

1 Long-term trends in seasonality and abundance of three key 2 zooplankters in the upper San Francisco Estuary

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19 samples.

20 Abstract (300 words)

21 Zooplankton provide critical food for threatened and endangered fish species in the
22 San Francisco Estuary (estuary). Reduced food supply has been implicated in the Pelagic
23 Organism Decline of the early 2000s and further changes in zooplankton abundance,

24 seasonality, and distribution may continue to threaten declining fishes. While we have a
25 wealth of monitoring data, we know little about the abundance trends of many estuary
26 zooplankton species. To fill these gaps., we reviewed past research and then examined trends
27 in seasonality and abundance from 1972 to present of three key but understudied zooplankton
28 species (*Bosmina longirostris*, *Acanthocyclops* spp., and *Acartiella sinensis*) that play
29 important roles in the estuary food web. All three species were found in the stomach contents
30 of important estuarine fishes. We fit Bayesian generalized additive mixed models of each
31 species' relationships with salinity, seasonality, year, and geography on an integrated
32 database of zooplankