Coefficient	P-Value
High Water Magnitude	0.742	0.009
Zooplankton Density	0.620	0.015
Low Water Duration	0.599	0.025
Waterway Distance to Miss River	0.579	0.026
Rise Count	0.545	0.029
Duration of Connectivity	0.609	0.043
River Fish	0.784	0.013
SGCN Wetland Fish	0.358	0.128

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1031 **Figure 4:** Hill-Shannon diversity estimates of wetland fish communities by wetland condition using incidence-based
 1032 rarefaction and extrapolation. Curves are based on electrofishing data collected from April 2019 to August 2020 in
 1033 western Kentucky, USA. All abundance-based extrapolation curves were plotted to achieve 95% coverage.
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Figure 5: Hill-Shannon diversity estimates of wetland fish communities by wetland condition using coverage-based rarefaction and extrapolation. Curves are based on electrofishing data collected from April 2019 to August 2020 in western Kentucky, USA. All abundance-based extrapolation curves were plotted to achieve 95% coverage.



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Figure 6: Photograph of a low water induced fish kill in a wetland with a short waterway distance to the Mississippi River (strongly influenced by the river) in western Kentucky, USA. Photograph was taken during August of 2020.

Chapter 2

Wetland elevation is an important determinant of larval fish community composition

Abstract

A major goal of the Wetlands Reserve Program (WRP) is to create wildlife benefits through wetland restoration. WRP, however, may overlook wetland elevation in their selection criteria which can have important implications on biological communities. The objectives of our study were to (1) examine if differences in wetland elevation led to variation in larval fish communities, (2) determine the relationships between larval fish community composition and wetland environmental conditions, and (3) determine the influence of differences in wetland condition (e.g., degraded, restored, and reference) on larval fish communities. From March through August 2020, we performed monthly dipnet surveys for larval fishes in lowland (n=3), transitional (n=2), and upland (n=7) wetlands throughout western Kentucky. Analysis of variance using distance matrices (ADONIS) and analysis of similarity percentages (SIMPER) were used in conjunction with non-metric multidimensional scaling to visualize larval fish community composition, quantify differences in composition amongst wetland elevation, and determine which species significantly contributed to differences in composition. Vector analysis was used to determine which environmental factors most affect larval fish community composition. Furthermore, larval fish diversity was quantified using Hill diversity (Hill-Shannon; $q=1$) and compared using 95% confidence intervals. Our findings suggest that differences in wetland elevation were characterized by differences in hydrologic conditions, which led to differences in larval fish community composition. Differences in community composition were driven by greater abundances of Hypophthalmichthys (bighead carp) and Pomoxis (crappie) in lowland wetlands when compared to upland wetlands. Wetland elevation did not lead to differences in diversity. Furthermore, wetland condition did not lead to differences in larval fish community composition or diversity. Our study demonstrated that wetland elevation and hydrology are important factors for wetland restoration managers to consider when selecting sites for future wetland restorations.

Introduction

Many of the United States non-coastal wetland resources exist as riverine wetlands throughout the Mississippi Alluvial Valley (MAV). Riverine wetlands are floodplain areas that are periodically inundated by direct overland flow, backwater flooding from adjacent rivers or streams, or surface runoff from precipitation (Brinson 1993; Hunter et al. 2008). As in other wetlands, the quantity and timing of hydrology define the structure and function of riverine wetlands (Faulkner & Patrick 1992; Corstanje & Reddy 2004). The hydrology of riverine

1085 wetlands throughout the MAV is largely influenced by the Mississippi River due to its seasonal
1086 flood pulse (winter and spring) and by local climate after flood waters recede (summer and fall)
1087 (Junk et al. 1989; Mitsch & Gosselink 2015). Like the biogeochemical processes and vegetation
1088 communities that rely on the hydrology of these wetlands, so do wildlife communities.

1089 The timing, duration, and magnitude of the MAV's hydrology is especially important to
1090 floodplain fishes as it fulfills a variety of life history requirements. The MAV's hydrology is
1091 flood pulsed in nature which promotes high levels of connectivity between the river channel and
1092 nearby wetlands and coincides with the seasonally increased energetic needs required for
1093 reproduction, i.e., spawning and healthy eggs (Welcomme 1985; Junk et al. 1989; Bayley 1991;
1094 Winemiller & Rose 1992; King & Keeland 1999; Tocker et al. 2000; Baber et al. 2002; Kluender
1095 et al 2015). The timing of connectivity, however, must directly correspond with optimal
1096 spawning temperatures to ensure reproductive success (Lubinski et al. 1991; Sparks et al. 1998).
1097 Many demersal spawners (i.e., those that require structure to lay eggs), including main channel
1098 obligates (e.g., blue sucker), require the habitat-rich littoral areas in riverine wetlands to
1099 successfully lay their eggs (Adams et al. 2006). Once hatched, many larval fish utilize food-rich
1100 (e.g., zooplankton) littoral areas to acquire energy required to grow and develop before
1101 emigrating back into their population (Chick & McIvor 1997; Dettmers et al. 2001; Adams et al.
1102 2008; Kluender et al. 2015).

1103 Despite the overwhelming importance of riverine wetlands for floodplain fishes
1104 throughout the MAV, the existence of these wetlands is in danger. Since European colonization,
1105 the MAV has lost 70-84% of its wetlands which once spanned across a vast 10 million ha
1106 (Haynes & Egan 2004; Frederickson 2005; Faulkner et al. 2011). Radical alterations to regional
1107 hydrology (e.g., dams, ditches, levees, and tile drains) paired with direct land-use conversion to

1108 agriculture have largely been responsible for these losses (Semlitsch 2000; Haynes & Egan 2004;
1109 King et al. 2006; Faulkner et al. 2011). Changes in hydrology and conversion to agriculture
1110 throughout a river-floodplain system are considered the most serious and pervasive
1111 anthropogenic threats to the system's ecological integrity because it separates the river from its
1112 floodplain (Tockner et al. 2000; Poff et al. 2007). Loss of lateral (e.g., river to floodplain)
1113 connectivity greatly diminishes riverine wetland function and wildlife benefits (Zedler 2000;
1114 Rewa 2005; Hunter et al. 2008; Moreno-Mateos et al. 2012; King & Keim 2019). Dramatic
1115 wetland loss throughout the MAV, however, has not gone unnoticed and the past 30 years have
1116 seen the implementation of large-scale wetland restorations performed at state and federal levels.

1117 The Wetlands Reserve Program (WRP) is one such large-scale federal-level restoration
1118 program. Implemented by the Natural Resources Conservation Service (NRCS), the WRP
1119 conducts wetland restorations throughout the United States. The goals of the WRP are to restore
1120 wetland ecological function and wildlife benefits (Natural Resources Conservation Service
1121 2013a). The WRP focuses on using a combination of reforestation and hydrologic restoration
1122 techniques to restore historic wetland function (King & Keeland 1999; Hayes & Egan 2004;
1123 Rewa 2005). Special emphasis is placed on hydrologic restoration as hydrology drives wetland
1124 function and wildlife benefits (Bedford 1996; Brinson & Rheinhardt 1998; Zedler 2000; Haynes
1125 & Egan 2004; Rewa 2005; Brauman et al. 2007; Hunter et al. 2008; Faulkner et al. 2011; King &
1126 Keim 2019).

1127 When selecting a site for wetland restoration, WRP ranking criteria are extensive and
1128 many hydrologic aspects are taken into consideration. Elevation of a wetland, however, is not
1129 always considered in selection criteria (NRCS WRP Ranking Criteria 2008). WRP typically
1130 employs restoration in riverine wetlands to maximize ecosystem function. Wetland elevation

1131 may result in large differences in environmental conditions, i.e., hydrology and water quality,
1132 even if hydrogeomorphic classifications are similar (Brinson 1993; Acreman & Holden 2013).
1133 For example, a riverine wetland located in the Mississippi River's floodplain may experience
1134 greater magnitude and duration of flooding during seasonal flooding events due to its massive
1135 watershed, while another riverine wetland located on an upstream tributary of the Mississippi
1136 River, i.e., smaller watershed, experiences relatively smaller magnitude and shorter duration
1137 flooding as its hydrology is less influenced by the seasonal flooding of the Mississippi River and
1138 more by inputs from local precipitation (Euliss et al. 2004; Acreman & Holden 2013). The
1139 effects of elevation on local wetland hydrology and water quality may have implications on the
1140 response of wildlife to wetland restoration as fish communities have been shown to be strongly
1141 influenced by these environmental conditions (Winter 2001; Euliss et al. 2004; Miranda & Lucas
1142 2004; Miranda 2005; Miranda 2010; Dembkowski & Miranda 2012).

1143 Understanding the factors that structure floodplain wetland larval fish communities is
1144 important because larval fish are reliant on specific hydrologic conditions and their recruitment
1145 is important in maintaining floodplain fish communities. Therefore, choosing wetland restoration
1146 sites that promote larval fish usage may indicate proper hydrologic function and ultimately
1147 benefit floodplain wildlife. The primary goals of this research were to (1) examine if differences
1148 in elevation led to differences in larval fish communities, (2) determine the relationships between
1149 larval fish community composition and environmental conditions, i.e., hydrology, water quality,
1150 and biota, in wetlands, and (3) examine if differences in wetland condition (e.g., degraded,
1151 restored, and reference) influenced larval fish communities. Wetland elevation was predicted to
1152 lead to differences in larval fish community composition and diversity, that differences in larval
1153 fish communities would reflect differences in environmental conditions and lastly, that larval

1154 fish communities would differ based on wetland condition. Hopefully, larval fish/environment
1155 relationships from this research can inform future wetland restoration management decisions and
1156 ensure the most effective management decisions (Merkle et al. 2019).

1157 **Methods**

1158 Study Area

1159 Our study was conducted in the Mississippi Alluvial Plain and Mississippi Valley Loess
1160 Plain ecoregions of western Kentucky (Omerink 1987). Historically, wetland resources in this
1161 region were characterized by bottomland hardwood forests and stream floodplains that
1162 experienced dramatic but predictable hydroperiods seasonally influenced by the upper
1163 Mississippi River watershed (King et al. 2006; Mitsch & Gosselink 2015). A substantial amount
1164 of wetland resources in this region still exist and are located along three major tributaries
1165 (Mayfield Creek, Obion Creek, and Bayou du Chien Creek) that drain approximately 2,800 km²
1166 into the Mississippi River. These wetlands are highly fragmented and suffer from changes to
1167 surrounding land use and regional hydrologic modifications (Frederickson 2005, King et al.
1168 2006). Despite these anthropogenic influences, the region retains some features of a large river
1169 floodplain; for example, seasonally high discharges reconnect the floodplain in the winter and
1170 spring (Mitsch & Gosselink 2015). Current land use of this region of western Kentucky is
1171 dominated by cultivated crops (64%) and forested floodplain wetlands (25%) (Dewitz 2019).

1172 Wetland Selection

1173 Twelve wetlands in far western Kentucky (Figure 1) were sampled. Wetland hydrology
1174 exhibited differences based on elevation, measured in meters above sea level (MASL) (Figure 2).
1175 Following differences in hydrology based on elevation, wetlands were classified as lowland

1176 (n=3) if their elevation was < 91 (MASL); transitional (n=2) if their elevation was > 91 but < 97
1177 MASL; and upland (n=7) if their elevation was > 97 MASL.

1178 Eight of our wetlands were restored by the WRP in western Kentucky and were used in
1179 analyses to compare differences in wetland elevation and condition (Figure 1). Before wetland
1180 selection, pertinent WRP easement information was obtained from the National Resources
1181 Conservation Service (NRCS) (including landowner contact, restoration type, restoration age,
1182 restoration plans). After obtaining easement information, the following criteria were used to
1183 select restored wetlands: (1) location on one of the regional tributaries to the Mississippi River
1184 (Figure 1), (2) similarity of hydrogeomorphic wetland class (riverine following Brinson et al.
1185 1993), (3) hydroperiod (semi-permanent to permanent following Cowardin et al. 1995), and (4)
1186 size (1-20 ha). Additionally, low (degraded; n=2) and high-quality wetlands (reference; n=2)
1187 were sampled to compare WRP restored wetlands with (Figure 1). Degraded wetlands were once
1188 natural wetlands that have experienced dramatic local hydrologic alterations for agricultural
1189 purposes, but still exhibit some wetland characteristics. Both of our degraded wetlands were in
1190 active agricultural fields. Reference wetlands were not subjected to local hydrologic alterations,
1191 but ultimately still exist within a highly altered landscape. One of our reference sites was a
1192 forested wetland located on a KY Wildlife Management Area; the other site was a bottomland
1193 hardwood swamp positioned on an upstream portion of one of our WRP easements.

1194 Larval fish sampling

1195 Larval fish were sampled monthly at all study sites from March 2020 through August
1196 2020 using a 20 jab dipnet (30.5 cm x 25.4 cm x 55.9 cm, 500µm) survey from all available areas
1197 (open water, vegetation, woody debris). Larval fish were anesthetized using clove oil, preserved
1198 in a 10% formalin solution, and were later enumerated and identified to genus using Auer

1199 (1982). Dipnet surveys were standardized by the number of jabs and catch per unit effort (CPUE)
1200 was considered as the abundance of each taxon captured from each survey. Monthly CPUE of
1201 each taxon was then averaged to give one CPUE value to be used for NMDS ordinations.

1202 Environmental influences on larval fish community composition

1203 Twenty-six environmental predictor variables (see below for explanations; Table 1) were
1204 quantified to examine their influence on larval fish community composition. Environmental
1205 predictor variables were to examine the influence of hydrology, water quality, landscape, and
1206 biota on community composition.

1207 Hydrology

1208 Surface water level changes were recorded in each wetland from March 2019 to August
1209 2020 using water level loggers (HOBO® U20-001-04, Onset Computer Corporation). One
1210 logger was deployed in each wetland in the deepest wade-able location. Depth was recorded
1211 every 15 minutes. Water level data was averaged per day and daily depths were then used to
1212 calculate mean depth, hydroperiod, and 13 indicators of hydrologic alteration (IHA) based on
1213 Richter et al. (1996) (Table 1). The percent of time each wetland experienced connectivity to its
1214 associated stream was estimated by determining the stream depth at which each wetland was
1215 connected to its stream and then calculating the percent of time the stream was greater than that
1216 depth. For wetlands on upstream portions of tributaries, stream depth was determined by using
1217 the United States Geological Survey stream gauge 07024000 (USGS 2019-2020) on the Bayou
1218 du Chien and for wetlands located on the Mississippi River floodplain stream depth was
1219 determined by using the United States Geological Survey stream gauge 07022000 (USGS 2019-
1220 2020) on the Mississippi River.

1221

1222 Hydrologic connectivity

1223 Two landscape variables were calculated to indirectly quantify hydrologic connectivity
1224 between wetlands and other permanent bodies of water. Distance to Main Channel was
1225 quantified by measuring the straight-line distance from the center of each wetland to the main
1226 channel of the nearest major stream in ArcGIS Pro (Version 2.7, Esri Inc.). The “elevation
1227 profile” tool in USGS Stream Stats was used to delineate each wetland’s boundary and assign
1228 elevation (m) values for 50 different locations within each wetland. Elevation was calculated by
1229 taking the median of each wetland’s 50 elevation values.

1230 Water quality

1231 Changes in water temperature (°C) and dissolved oxygen (DO) (mg/L) were recorded in
1232 each wetland from March 2019 to August 2020 using multi-parameter sondes (YSI® EXO2,
1233 Xylem Incorporated). One sonde was suspended in each wetland approximately midway in the
1234 water column in the deepest accessible location. Sondes recorded data at 15-minute intervals.
1235 Water temperature and DO were averaged per day and then used to calculate minimum
1236 temperature, maximum temperature, mean temperature, minimum DO, and mean DO for each
1237 wetland.

1238 Biotic variables

1239 Zooplankton communities were sampled monthly from April 2019 through August 2020
1240 at each wetland using a 9-cm diameter littoral sampling tube following Pennak (1962). Each
1241 sample consisted of three replicates averaged together, one each taken from open water, edge of
1242 wetland, and dense vegetation. Samples poured into a volumetric container where volume (L)
1243 was recorded, rinsed through a 43-µm sieve, and preserved in 4% buffered formalin solution. In
1244 the laboratory, samples were subsampled to a maximum 1/8 using a Folsom Plankton Splitter

1245 (WILDSCO, Saginaw, MI) so that at least 50 zooplankters were found per sample. Cladocerans
1246 and copepods were enumerated but rotifers were excluded. After scaling back up to 100% from
1247 the subsampled fraction, density was calculated by dividing the abundance of each sample by the
1248 original volume of water.

1249 Aquatic vegetation was sampled monthly at each site between April 2020 and August
1250 2020, which corresponded with the region's growing season, by using a 1m-by-1m quadrat. Each
1251 sample consisted of nine replicates averaged together: three each taken from open water, wetland
1252 edge, and dense vegetation. Percent cover of three aquatic vegetation groups (aquatic emergent,
1253 aquatic submergent, aquatic floating) was estimated using six cover categories (1 = 0-10%, 2=
1254 11-20%, 3 = 21-40%, 4 = 41-60%, 5= 61-80%, 6 = 81-100%). The midpoint of the cover
1255 categories was used for percent cover calculations.

1256 The Kentucky Wetland Rapid Assessment Method (KY-WRAM) (Kentucky Division of
1257 Water, 2016) was used to assess for general wetland integrity and function. The KY-WRAM is
1258 used to evaluate overall wetland function and ecological integrity and was developed for use by
1259 regulatory agencies in wetland permitting decisions. The KY-WRAM assumes that wetlands
1260 with high ecological integrity also have high wetland function. The KY-WRAM metrics were
1261 developed to capture a full range of potential disturbances to wetland integrity and fall into six
1262 categories: size and distribution; buffer and intensity of surrounding land uses; hydrology;
1263 habitat alteration and habitat structure development; special wetlands; vegetation, interspersions,
1264 and habitat features. Each category is subdivided into additional metrics. The special wetlands
1265 category was omitted because we were unable to gain access to the required information (all
1266 wetlands received a score of 0 for this category). The final KY-WRAM score is the sum of all
1267 the metric scores and ranges from 0 (very poor condition) to 90 (reference condition). Out of the

1268 metrics, 10 were evaluated using ArcGIS Pro® software (Version 2.7, Esri Inc.) and high
1269 resolution orthoimages (USDA 2018), and 10 were assessed in the field during a one-hour site
1270 visit per wetland during July - September 2020.

1271 Statistical analyses

1272 Statistical analyses were carried out using R statistical software (version 4.0.5) (R core
1273 team 2021) and the ‘Vegan’ software package (Oskansen et al. 2013). Nonmetric
1274 multidimensional scaling (NMDS) was used to visualize variation in larval fish community
1275 composition among wetland elevation (lowland, transitional, upland) and wetland condition
1276 (degraded, restored, reference). Experimental units in ordinations were defined as the average
1277 CPUE of all sampling events at each wetland in multivariate space. The input data for NMDS
1278 were defined by the CPUE of each larval fish taxa present from each wetland. No fish species
1279 were removed prior to ordination. Ordinations were constructed with Bray–Curtis distances, run
1280 for the maximum number of iterations (300), and chosen for minimum stress values.
1281 Dimensional solutions, stepping down from six to one, were tested and determined by the use
1282 and examination of individual scree plots (McCune & Grace 2002).

1283 Analysis of variance using distance matrices (ADONIS) was used to statistically test
1284 larval fish community composition for differences among *a priori* groups of wetland elevation
1285 and condition. The Bray-Curtis distance was used in ADONIS and run for 9999 maximum
1286 iterations generating *r*-values between -1 and 1. Positive values indicate differences among
1287 groups and significance was tested at $\alpha = 0.05$.

1288 Additionally, analysis of similarity percentages (SIMPER; Clarke 1993) was performed
1289 to make pairwise comparisons amongst wetland elevations. SIMPER assesses the contribution of
1290 individual species to the dissimilarity between objects in a Bray-Curtis dissimilarity matrix. P-

1291 values were considered significant at the $\alpha = 0.05$ level This allows the identification of species
1292 that are likely to be major contributors to differences between groups detected by methods such
1293 as ADONIS (Clarke and Warwick 2001).

1294 The influence of environmental predictor variables (Table 1) was examined on larval fish
1295 community composition using vector fitting analysis with the function ‘ENVFIT’ (R package
1296 ‘vegan’) on NMDS ordinations. Each variable’s association with each experimental unit was
1297 indicated by the vector’s direction while the strength of each variable’s association was indicated
1298 by the vector’s length (McCune & Grace 2002). Vector significance ($\alpha = 0.05$) was estimated
1299 using 999 random permutations of the data (Faith & Norris 1989). For ease of interpretation, the
1300 hydrologic variables 1-Day Maximum (m), 30-Day Maximum (m), and 90-Day Maximum were
1301 grouped into one vector labeled High Water Magnitude on ordination plots because of high
1302 correlation (overlapping angle and vector magnitude) among variables (Flinn et al. 2008).

1303 Diversity

1304 Hill diversity was calculated for each wetland. Hill diversity was chosen due to its
1305 numerous advantages over other diversity indices (see Chao et al. 2014) and were calculated
1306 according to Hsieh et al. (2016), and Chao et al. (2014). See Jost (2006) for a more thorough
1307 review of the benefits of Hill diversity in relation to other indices or Roswell et al. (2021) for a
1308 current consensus within the ecological community.

1309 Before calculating Hill diversity, samples were standardized by 90% coverage (Chao &
1310 Jost 2012; Chao et al. 2014; Roswell et al. 2021; R package iNEXT). Coverage is a relatively
1311 new method of sample standardization in ecology that measures sample completeness and
1312 accounts for the abundance of species in the sampled community. Coverage estimates the
1313 proportion of individuals in the community that belong to species present in a sample (Roswell et

1314 al. 2021). For example, achieving coverage of 90% means that 10% of individuals in the
1315 community were not sampled.

1316 Hill diversity varies based on the choice of exponent used. Ecologists most commonly
1317 use $q = 0$ (species richness), $q = 1$ (Hill-Shannon diversity), and $q = 2$ (Hill-Simpson diversity).
1318 Hill-Shannon diversity ($q = 1$) was used because it results in all individuals being considered
1319 equally as it counts species proportionately to their abundance of incidence (Chao et al. 2014).
1320 Genus-level richness was used as opposed to species-level richness because of limitations in
1321 identification. Hill-Shannon diversity was calculated using the iNEXT package which provides
1322 functions to compute the most widely used Hill numbers ($q = 1$, $q = 2$, $q = 3$) for individual-
1323 based abundance data or sampling-unit based incidence data. Incidence data was used because it
1324 suitably represents our sampling methods and because Colwell et al. 2012, Chao et al. (2014),
1325 and Chao & Colwell (2017) demonstrated that incidence data allows for biological inference just
1326 as powerful as abundance-data based approaches. Our input data for the iNEXT package
1327 consisted of genus-specific incidence data from each sample from each wetland which was
1328 categorized by each sample's wetland elevation and condition.

1329 To describe patterns in Hill-Shannon diversity, sample- and coverage-based rarefaction
1330 and extrapolation curves were generated using the "estimate d" function (R package iNEXT) to
1331 determine how diversity increases with increasing sampling effort and completeness. Rarefaction
1332 and extrapolation of Hill-Shannon diversity were conducted according to Hsieh et al. (2016) and
1333 further discussed in Colwell et al. (2012), Chao & Jost (2012), and Chao et al. (2014). Sample-
1334 based curves evaluated the number of individuals in a sample by plotting diversity estimates in
1335 relation to the number of sampling units. Coverage-based curves were plotted against rarefied
1336 sample completeness to illustrate diversity estimates in relation to sample coverage. All

1337 extrapolation curves were plotted using a doubling in sample size, and 500 bootstrap replicates
1338 were used to estimate 95% confidence intervals. Confidence intervals, a known alternative to
1339 standard statistical testing (Magurran 2004; Colwell et al. 2004), were used to determine if
1340 differences between wetland elevation and condition were statistically significant.
1341 Nonoverlapping 95% confidence intervals, whether rarefied or extrapolated, were considered to
1342 indicate significant differences at $\alpha = 0.05$ (Chao & Jost, 2012; Chao et al., 2014).

1343 **Results**

1344

1345 Larval fish sampling

1346

1347 Larval fish were present at each wetland. In total, 2081 larval fish from 11 families and
1348 16 genera were collected across all wetlands. The mean CPUE (abundance of taxa per survey) at
1349 lowland wetlands was 98 ± 76.3 (SE), 10 ± 3.00 (SE) at upland wetlands, and 8 ± 2.28 (SE) at
1350 transitional wetlands. Twelve of the 16 genera were collected in uplands wetlands, 10 genera in
1351 lowland wetlands, and 9 genera in transitional wetlands. *Dorosoma* (shads), *Ictiobus* (buffalo),
1352 *Lepisosteus* (gars), and *Pomoxis* (crappies) were only found in lowland wetlands. *Ameiurus*
1353 (bullheads), *Erimyzon* (chubsuckers), and *Umbra* (mudminnows) were only found in upland
1354 wetlands. The mean CPUE at restored wetlands was 43 ± 24.7 (SE), 8 ± 2.03 (SE) at reference
1355 wetlands, and 6 ± 3.33 (SE) at degraded wetlands. Fifteen of the 16 genera were collected in
1356 restored wetlands, 10 genera were collected in reference wetlands, and 5 genera were collected in
1357 degraded wetlands. Bullheads, shads, *Hypophthalmichthys* (bighead carp: invasive genus),
1358 buffalo, gars, and crappies were only found in restored wetlands. Mudminnows were only found
1359 in reference wetlands. *Lepomis* (true sunfish) was by far the most ubiquitous genus across
1360 wetland elevation and condition. Overall community composition of all wetlands was dominated
1361 by bighead carp (64%) and true sunfish (27%) with all other genera contributing $\leq 2\%$ each.

1362 Differences in larval fish community composition among wetland elevation and condition

1363 NMDS ordination based on larval fish CPUE data revealed considerable separation by
1364 wetland elevation. Final NMDS solutions consisted of two dimensions with a relatively low final
1365 stress values (0.137) and high interpretability compared to alternative solutions (Figure 3).
1366 Results of ADONIS analysis confirmed that larval fish community composition among lowland,
1367 transitional, and upland wetlands was significantly different ($r^2 = 0.282$, $p = 0.047$).

1368 SIMPER analysis identified genera that led to differences among wetland elevation
1369 (Table 2a, b, c). Bighead carp (54.2 %) and true sunfish (9.3%) contributed the most to the
1370 differences between lowland and transitional wetlands, although no genera's contribution was
1371 significant ($p > 0.05$) (Table 2a). True sunfish (28.1%) and *Elassoma* (pygmy sunfish) (8.5%)
1372 contributed the most to the differences between upland and transitional wetlands although no
1373 genera's contribution was significant ($p > 0.05$) (Table 2b). Bighead carp (50.2%) and true
1374 sunfish (25.8%) contributed the most to differences between lowland and upland wetlands.
1375 Bighead carp ($p = 0.048$) and crappies ($p = 0.013$) were found to be more important in lowland
1376 wetlands (Table 2c) and their contribution was significant to the differences between lowland
1377 and upland wetlands.

1378 NMDS ordination based on larval fish CPUE data revealed little separation by wetland
1379 condition. Results of ADONIS confirmed that degraded, restored, and reference wetlands were
1380 not significantly different ($r^2 = 0.109$, $p = 0.955$).

1381 Environmental influences on larval fish community composition

1382 Significant environmental variables that helped explain differences in larval fish
1383 community composition were overlaid as vectors onto NMDS ordination (Figure 3, Table 3).
1384 Environmental vectors displayed a strong vertical gradient: increasing High Water Magnitude,

1385 Low Water Duration, and Percent Time Connected were all associated with each other and were
1386 negatively associated with Elevation, which also displayed a vertical gradient. Larval fish
1387 community composition based on wetland elevation displayed more of a horizontal gradient than
1388 vertical gradient, but still, followed the vertical gradient of Elevation. Larval fish community
1389 composition found in lowland wetlands were most associated with High Water Magnitude, Low
1390 Water Duration, and Percent Time Connected and negatively associated with Elevation. Larval
1391 fish community composition in upland wetlands had varying, but mostly negative association
1392 with these vectors. Transitional wetlands had little to no association with any vectors.

1393 *Influence of elevation on fish diversity*

1394 Rarefaction curve analysis did not detect significant differences in Hill-Shannon diversity
1395 among wetland elevation as overlap existed in the 95% confidence intervals among lowland
1396 11.1, CI [7.3, 18.7], transitional 6.5, CI [5.5, 8.9], and upland wetlands 7.4, CI [6.6, 9.1] as the
1397 number of sampling units increased (Figure 4). The estimated curve patterns of Hill-Shannon
1398 diversity accumulation per sampling unit for transitional and upland wetlands were approaching
1399 asymptotic plateau, suggesting that the sampling strategy was sufficient in revealing true
1400 diversity associated with these wetlands. The estimated curve patterns of Hill-Shannon diversity
1401 accumulation per sampling unit for lowland wetlands, however, did not approach asymptotic
1402 plateau suggesting that the sampling strategy was insufficient in revealing the true diversity,
1403 likely leading to an underrepresentation of diversity. Coverage-based rarefaction and
1404 extrapolation further indicated that sample completeness was sufficient for transitional and
1405 upland wetlands as coverage values were greater than 90% (92% and 95% respectively) but was
1406 insufficient for lowland wetlands as coverage values (74%) were less than 90%.

1407

1408 *Influence of wetland condition on diversity*

1409 Rarefaction curve analysis did not detect significant differences in Hill-Shannon diversity
1410 of larval fish communities among wetland condition as overlap existed in the 95% confidence
1411 intervals among degraded 7.8, CI [4.0, 16.0], restored 12.5, CI [9.4, 17.2], and reference
1412 wetlands 10.2, CI [7.4, 14.6] as the number of sampling units increased (Figure 5). The estimated
1413 curve pattern of Hill-Shannon diversity accumulation per sampling unit for restored wetlands
1414 was approaching asymptotic plateau, suggesting that the sampling strategy was sufficient in
1415 revealing true diversity associated with these wetlands. The estimated curve patterns of Hill-
1416 Shannon diversity accumulation per sampling unit for degraded and reference wetlands,
1417 however, did not approach asymptotic plateau suggesting that the sampling strategy was
1418 insufficient in revealing the true diversity associated with these wetlands leading to an under
1419 representation of degraded and reference wetland larval fish diversity. Underrepresentation of
1420 true diversity was also apparent in coverage-based rarefaction and extrapolation estimates, which
1421 indicated that sample completeness was insufficient for degraded (54%), restored (87.5%), and
1422 reference (83%) wetlands.

1423 **Discussion**

1424 Our results indicated that wetland elevation was an important factor in influencing larval
1425 fish community composition, likely via differences in wetland hydrology along elevation
1426 gradients. Other studies have found that elevation is important in determining wetland hydrology
1427 (Brinson 1993; Euliss et al. 2004). Lowland wetlands were greatly affected by the Mississippi
1428 River during its spring seasonal flooding due to their similar elevations (Figure 3). During
1429 periods of direct connectivity to the Mississippi River, lowland wetlands experienced high
1430 magnitude long-lasting flooding and most likely had lotic conditions during flooding. Upland

1431 wetlands experienced flooding but were never connected to the Mississippi River and, therefore,
1432 experienced lower magnitude and shorter duration flooding and were likely more lentic. Even
1433 though conditions in upland wetlands may lead to distinct communities, it was most likely high
1434 magnitude long-lasting flooding present in lowland wetlands drove the community differences
1435 we observed.

1436 Bighead carp receive spawning cues from increased discharge typically associated with
1437 spring seasonal flooding (Hintz et al. 2017; but see Coulter et al. 2013). This association likely
1438 led to bighead carp's 50.2% contribution to the dissimilarity between lowland and upland
1439 wetlands ($p = 0.048$, Table 2c). It is unlikely that the bighead carp larvae found were the result of
1440 spawning that took place in lowland wetlands, as their eggs require flowing water to develop and
1441 may drift downstream over 100 km before hatching (George & Chapmann 2013; George et al.
1442 2017). If spawning directly occurred in lowland wetland sites, eggs and larvae would surely drift
1443 much further downstream. Therefore, spawning likely occurred upstream of lowland wetlands,
1444 eggs drifted downstream, and eventually hatched in our wetlands. Once hatched, however,
1445 lowland wetlands likely provided larvae with suitable nursery habitat. Varble et al. (2007) found
1446 that floodplain environments are commonly utilized by bighead carp larvae as floodplains are
1447 productive environments that offer abundant food and warm temperature which result in fast
1448 growth rates. The pervasiveness of bighead carp throughout the Mississippi River is well
1449 documented (Chick & Pegg 2001; Pongruktham et al. 2010; Sass et al. 2010) and lowland
1450 wetlands may inherently be at greater risk of bighead carp invasion solely due to their high levels
1451 of connectivity to the river.

1452 Crappie, which significantly contributed to the dissimilarity between lowland and upland
1453 wetlands ($p = 0.013$, Table 2c), are typically associated with lentic conditions, but populations

1454 are common in large river systems (Etnier & Starnes 1993). Despite many studies describing a
1455 strong dependence of many fishes on increased depth and lateral connectivity during seasonal
1456 flooding (Welcomme 1985; Junk et al. 1989; Zeug et al. 2005; Dembkowski & Miranda 2012),
1457 other studies have found that crappie are at best weakly correlated with increased depth and
1458 connectivity and instead are strongly correlated with shallower disconnected floodplain lakes
1459 (Miyazono et al. 2010; Alfermann & Miranda 2013). Our results suggest that crappie may be
1460 positively associated with the high connectivity present in lowland wetlands (Figure 3, Table 2c),
1461 which was most likely due to riverine populations utilizing floodplain habitat to spawn. Riverine
1462 crappie populations are commonly known to utilize the floodplain during spawning because they
1463 are demersal spawners, i.e., they require structure such as submergent aquatic or flooded
1464 terrestrial vegetation that the main channel does not afford (Phelps et al. 2009; Miranda et al.
1465 2015). Recruitment of age-zero crappie has been found to suffer with deeper depths (Dagel &
1466 Miranda 2012). Even if spawning was successful in lowland wetlands, exceptional depths during
1467 seasonal flooding may have imposed negative implications on larval crappie recruitment.

1468 Lowland wetlands experienced prolonged low water and eventually dried during the
1469 summer and fall (Figure 6) because they only flooded from the Mississippi River's seasonal
1470 flood pulse (winter and spring) and water levels were little affected by local precipitation events
1471 throughout the rest of the year (USGS Watershed Science School 2019; Berkowitz et al. 2020).
1472 Drying has obvious negative consequences on fishes such as physical stress, predation, and
1473 mortality. Additionally, Dembkowski & Miranda (2012) found that shallow depth is associated
1474 with harsh environmental conditions, i.e., low dissolved oxygen, high temperatures, which may
1475 cause depauperate fish assemblage composition and prevent larval fish recruitment (Beesley et

1476 al. 2012). During the summer of 2020, lowland wetlands experienced drying before they could
1477 hydrologically reconnect with the river causing local extinctions and prevented recruitment.

1478 Lotic conditions serve as spawning cues for many fishes and certain species' eggs, or
1479 larvae require flows to drift downstream while developing (Welcomme 1985; Junk et al. 1989;
1480 Kluender et al 2015). It is probable that lotic conditions in lowland wetlands allowed other
1481 riverine species to access suitable spawning habitat on the floodplain. Lack of river fishes in our
1482 samples may have resulted from our sampling strategy or these fish may have emigrated from
1483 our wetlands before sampling occurred.

1484 Similarities found in larval fish community composition among wetland condition may
1485 have resulted from exceptional spring flooding experienced throughout our study area during the
1486 spring of 2020. Community composition of fishes is more similar during floods than during low
1487 water periods because floods promote high levels of lateral connectivity which allows the
1488 exchange of fishes among river and floodplain habitat (Hamilton & Lewis 1990; Thomasz et al.
1489 1997; Miranda 2005). Lack of differences in larval fish community composition among wetland
1490 condition may have also resulted from the identical fish species from similar source pools (e.g.,
1491 Bayou du Chien Creek, Mayfield Creek, and Obion Creek) utilizing our wetlands as spawning
1492 habitat during spring flooding. For example, Centrarchidae are common in Mississippi River
1493 tributaries and are prolific floodplain dispersers during flooding (Alfermann & Miranda 2013).
1494 Furthermore, many of our sites were located very near one another, which most likely led to
1495 similarities in colonizing species.

1496 The lack of differences in larval fish diversity amongst wetland elevation and condition
1497 (Figures 5, 6) were most likely due low taxonomic resolution and small sample sizes resulting in

1498 an underrepresentation of diversity. Sample sizes could have been increased by incorporating bi-
1499 monthly sampling or by sampling multiple spawning seasons over two or more years.

1500 Management implications

1501 Wetland elevation's influence on hydrologic conditions and its consequent influences on
1502 larval fish communities was evident throughout our study. When choosing sites for wetland
1503 restoration, restoration managers must consider meaningful tradeoffs that result from wetland
1504 elevation. The elevation of a restored wetland plays a large role in determining local wetland
1505 hydrology which is likely to in turn affect fish communities. Lowland elevation wetlands may be
1506 more strongly influenced by the Mississippi River, experience lotic conditions during seasonal
1507 floods, and dry more often. These conditions can promote spawning habitat for riverine species
1508 and important sport fish, i.e., crappie, but may also promote species invasions or prevent fish
1509 recruitment back into the Mississippi River. If managers are less concerned with providing
1510 benefits for fish, drying and subsequent large-scale die-offs in lowland wetlands may be
1511 beneficial to other wildlife or vegetation communities (Gawlik 2002; Benbow et al. 2020).
1512 Restoring wetlands at higher elevations may minimize the influence of the Mississippi River on
1513 wetland hydrology, which may promote spawning habitat for lentic fishes, nest, or demersal
1514 spawners. Lack of drying can help prevent large-scale die offs of larval fish and allow them to
1515 successfully recruit back into their respective populations during periods of higher hydrologic
1516 connectivity.

1517 Limitations

1518 Low abundances of larval fish were collected and may have been due to only employing
1519 dipnet surveys. Greater larval fish abundances could be achieved in the future by employing
1520 multiple sampling gears (e.g., dipnet and light trap surveys). Similar studies in the future may

1521 consider collecting data over multiple years to allow inter-year comparisons and provide greater
1522 inference into larval fish community patterns throughout the study area. Additionally, future
1523 studies may consider increasing the number of wetlands sampled and employing a balanced data
1524 set, i.e., equal number of wetlands based on elevation or condition categories.

1525 Conclusions

1526 The clear influence of elevation on larval fish communities found in this study may have
1527 implications for wetland restorations throughout the MAV and other large river floodplain
1528 ecosystems. Knowledge gaps associated with the consideration of landscape-level factors exist
1529 within large wetland restoration programs and therefore, quantifying the influence of wetland
1530 elevation on larval fish communities may provide wetland restoration managers with insight and
1531 direction when choosing sites for future restoration. Furthermore, wetland restoration managers
1532 may want to focus on wetland environmental conditions if wildlife response is a goal as our
1533 study found that hydrologic conditions associated with lowland wetlands may have assisted in
1534 promoting unique larval fish community composition. Even though wetland restoration did not
1535 lead to clear community differences when compared with degraded or reference wetlands, our
1536 study provides wetland restoration managers with important criteria to consider when wildlife
1537 response is a goal of restoration. Certainly, restoration practitioners will face and must consider
1538 tradeoffs associated with wetland restoration practices and, hopefully, our results may better
1539 inform future recommendations and restoration projects. Regardless, the need for future studies
1540 that span across multiple temporal and spatial scales to better understand how wetland
1541 restoration practices can influence the entire MAV regions still exists.

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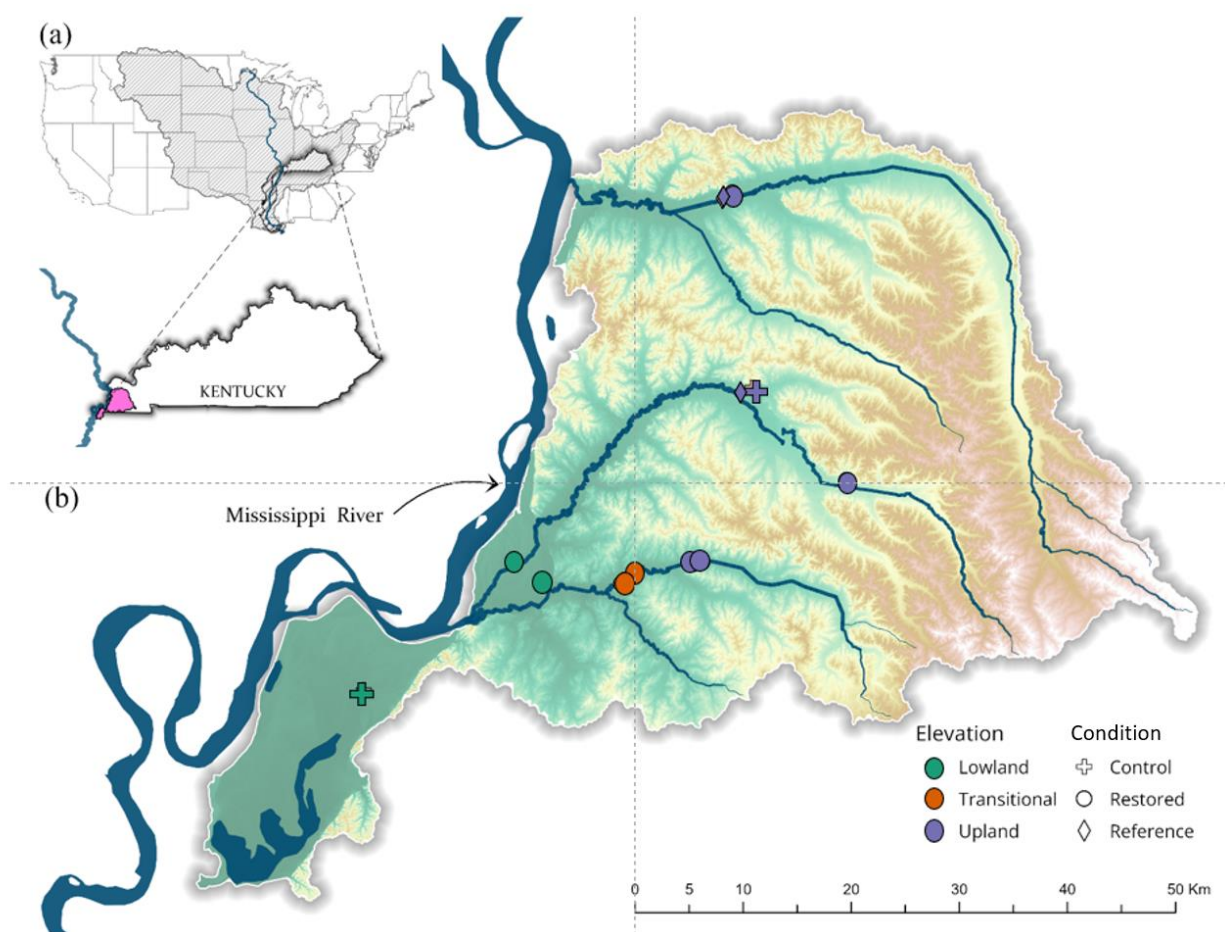
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Figures and tables



1719 **Figure 1.** (A) Location of study area in western Kentucky and part of the Mississippi River watershed, USA. (B)
 1720 Twelve study wetlands were classified by wetland elevation (lowland (n=3), transitional (n=2), upland (n=7))
 1721 denoted by symbol color and wetland condition denoted by shape (degraded (n=2), restored (n=8), reference (n=2))
 1722 (land use classifications from 2018 USDA National Imagery Program).
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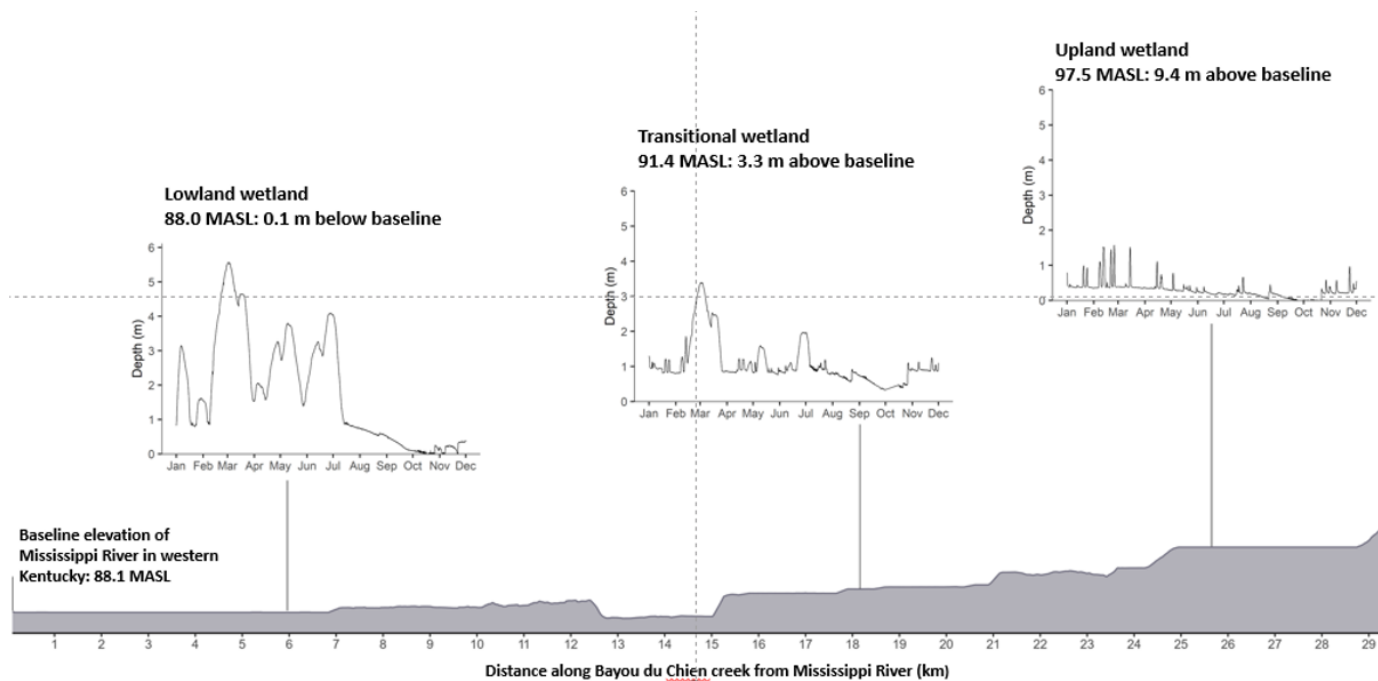


Figure 2: Location of a lowland (left, elevation < 91 MASL), transitional (middle, elevation > 91 but < 97 MASL), and upland (right, elevation > 97 MASL) along the elevation gradient of a tributary (Bayou du Chien) to the Mississippi River, USA. Distance from the Mississippi River (km) is measured along the tributary. One year of mean daily surface water depth (m) readings (taken during 2020) are pictured above each wetland location.

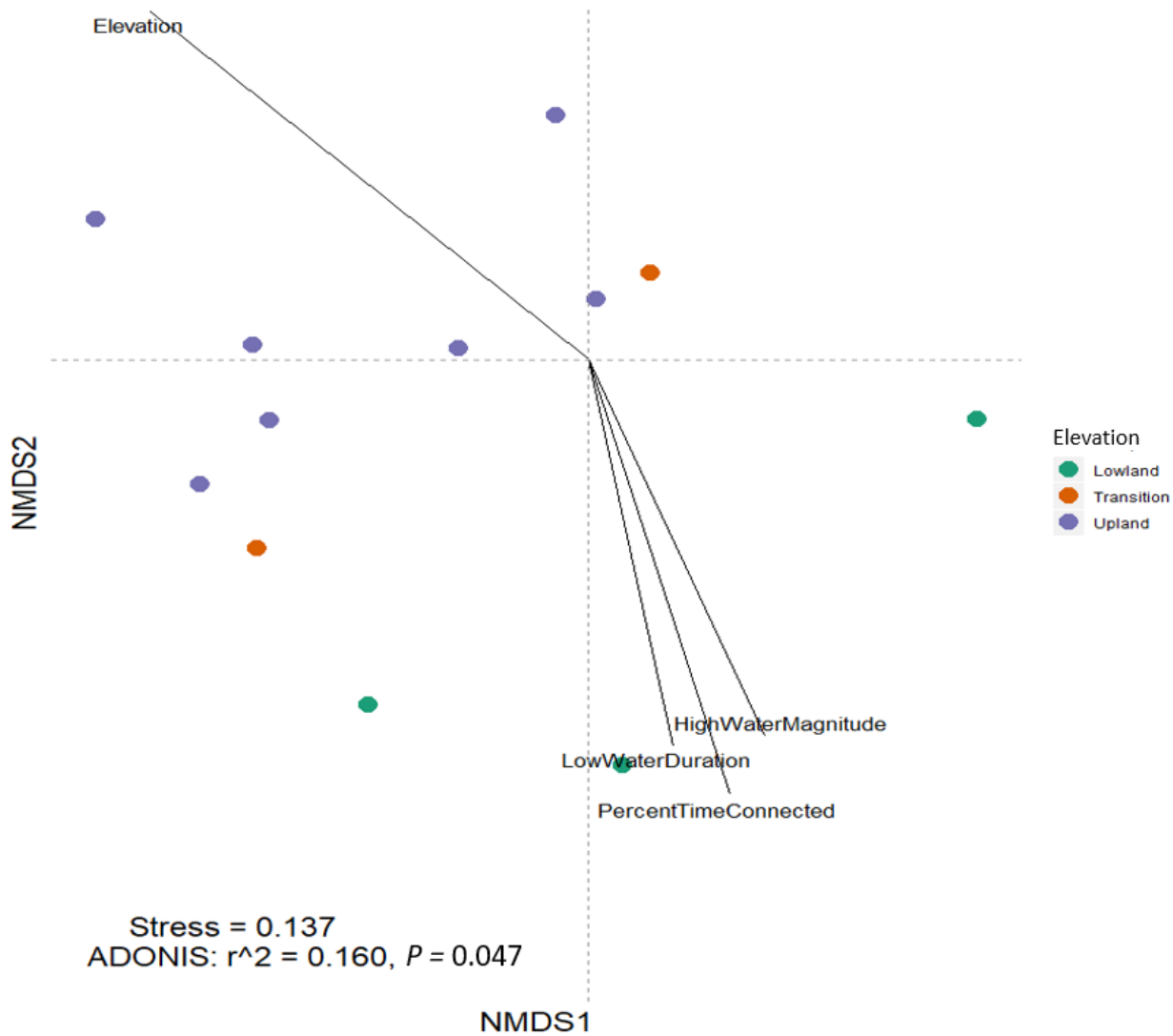
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1760 **Table 1:** Environmental metrics (26) considered for vector analysis in NMDS ordinations measured in wetlands
 1761 throughout western Kentucky, USA. Each metric was calculated using data from the entire sampling period for each
 1762 wetland in our study. Indicators of hydrologic alteration following Richter et al. (1996) are denoted with the
 1763 abbreviation 'IHA'.
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Metric	Definition	Measures
<u>Hydrologic metrics</u>		
Mean Depth (m)	The average wetland depth.	Magnitude
Hydroperiod	The number of days a wetland had water.	Duration
Percent Time Connected	The percent of time a wetland exhibited lateral connectivity to its nearest stream (USGS-NWIS).	Connectivity
1-Day Maximum (m) (IHA)	The maximum single day depth recorded.	Magnitude/Duration
1-Day Minimum (m) (IHA)	The minimum single day depth recorded.	Magnitude/Duration
7-Day Maximum (m) (IHA)	The maximum 7-day rolling average recorded.	Magnitude/Duration
7-Day Minimum (m) (IHA)	The minimum 7-day rolling average recorded.	Magnitude/Duration
30-Day Maximum (m) (IHA)	The maximum 30-day rolling average recorded.	Magnitude/Duration
30-Day Minimum (m) (IHA)	The minimum 30-day rolling average recorded.	Magnitude/Duration
90-Day Maximum (m) (IHA)	The maximum 90-day rolling average recorded.	Magnitude/Duration
90-Day Minimum (m) (IHA)	The minimum 90-day rolling average recorded.	Magnitude/Duration
Minimum Date (day of the year) (IHA)	The date the lowest single recorded depth occurred.	Timing
Maximum Date (day of the year) (IHA)	The date the greatest single recorded depth occurred.	Timing
Low Water Duration (days) (IHA)	The number of consecutive days depths stayed below the 25th percentile.	Duration
High Water Duration (days) (IHA)	The number of consecutive days depths stayed above the 75th percentile.	Duration
Rise Count (IHA)	The number of occurrences that wetland depth rose from the previous day.	Connectivity
<u>Water quality metrics</u>		
Mean Dissolved Oxygen (DO) (mg/L)	The average DO measurement.	Magnitude
Minimum Dissolved Oxygen (DO) (mg/L)	The minimum single day DO recorded.	Magnitude
Mean Temperature (C°)	The average temperature recorded.	Magnitude

1-Day Maximum (C°)	The maximum single day temperature recorded.	Magnitude
1-Day Minimum (C°)	The minimum single day temperature recorded.	Magnitude
<u>Biotic metrics</u>		
Vegetation	The average aquatic vegetation scores.	Habitat
Zooplankton Density	Average of each wetland's zooplankton density	Food resources
KY-WRAM Score	The KY-WRAM score indication ecological integrity (0-100) (100 indicates highest quality).	Disturbance
<u>Hydrologic connectivity</u>		
Distance to Main Channel (m)	The distance to the main channel of the nearest stream.	Connectivity
Elevation (m)	The median of elevation (n=50) (USGS Stream Stats).	Connectivity

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Figure 3: NMDS ordination of larval fish community composition in western Kentucky, USA wetlands. Ordination is based on per taxa CPUE from dipnet surveys that occurred monthly from March 2020 to August 2020. Symbol colors indicate wetland elevation (lowland, transitional, upland). All variables from Table 1 were tested and only significant variables were placed onto ordination as vectors. The vector High Water Magnitude is a combination of the metrics 1-Day Maximum (m), 30-Day Maximum (m), 90-Day Maximum (m). See table 1 for definitions of other variables used as vectors.

1790 **Table 2:** Summary of SIMPER results for each fish genera from pairwise comparisons amongst (a) lowland vs
 1791 transitional wetlands, (b) upland vs transitional wetlands, and (c) lowland vs upland wetlands: average abundance of
 1792 individual species from each wetland, their average contribution (%) to overall dissimilarity, and each species
 1793 associated p-value. P-values were considered significant at the $\alpha = 0.05$ level.
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(a) Lowland vs transitional	Lowland average abundance	Transitional average abundance	Contribution	P-value
<i>Hypophthalmichthys</i>	109.7	0.67	54.2	0.068
<i>Lepomis</i>	1.08	3.33	9.3	0.936
<i>Elassoma</i>	0.11	1.75	6.9	0.222
<i>Notemigonus</i>	0.67	0.42	2.1	0.233
<i>Micropterus</i>	0.42	0.00	1.5	0.775
<i>Pomoxis</i>	0.14	0.00	0.9	0.060
<i>Fundulus</i>	0.00	0.17	0.8	0.449
<i>Esox</i>	0.00	0.17	0.8	0.348
<i>Aphredoderus</i>	0.00	0.08	0.4	0.798
<i>Dorosoma</i>	0.08	0.00	0.3	0.262
<i>Ictiobus</i>	0.25	0.00	0.1	0.288
<i>Lepisosteus</i>	0.17	0.00	0.1	0.288
<i>Umbra</i>	0.00	0.00	0.0	1.00
<i>Ameiurus</i>	0.00	0.00	0.0	1.00
<i>Erimyzon</i>	0.00	0.00	0.0	1.00

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(b) Upland vs transitional	Upland average abundance	Transitional average abundance	Contribution	P-value
<i>Lepomis</i>	11.7	3.33	28.1	0.476
<i>Elassoma</i>	0.48	1.75	8.5	0.081
<i>Hypophthalmichthys</i>	0.00	0.67	3.8	0.958
<i>Notemigonus</i>	0.21	0.42	2.4	0.107
<i>Micropterus</i>	0.55	0.0	2.0	0.650
<i>Aphredoderus</i>	0.29	0.08	1.6	0.419
<i>Ameiurus</i>	0.59	0.0	1.5	0.410
<i>Fundulus</i>	0.95	0.17	1.2	0.202
<i>Esox</i>	0.17	0.17	1.0	0.099
<i>Erimyzon</i>	0.07	0.0	0.4	0.356
<i>Umbra</i>	0.24	0.0	0.2	0.379
<i>Pomoxis</i>	0.00	0.0	0.0	1.00
<i>Dorosoma</i>	0.00	0.0	0.0	1.00
<i>Lepisosteus</i>	0.00	0.0	0.0	1.00
<i>Ictiobus</i>	0.00	0.0	0.0	1.00

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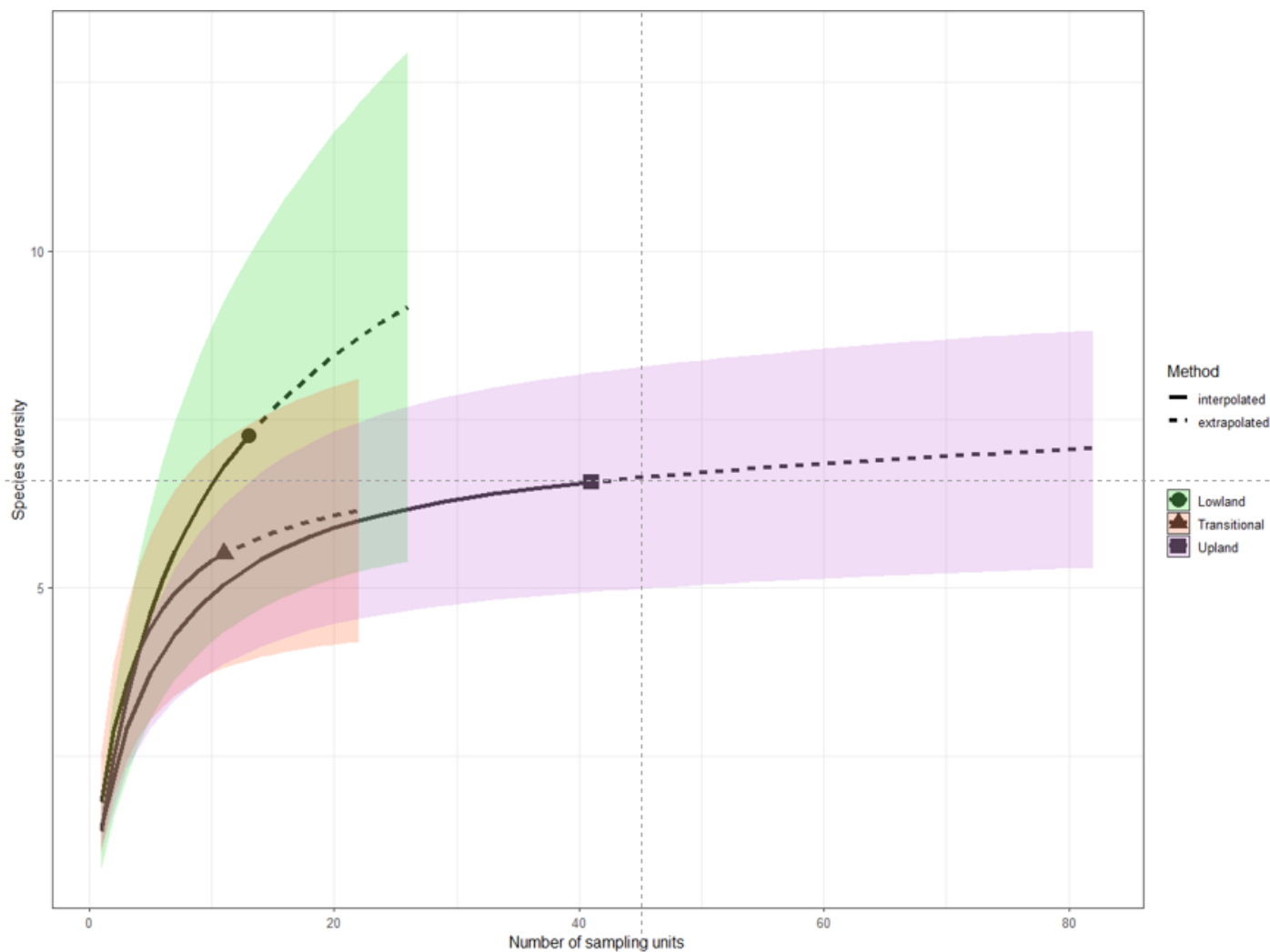
(c) Upland vs lowland	Upland average abundance	Lowland average abundance	Contribution	P-value
<i>Hypophthalmichthys</i>	0.00	109.7	50.2	0.048*
<i>Lepomis</i>	11.7	1.08	25.8	0.602
<i>Micropterus</i>	0.55	0.42	1.98	0.642
<i>Elassoma</i>	0.47	0.11	1.48	0.943
<i>Aphredoderus</i>	0.29	0.00	1.29	0.630
<i>Ameiurus</i>	0.56	0.00	1.01	0.759
<i>Notemigonus</i>	0.21	0.67	1.00	0.911
<i>Pomoxis</i>	0.00	0.14	0.80	0.013*
<i>Esox</i>	0.17	0.00	0.04	0.974
<i>Fundulus</i>	0.10	0.00	0.04	0.849
<i>Erimyzon</i>	0.07	0.00	0.04	0.561
<i>Dorosoma</i>	0.00	0.08	0.03	0.091
<i>Umbra</i>	0.24	0.00	0.02	0.552
<i>Ictiobus</i>	0.00	0.25	0.01	0.212
<i>Lepisosteus</i>	0.00	0.17	0.01	0.212

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1798 **Table 3:** Correlation coefficients and p-values for vectors placed onto NMDS ordination that had significant
 1799 associations with larval fish community composition.
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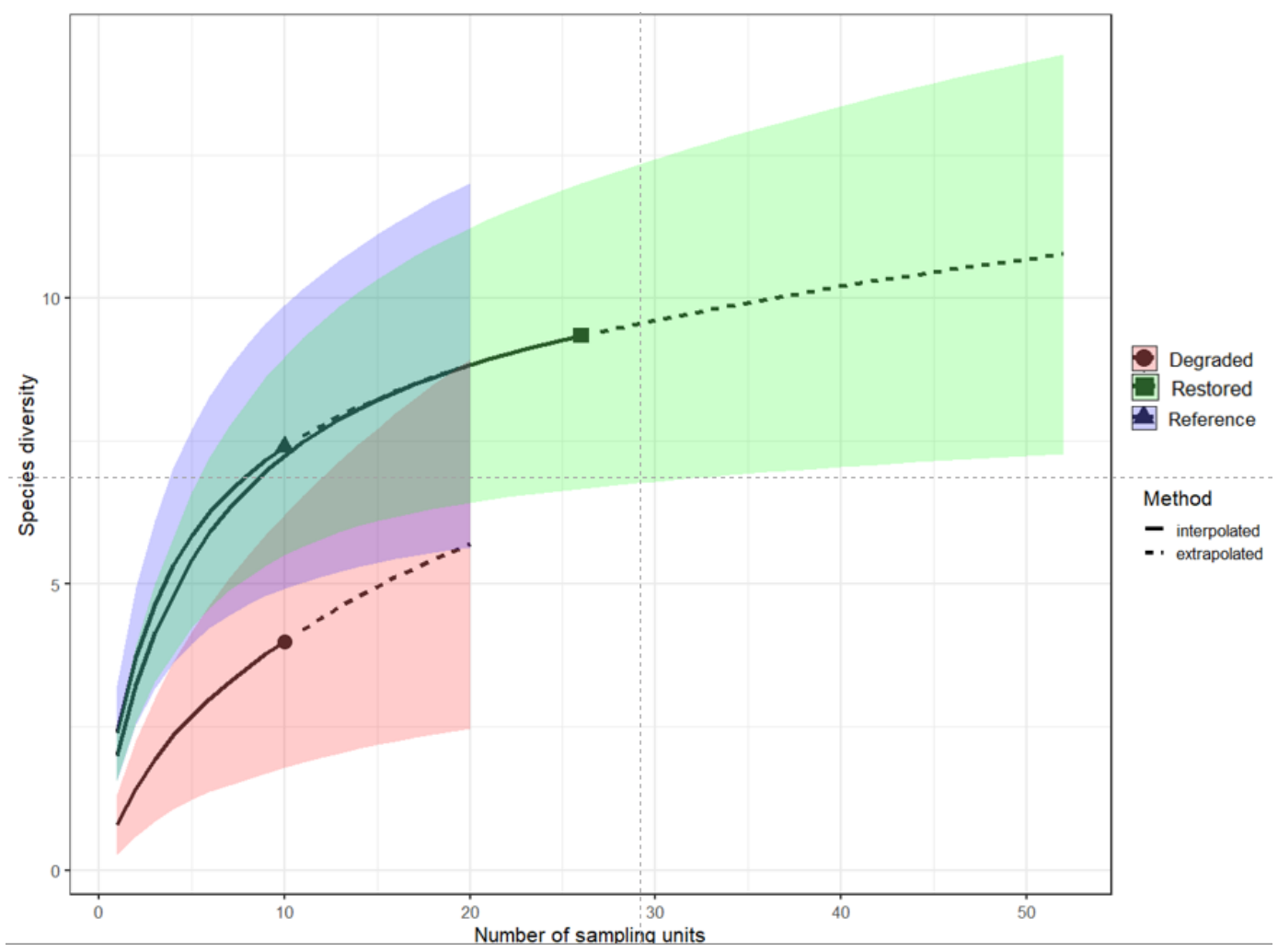
Vector	Correlation Coefficient (r^2)	P-Value
Percent time connected	0.736	0.001
Elevation	0.777	0.002
High water magnitude	0.669	0.009
Low water duration	0.529	0.032

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Figure 4: Hill-Shannon diversity estimates of wetland larval fish communities by wetland elevation using incidence-based rarefaction and extrapolation. Curves are based on larval fish dipnet survey data collected from March 2020 through August 2020 in western Kentucky, USA. All extrapolation curves were plotted to achieve 90% coverage.



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Figure 5: Hill-Shannon diversity estimates of wetland larval fish communities by wetland condition using incidence-based rarefaction and extrapolation. Curves are based on larval fish dipnet survey data collected from March 2020 through August 2020 in western Kentucky, USA. All extrapolation curves were plotted to achieve 90% coverage.



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Figure 6: Photograph of dramatic low water/drying in a lowland wetland in western Kentucky, USA. Photograph was taken in August 2020.

Appendix

Supplemental tables

Table 1: Site name, wetland condition, wetland elevation, county, easement acreage, and wetland acreage of wetland sampled in western Kentucky, USA.

Site	Wetland condition	Wetland elevation	County	Acreage	Wetland acreage
ALEN	Restored	Upland	Hickman	68	49
BCYP	Degraded	Lowland	Fulton	NA	2
COFY	Restored	Lowland	Fulton	251	34
GDMN	Restored	Lowland	Fulton	115	24
GUTH	Restored	Upland	Graves	141	2
HEST	Restored	Transitional	Hickman	39	20
HOPK	Restored	Upland	Hickman	44	4
HWST	Restored	Transitional	Hickman	35	19
OBOT	Reference	Upland	Carlisle	NA	72
OWMA	Degraded	Upland	Carlisle	NA	5
SARC	Reference	Upland	Carlisle	NA	2
SWAN	Restored	Upland	Carlisle	784	9

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Table 2: Presence/absence of every taxon collected at each wetland. White rows represent species-level presence/absence data of adult fish community collected with backpack electrofishing. Green rows represent genus-level presence/absence data of larval fish community collected with dipnet surveys. Orange row represents family-level presence/absence data (due to difficulty in identification) of larval fish community collected with dipnet surveys. ‘X’ indicates species was present at a wetland. ‘*’ indicates larval genus was present at a wetland. ‘***’ indicates special note: identification of *Hybognathus hayi* is pending on verification from Kentucky’s state ichthyologist; the *Lepomis marginatus* we collected within the Bayou du Chien watershed were the first species records collected within that watershed (verified by Kentucky’s state ichthyologist).

Genus/Species	ALEN	BCYP	COFY	GDMN	GUTH	HEST	HOPK	HWST	OBOT	OWMA	SARC	SWAN
<i>Ameiurus</i>												*
<i>Ameiurus melas</i>	X	X		X		X	X	X		X	X	X
<i>Ameiurus nebulosus</i>		X				X		X				
<i>Ameiurus natalis</i>	X			X	X	X	X	X		X		X
<i>Amia</i>												
<i>Amia calva</i>	X	X				X	X	X	X	X	X	X
<i>Aphredoderus</i>					*	*			*		*	*
<i>Aphredoderus sayanus</i>	X	X	X	X		X	X	X	X	X	X	X
<i>Aplodinotus</i>												
<i>Aplodinotus grunniens</i>				X								
<i>Centrarchus</i>												
<i>Centrarchus macropterus</i>	X	X		X		X	X	X	X	X	X	X
<i>Ctenopharyngodon</i>												
<i>Ctenopharyngodon idella</i>			X	X								
<i>Cycleptus</i>												
<i>Cycleptus elongatus</i>				X								
<i>Cyprinella</i>												
<i>Cyprinella lutrensis</i>											X	
<i>Cyprinella venusta</i>						X						
<i>Cyprinidae</i>			*	*		*					*	
<i>Cyprinus</i>												
<i>Cyprinus carpio</i>			X	X		X	X					X

Genus/Species	ALEN	BCYP	COFY	GDMN	GUTH	HEST	HOPK	HWST	OBOT	OWMA	SARC	SWAN
Dorosoma			*									
<i>Dorosoma cepedianum</i>			X	X		X	X	X				X
Elassoma	*	*				*	*	*	*		*	*
<i>Elassoma zonatum</i>	X	X			X	X	X	X	X	X	X	X
Erimyzon									*			*
<i>Erimyzon sucetta</i>	X					X	X	X	X	X		X
Esox							*	*		*	*	*
<i>Esox americanus</i>	X	X				X	X	X	X	X	X	X
Etheostoma												
<i>Etheostoma asprigene</i>						X					X	
<i>Etheostoma chlorosomum</i>			X			X					X	
<i>Etheostoma gracile</i>					X	X	X	X	X	X	X	X
Fundulus						*					*	
<i>Fundulus chrysotus</i>		X										
<i>Fundulus olivaceus</i>			X			X		X	X		X	X
Gambusia												
<i>Gambusia affinis</i>	X	X	X	X	X	X	X	X	X	X	X	X
Hybognathus												
<i>Hybognathus hayi</i> ***											X	X
Hypophthalmichthys			*	*		*						
<i>Hypophthalmichthys molitrix</i>						X						
Ictalurus												
<i>Ictalurus punctatus</i>				X								
Ictiobus				*								
<i>Ictiobus bubalus</i>			X	X		X	X	X		X	X	X
Labidesthes												
<i>Labidesthes sicculus</i>	X					X					X	

Genus/Species	ALEN	BCYP	COFY	GDMN	GUTH	HEST	HOPK	HWST	OBOT	OWMA	SARC	SWAN
<i>Lepisosteus</i>				*								
<i>Lepisosteus oculatus</i>	X					X		X	X		X	
<i>Lepisosteus osseus</i>			X	X		X					X	
<i>Lepomis</i>	*	*	*	*	*	*	*	*	*	*	*	*
<i>Lepomis cyanellus</i>	X	X	X	X	X	X	X	X	X	X	X	X
<i>Lepomis humilis</i>			X	X			X				X	
<i>Lepomis macrochirus</i>	X	X	X	X		X	X	X	X		X	X
<i>Lepomis marginatus</i>	X		X	X		X	X	X	X	X		
<i>Lepomis megalotis</i>			X	X		X					X	
<i>Lepomis microlophus</i>	X			X							X	
<i>Lepomis miniatus</i>	X										X	X
<i>Lepomis symmetricus</i>	X	X		X		X	X	X	X	X	X	X
<i>Lepomis gulosus</i>	X	X	X	X	X	X	X	X	X	X	X	X
<i>Menidia</i>												
<i>Menidia beryllina</i>				X								
<i>Micropterus</i>	*		*				*				*	*
<i>Micropterus salmoides</i>	X	X	X	X		X	X	X	X	X	X	X
<i>Minytrema</i>												
<i>Minytrema melanops</i>	X					X			X		X	
<i>Morone</i>												
<i>Morone mississippiensis</i>				X								
<i>Notemigonus</i>						*			*	*	*	*
<i>Notemigonus crysoleucas</i>	X	X	X	X		X	X	X	X	X	X	X
<i>Notropis</i>												
<i>Notropis maculatus</i>											X	
<i>Noturus</i>												
<i>Noturus gyrinus</i>				X					X			

Genus/Species	ALEN	BCYP	COFY	GDMN	GUTH	HEST	HOPK	HWST	OBOT	OWMA	SARC	SWAN
<i>Opsopoeodus</i>												
<i>Opsopoeodus emiliae</i>											X	
<i>Percina</i>												
<i>Percina caprodes</i>			X									
<i>Pimephales</i>												
<i>Pimephales vigilax</i>											X	
<i>Pomoxis</i>												
<i>Pomoxis annularis</i>			X			X	X	X			X	X
<i>Pomoxis nigromaculatus</i>	X	X	X	X		X	X	X			X	X
<i>Sander</i>												
<i>Sander canadensis</i>			X									
<i>Semotilus</i>												
<i>Semotilus atromaculatus</i>						X					X	
<i>Umbra</i>												
<i>Umbra limi</i>	X	X			X	X	X	X	X	X	X	

ALEN												
Species	2019-05-29	2019-06-25	2019-07-24	2019-08-27	2019-10-24	2020-02-01	2020-03-01	2020-04-01	2020-05-01	2020-06-02	2020-07-06	2020-08-05
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	0.12	0.17	0.08	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	—	0.09	—	—	—	0.09	—

BCYP

Species	2019															
	-04-29	2019-05-31	2019-06-25	2019-07-24	2019-09-06	2019-10-03	2019-11-06	2019-12-13	2020-01-08	2020-02-06	2020-03-13	2020-03-31	2020-05-11	2020-06-10	2020-07-14	2020-08-18
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	—	—	0.11	0.30	0.44	0.30	0.10	—	0.17	0.09	—	—	—	—	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	0.24	—	0.36	0.55	0.20	—	—	1.60	0.60	0.17	—	0.17	0.08	0.34	1.98	2.40
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	—	—	0.20	0.09	—	—	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	0.11	—	—	—	—	0.09	—	—	0.09	—	—

1864 **Table 3c:**

Species	COFY						
	2019-08-02	2019-09-03	2019-12-13	2020-05-18	2020-06-16	2020-07-14	2020-08-18
<i>Ameiurus melas</i>	—	—	—	—	—	—	—
<i>Ameiurus natalis</i>	—	—	—	—	—	—	—
<i>Ameiurus nebulosus</i>	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	—	—	—	—	—
<i>Aphredoderus sayanus</i>	0.34	0.10	—	0.24	0.08	—	—
<i>Aplodinotus grunniens</i>	—	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	—	—	—	—	—	—
<i>Ctenopharyngodon idella</i>	—	—	—	—	—	—	0.11
<i>Cycleptus elongatus</i>	—	—	—	—	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—
<i>Cyprinus carpio</i>	—	—	—	—	—	—	0.22
<i>Dorosoma cepedianum</i>	0.60	—	—	0.24	—	—	0.33
<i>Elassoma zonatum</i>	—	—	—	—	—	—	—
<i>Erimyzon sucetta</i>	—	—	—	—	—	—	—
<i>Esox americanus</i>	—	—	—	—	—	—	—
<i>Etheostoma asprigene</i>	—	—	—	—	—	—	—
<i>Etheostoma chlorosoma</i>	0.43	0.10	—	—	—	—	—
<i>Etheostoma gracile</i>	—	—	—	—	—	—	—
<i>Fundulus chrysotus</i>	—	—	—	—	—	—	—
<i>Fundulus olivaceus</i>	0.09	—	0.20	—	—	0.08	1.20
<i>Gambusia affinis</i>	—	1.00	0.20	—	—	—	—
<i>Hybognathus hayi</i>	—	—	—	—	—	—	—
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	—	—	—	—	—	—
<i>Ictiobus bubalus</i>	0.17	0.20	0.70	—	—	—	—

COFY

Species	2019-08-02	2019-09-03	2019-12-13	2020-05-18	2020-06-16	2020-07-14	2020-08-18
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—
<i>Lepisosteus oculatus</i>	—	—	—	—	—	—	—
<i>Lepomis cyanellus</i>	0.09	—	—	—	—	—	—
<i>Lepomis gulosus</i>	0.17	—	—	0.08	0.08	—	—
<i>Lepomis humilis</i>	0.09	—	—	0.08	—	—	—
<i>Lepomis macrochirus</i>	0.09	0.30	0.30	0.40	0.08	0.32	0.33
<i>Lepomis marginatus</i>	—	—	0.20	—	—	—	—
<i>Lepomis megalotis</i>	0.09	0.50	—	0.08	0.08	—	0.22
<i>Lepomis microlophus</i>	—	—	—	—	—	—	—
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	—	—	—	—	—	—	—
<i>Menidia beryllina</i>	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	—	—	0.08	0.32	0.64	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	—	—	—	—	0.08	0.88	0.11
<i>Notropis maculatus</i>	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—
<i>Percina caprodes</i>	0.09	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	0.17	0.10	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	0.26	1.00	0.60	0.08	0.56	0.16	0.11
<i>Sander canadensis</i>	—	0.10	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	—	—

1866 **Table 3d:**

Species	GDMN						
	2019-07-18	2019-09-03	2019-09-24	2020-05-14	2020-06-16	2020-07-14	2020-08-18
<i>Ameiurus melas</i>	—	—	—	—	—	4.80	1.85
<i>Ameiurus natalis</i>	—	0.60	—	0.09	0.09	—	—
<i>Ameiurus nebulosus</i>	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	—	—	—	—	—
<i>Aphredoderus sayanus</i>	—	—	—	0.17	0.17	—	—
<i>Aplodinotus grunniens</i>	0.10	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	0.40	—	—	0.34	—	—
<i>Ctenopharyngodon idella</i>	—	—	—	—	0.09	—	—
<i>Cycleptus elongatus</i>	—	—	—	0.09	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—
<i>Cyprinus carpio</i>	—	—	0.15	—	1.20	1.37	0.87
<i>Dorosoma cepedianum</i>	0.20	—	—	—	—	—	—
<i>Elassoma zonatum</i>	—	—	—	—	—	—	—
<i>Erimyzon sucetta</i>	—	—	—	—	—	—	—
<i>Esox americanus</i>	—	—	—	—	—	—	—
<i>Etheostoma asprigene</i>	—	—	—	—	—	—	—
<i>Etheostoma chlorosoma</i>	—	—	—	—	—	—	—
<i>Etheostoma gracile</i>	—	—	—	—	—	—	—
<i>Fundulus chrysotus</i>	—	—	—	—	—	—	—
<i>Fundulus olivaceus</i>	—	—	—	—	—	—	—
<i>Gambusia affinis</i>	—	—	1.05	—	—	1.89	3.93
<i>Hybognathus hayi</i>	—	—	—	—	—	—	—
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	0.10	0.75	—	—	—	—
<i>Ictiobus bubalus</i>	—	—	10.20	—	1.11	—	—

GDMN

Species	2019-07-18	2019-09-03	2019-09-24	2020-05-14	2020-06-16	2020-07-14	2020-08-18
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—
<i>Lepisosteus oculatus</i>	—	—	—	—	—	—	—
<i>Lepomis cyanellus</i>	—	0.60	0.60	0.09	1.20	0.86	—
<i>Lepomis gulosus</i>	—	0.20	—	0.34	—	—	0.55
<i>Lepomis humilis</i>	—	—	—	0.09	—	0.09	—
<i>Lepomis macrochirus</i>	—	0.40	0.30	—	0.17	0.86	0.76
<i>Lepomis marginatus</i>	—	—	—	0.26	—	—	—
<i>Lepomis megalotis</i>	0.10	1.10	0.45	0.43	0.09	0.60	0.33
<i>Lepomis microlophus</i>	—	—	0.15	—	—	—	—
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	—	—	0.15	—	—	—	—
<i>Menidia beryllina</i>	—	—	—	—	—	—	0.11
<i>Micropterus salmoides</i>	—	—	—	—	0.26	0.17	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	0.10	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	—	—	—	—	0.17	—	—
<i>Notropis maculatus</i>	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	0.10	0.15	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	0.50	—	—	—	0.69	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	—	—

GUTH

Species	2019-04-09	2019-05-08	2019-06-03	2019-07-01	2019-08-01	2019-09-05	2019-10-08	2019-11-05	2019-12-12	2020-01-08	2020-02-07	2020-03-04	2020-04-01	2020-05-07	2020-06-11	2020-07-09	2020-08-04
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	0.12	—	—	—	—	—	—	—	—	—	—	—	—

1870 Table 3f:

Species	HEST												
	2019-05-31	2019-07-18	2019-08-28	2019-09-24	2019-12-11	2020-01-06	2020-02-03	2020-03-06	2020-04-15	2020-05-11	2020-06-10	2020-07-15	2020-08-18
<i>Ameiurus melas</i>	—	0.30	0.80	52.55	—	0.10	0.32	0.09	—	0.28	0.27	0.32	—
<i>Ameiurus natalis</i>	—	—	1.50	28.83	—	—	—	—	—	—	—	0.24	0.87
<i>Ameiurus nebulosus</i>	—	—	0.30	4.20	—	—	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	0.10	—	—	0.10	—	0.09	0.16	—	—	—	—
<i>Aphredoderus sayanus</i>	0.48	1.10	0.10	0.60	0.36	0.20	0.56	0.26	—	—	1.47	—	0.22
<i>Aplodinotus grunniens</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	0.20	0.30	0.30	0.24	0.10	0.24	—	0.08	—	1.60	0.32	0.44
<i>Ctenopharyngodon idella</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cycleptus elongatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—	0.17	—	—	—	—	—
<i>Cyprinus carpio</i>	0.12	0.20	0.80	—	—	—	—	—	—	—	0.67	—	0.76
<i>Dorosoma cepedianum</i>	—	0.50	0.10	—	—	—	—	0.09	—	—	—	—	0.22
<i>Elassoma zonatum</i>	0.96	0.20	—	—	0.12	0.10	0.48	0.17	0.16	0.09	0.27	—	—
<i>Erimyzon sucetta</i>	—	—	—	—	—	—	—	0.09	—	—	2.40	—	0.33
<i>Esox americanus</i>	0.72	—	0.20	—	0.12	0.20	0.08	0.09	0.64	0.55	1.60	—	0.11
<i>Etheostoma asprigene</i>	—	—	—	—	—	—	—	—	—	—	0.13	—	—
<i>Etheostoma chlorosoma</i>	—	—	—	—	—	—	—	0.09	—	—	—	—	—
<i>Etheostoma gracile</i>	0.36	0.10	—	—	—	—	0.64	0.09	—	0.18	0.53	—	0.11

HEST

Species	2019-05-31	2019-07-18	2019-08-28	2019-09-24	2019-12-11	2020-01-06	2020-02-03	2020-03-06	2020-04-15	2020-05-11	2020-06-10	2020-07-15	2020-08-18
<i>Fundulus chrysotus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Fundulus olivaceus</i>	—	—	0.10	—	0.48	0.10	0.08	0.17	—	0.09	—	—	0.44
<i>Gambusia affinis</i>	0.24	0.40	0.10	69.37	0.48	0.10	0.16	0.17	0.40	0.65	15.33	0.16	1.64
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	0.12	—	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictiobus bubalus</i>	4.32	0.10	0.20	0.60	—	—	—	0.77	—	—	—	—	—
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—	—	—	—	—	—	0.33
<i>Lepisosteus oculatus</i>	—	0.10	—	—	—	—	—	—	—	—	—	—	—
<i>Lepisosteus osseus</i>	—	—	—	0.30	—	—	—	—	—	—	—	—	—
<i>Lepomis cyanellus</i>	—	—	—	—	—	—	—	0.17	0.08	0.09	—	0.08	—
<i>Lepomis gulosus</i>	0.72	0.10	0.10	0.90	0.60	0.30	0.08	0.34	0.40	0.46	0.40	—	0.11
<i>Lepomis humilis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis macrochirus</i>	0.36	—	—	—	0.48	0.90	0.16	0.34	0.24	0.28	—	0.24	0.44
<i>Lepomis marginatus</i>	—	—	—	—	—	—	—	0.09	—	—	—	—	—
<i>Lepomis megalotis</i>	—	—	—	—	—	—	—	—	—	—	—	0.08	0.11
<i>Lepomis microlophus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	0.24	0.20	0.10	0.60	—	—	0.16	0.09	0.40	0.18	—	—	0.76

HEST

Species	2019-05-31	2019-07-18	2019-08-28	2019-09-24	2019-12-11	2020-01-06	2020-02-03	2020-03-06	2020-04-15	2020-05-11	2020-06-10	2020-07-15	2020-08-18
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	0.24	—	0.30	—	0.12	0.30	0.16	0.43	—	—	—	—	0.11
<i>Minytrema melanops</i>	—	—	—	—	0.12	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	3.24	1.90	—	—	0.12	0.10	—	0.09	0.08	0.18	4.13	0.16	0.33
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	0.20	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	0.20	0.40	0.60	—	0.10	—	0.09	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	0.24	—
<i>Umbra limi</i>	—	—	—	—	—	—	—	—	—	—	—	—	0.22

1872 Table 3g:

Species	HOPK														
	2019-05-29	2019-06-25	2019-07-24	2019-08-27	2019-09-19	2019-10-24	2019-12-11	2020-01-06	2020-02-01	2020-03-01	2020-04-01	2020-05-01	2020-06-02	2020-07-06	2020-08-05
<i>Ameiurus melas</i>	0.24	—	—	—	—	—	0.20	0.28	0.48	0.17	—	—	0.18	0.17	—
<i>Ameiurus natalis</i>	—	—	—	—	0.12	0.13	—	—	—	—	—	—	—	—	—
<i>Ameiurus nebulosus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	—	—	0.12	—	—	—	—	—	0.08	—	—	—	—
<i>Aphredoderus sayanus</i>	—	—	0.12	—	0.12	—	0.10	0.18	0.08	—	—	—	—	—	—
<i>Aplodinotus grunniens</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	—	—	—	0.24	0.13	0.10	0.18	0.08	0.34	0.08	—	—	—	—
<i>Ctenopharyngodon idella</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cycleptus elongatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinus carpio</i>	—	—	0.12	0.10	—	—	—	—	—	—	—	—	—	—	—
<i>Dorosoma cepedianum</i>	—	—	0.06	—	—	—	—	—	—	—	—	—	—	—	—
<i>Elassoma zonatum</i>	—	—	—	0.10	—	0.13	—	0.09	0.40	1.54	1.28	—	—	—	—
<i>Erimyzon sucetta</i>	—	0.12	—	0.10	0.12	—	0.20	0.18	0.96	0.69	—	—	—	—	—
<i>Esox americanus</i>	—	—	—	—	0.36	0.13	0.20	0.28	0.48	0.26	0.16	0.09	—	—	—
<i>Etheostoma asprigene</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Etheostoma chlorosoma</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Etheostoma gracile</i>	—	—	—	—	—	—	—	—	0.48	0.69	—	—	—	—	—

HOPK

Species	2019-05-29	2019-06-25	2019-07-24	2019-08-27	2019-09-19	2019-10-24	2019-12-11	2020-01-06	2020-02-01	2020-03-01	2020-04-01	2020-05-01	2020-06-02	2020-07-06	2020-08-05
<i>Fundulus chrysotus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Fundulus olivaceus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Gambusia affinis</i>	0.12	0.24	0.24	—	—	0.13	—	0.09	0.08	—	0.16	1.94	0.09	—	—
<i>Hybognathus hayi</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictiobus bubalus</i>	—	—	—	—	0.24	—	—	—	—	—	—	—	—	—	—
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepisosteus oculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepisosteus osseus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis cyanellus</i>	—	0.12	—	—	0.72	—	—	—	—	—	—	0.18	0.09	—	0.10
<i>Lepomis gulosus</i>	0.36	0.72	0.18	0.40	3.48	1.60	0.20	0.28	0.24	0.86	0.88	0.92	0.37	—	0.10
<i>Lepomis humilis</i>	—	—	—	—	—	—	—	0.09	—	—	—	—	—	—	—
<i>Lepomis macrochirus</i>	1.20	1.08	0.18	0.20	1.44	0.67	0.60	0.92	0.72	0.77	0.56	0.37	0.09	0.69	1.15
<i>Lepomis marginatus</i>	0.24	0.72	0.18	0.10	1.20	0.80	—	0.09	0.16	0.43	0.32	0.28	—	—	0.10
<i>Lepomis megalotis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis microlophus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	0.24	1.32	0.12	0.10	1.68	1.20	—	—	0.32	0.34	0.96	0.28	0.09	0.17	0.10

HOPK

Species	2019- 05-29	2019- 06-25	2019- 07-24	2019- 08-27	2019- 09-19	2019- 10-24	2019- 12-11	2020- 01-06	2020- 02-01	2020- 03-01	2020- 04-01	2020- 05-01	2020- 06-02	2020- 07-06	2020- 08-05
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	0.12	—	0.10	—	0.13	0.10	—	—	—	—	—	—	0.34	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	0.24	0.12	0.36	1.60	1.08	0.13	0.10	—	0.16	0.17	—	0.28	0.09	0.51	0.31
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	0.06	0.20	0.12	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	0.06	—	0.48	—	—	—	—	—	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	0.12	—	—	0.12	0.27	—	—	0.16	—	—	—	—	—	—

1874 **Table 3h:**

Species	HWST												
	2019-05-31	2019-07-18	2019-08-28	2019-09-24	2019-10-14	2019-12-11	2020-01-06	2020-02-03	2020-03-06	2020-04-15	2020-05-11	2020-06-10	2020-07-15
<i>Ameiurus melas</i>	—	—	—	—	—	0.26	0.26	0.17	0.40	0.16	0.08	—	—
<i>Ameiurus natalis</i>	—	0.24	—	—	—	—	0.17	—	—	—	—	—	—
<i>Ameiurus nebulosus</i>	—	—	—	0.08	—	0.09	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	—	—	0.09	—	0.09	—	—	0.16	0.16	—	—
<i>Aphredoderus sayanus</i>	0.48	0.24	—	0.15	—	0.26	0.26	0.51	0.32	—	—	—	—
<i>Aplodinotus grunniens</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	—	1.29	0.69	0.77	0.26	0.60	0.43	0.08	—	1.28	0.90	—
<i>Ctenopharyngodon idella</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cycleptus elongatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinus carpio</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Dorosoma cepedianum</i>	—	—	—	0.08	—	—	—	—	—	—	—	0.60	—
<i>Elassoma zonatum</i>	0.12	—	—	—	0.09	0.69	1.03	0.51	1.12	0.40	—	—	0.16
<i>Erimyzon sucetta</i>	0.24	—	—	0.08	—	0.09	0.09	—	0.24	0.16	0.24	0.15	0.16
<i>Esox americanus</i>	—	—	0.18	—	—	—	—	0.17	0.32	1.04	—	—	—
<i>Etheostoma asprigene</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Etheostoma chlorosoma</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Etheostoma gracile</i>	—	—	—	—	—	—	0.09	0.09	0.08	—	—	—	—

HWST													
Species	2019-05-31	2019-07-18	2019-08-28	2019-09-24	2019-10-14	2019-12-11	2020-01-06	2020-02-03	2020-03-06	2020-04-15	2020-05-11	2020-06-10	2020-07-15
<i>Micropterus salmoides</i>	0.12	0.12	0.46	0.38	0.43	0.51	0.34	0.17	0.08	0.08	0.08	0.75	0.16
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	3.60	3.00	—	—	—	—	—	0.26	—	—	—	—	0.48
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	0.24	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	0.46	0.26	0.43	0.09	0.26	—	—	0.16	0.15	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	0.28	—	0.60	0.60	0.09	0.17	—	0.16	0.08	—	—

OBOT

Species	2019-04-09	2019-05-08	2019-06-03	2019-07-01	2019-08-01	2019-10-08	2019-11-05	2019-12-12	2020-01-08	2020-02-07	2020-03-04	2020-04-01	2020-05-07	2020-06-11	2020-07-09	2020-08-11
<i>Fundulus olivaceus</i>	—	0.48	—	—	—	0.09	0.09	0.23	0.17	—	—	—	—	0.08	0.08	—
<i>Gambusia affinis</i>	0.30	2.04	0.24	1.20	1.56	1.71	1.66	0.60	0.69	0.24	0.69	1.09	0.24	0.24	0.96	—
<i>Hybognathus hayi</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictiobus bubalus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepisosteus oculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	0.08	—	—	—
<i>Lepisosteus osseus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis cyanellus</i>	—	—	0.12	—	—	—	—	—	—	0.06	—	—	0.08	—	—	—
<i>Lepomis gulosus</i>	0.10	0.36	—	—	0.60	—	0.18	0.08	—	0.06	0.17	0.22	1.52	0.64	0.40	—
<i>Lepomis humilis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis macrochirus</i>	—	0.36	0.12	—	—	0.09	0.18	0.15	0.17	0.06	0.26	0.22	0.08	—	0.48	0.53
<i>Lepomis marginatus</i>	—	0.72	1.80	0.28	—	0.43	0.28	0.08	0.09	0.06	—	0.87	2.96	1.68	0.24	—
<i>Lepomis megalotis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis microlophus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	0.30	1.80	1.20	0.37	0.12	1.63	0.92	0.98	0.43	0.54	1.29	0.98	5.04	0.72	1.68	0.40
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	—	0.12	—	—	—	—	0.08	—	0.06	—	—	—	0.16	0.08	—

OBOT

Species	2019-04-09	2019-05-08	2019-06-03	2019-07-01	2019-08-01	2019-10-08	2019-11-05	2019-12-12	2020-01-08	2020-02-07	2020-03-04	2020-04-01	2020-05-07	2020-06-11	2020-07-09	2020-08-11
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	0.09	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	—	—	0.24	—	—	—	—	0.08	—	—	—	—	—	—	—	—
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	0.11	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	0.09	—	—	0.09	—	—	0.33	0.08	—	0.24	—

OWMA

Species	2019-04-09	2019-05-08	2019-06-03	2019-07-01	2019-08-01	2019-09-05	2019-10-08	2019-11-05	2020-01-08	2020-02-07	2020-03-04	2020-04-01	2020-05-07	2020-06-11	2020-07-09	2020-08-11
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis symmetricus</i>	—	—	0.60	—	—	—	—	—	—	—	—	—	0.60	0.90	1.65	0.53
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	—	—	—	—	0.14	—	—	—	—	—	—	—	—	—	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	0.12	—	0.10	3.24	14.83	14.78	16.48	5.10	—	—	—	0.24	0.60	1.50	2.55	2.13
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	0.48	—	—	—	—	—	—	—	0.08	0.17	0.32	—	—	—	—

1880 Table 3k:

Species	SARC															
	2019-04-01	2019-05-07	2019-06-04	2019-06-28	2019-07-26	2019-09-04	2019-10-01	2019-12-05	2020-01-07	2020-02-03	2020-03-05	2020-04-02	2020-05-05	2020-06-03	2020-07-08	2020-08-04
<i>Ameiurus melas</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	0.08	—
<i>Ameiurus natalis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ameiurus nebulosus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Amia calva</i>	—	—	—	—	—	—	—	—	0.17	0.08	—	—	0.08	0.08	—	—
<i>Aphredoderus sayanus</i>	—	—	—	—	—	0.60	0.17	—	0.26	0.08	0.09	0.08	0.32	0.08	0.16	—
<i>Aplodinotus grunniens</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Centrarchus macropterus</i>	—	0.10	0.50	0.12	0.10	—	0.09	—	—	0.16	—	—	—	—	—	—
<i>Ctenopharyngodon idella</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cycleptus elongatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinella lutrensis</i>	—	—	—	—	—	—	—	—	—	—	0.17	—	—	—	—	—
<i>Cyprinella venusta</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Cyprinus carpio</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Dorosoma cepedianum</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Elassoma zonatum</i>	—	—	—	—	—	0.20	0.09	1.33	1.97	1.36	1.63	0.08	0.16	—	0.08	—
<i>Erimyzon sucetta</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Esox americanus</i>	—	1.00	0.20	—	—	—	—	—	—	—	0.17	0.08	0.16	—	—	0.08
<i>Etheostoma asprigene</i>	—	—	—	—	—	0.10	0.17	—	—	0.08	—	—	—	—	—	—
<i>Etheostoma chlorosoma</i>	—	0.10	—	0.96	0.10	0.50	0.34	—	—	0.16	—	0.16	—	—	0.08	0.08

SARC

Species	2019-04-01	2019-05-07	2019-06-04	2019-06-28	2019-07-26	2019-09-04	2019-10-01	2019-12-05	2020-01-07	2020-02-03	2020-03-05	2020-04-02	2020-05-05	2020-06-03	2020-07-08	2020-08-04
<i>Etheostoma gracile</i>	—	—	0.10	0.84	—	0.40	0.77	—	0.34	0.24	—	—	—	—	—	0.08
<i>Fundulus chrysotus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Fundulus olivaceus</i>	—	0.10	—	0.60	0.80	0.40	0.09	0.53	0.51	0.16	0.17	0.08	—	0.16	0.24	0.72
<i>Gambusia affinis</i>	0.30	0.30	—	0.48	0.70	0.20	0.69	1.47	0.34	0.48	0.43	—	0.24	0.32	0.24	0.16
<i>Hybognathus hayi</i>	—	—	—	0.12	—	—	—	—	—	—	—	—	0.16	—	—	—
<i>Hypophthalmichthys molitrix</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictalurus punctatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Ictiobus bubalus</i>	—	—	—	—	—	0.10	—	—	—	—	—	—	—	—	—	—
<i>Labidesthes sicculus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	0.08
<i>Lepisosteus oculatus</i>	—	—	0.10	—	—	—	—	—	—	—	—	—	0.08	—	—	—
<i>Lepisosteus osseus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	0.08	—
<i>Lepomis cyanellus</i>	—	—	—	—	—	0.20	0.09	—	—	—	—	—	—	—	—	—
<i>Lepomis gulosus</i>	0.40	1.00	0.20	0.24	0.10	0.40	0.69	0.80	0.17	0.48	0.77	0.80	0.48	0.32	0.08	0.40
<i>Lepomis humilis</i>	—	—	—	0.24	0.10	—	—	—	—	—	—	0.08	0.08	0.08	0.08	0.08
<i>Lepomis macrochirus</i>	0.70	2.10	0.70	0.12	0.60	1.10	0.60	0.27	0.09	0.16	1.29	1.36	1.04	0.64	0.80	2.08
<i>Lepomis marginatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Lepomis megalotis</i>	—	—	—	—	—	0.10	—	—	—	—	—	—	—	—	0.24	0.64
<i>Lepomis microlophus</i>	—	0.10	0.10	—	—	—	—	—	—	—	0.09	—	0.08	—	—	—

SARC

Species	2019-04-01	2019-05-07	2019-06-04	2019-06-28	2019-07-26	2019-09-04	2019-10-01	2019-12-05	2020-01-07	2020-02-03	2020-03-05	2020-04-02	2020-05-05	2020-06-03	2020-07-08	2020-08-04
<i>Lepomis miniatus</i>	—	0.10	—	—	0.10	—	—	—	—	—	0.09	0.08	0.08	0.16	—	—
<i>Lepomis symmetricus</i>	0.30	0.20	0.10	—	0.40	—	—	0.53	0.09	0.32	0.34	0.40	0.24	—	0.08	0.08
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	—	0.10	—	0.12	0.10	0.20	—	—	—	—	0.09	0.24	0.08	—	0.32	0.24
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	0.08
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	—	—	—	—	0.10	—	—	—	—	—	—	—	—	—	—	0.08
<i>Notropis maculatus</i>	—	—	—	0.24	0.10	0.20	0.09	—	—	—	0.09	0.24	0.16	0.16	0.16	0.40
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	0.60	—	—	—	—	—	—	—	—	—	0.08	0.08	0.16	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	0.40	0.08	—	—
<i>Pomoxis annularis</i>	—	—	—	—	0.20	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	0.10	0.17	0.13	—	—	—	0.08	0.16	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	0.10	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	0.08	—

SWAN

Species	2019-04-01	2019-05-07	2019-06-04	2019-06-28	2019-07-26	2019-09-04	2019-10-01	2019-12-05	2020-01-07	2020-02-03	2020-03-05	2020-04-02	2020-05-05	2020-06-03	2020-07-08	2020-08-04
<i>Lepomis miniatus</i>	—	—	—	—	—	—	—	0.17	—	—	—	0.08	—	—	—	—
<i>Lepomis symmetricus</i>	—	—	—	0.36	0.40	0.70	8.42	—	—	—	0.10	—	0.64	0.90	0.96	0.67
<i>Menidia beryllina</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Micropterus salmoides</i>	0.20	—	—	—	0.30	0.20	1.08	0.26	0.26	0.09	—	—	—	—	0.32	—
<i>Minytrema melanops</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Morone mississippiensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Notemigonus crysoleucas</i>	—	—	0.10	0.72	0.60	0.60	4.84	—	0.09	0.09	0.10	0.80	0.08	0.75	0.48	0.40
<i>Notropis maculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Noturus gyrinus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Opsopoeodus emiliae</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Percina caprodes</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pimephales vigilax</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis annularis</i>	—	—	—	—	0.10	—	—	—	—	—	—	—	—	—	—	—
<i>Pomoxis nigromaculatus</i>	—	—	—	—	—	0.30	1.61	0.17	0.09	0.09	0.30	0.32	—	—	—	—
<i>Sander canadensis</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Semotilus atromaculatus</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—
<i>Umbra limi</i>	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—	—

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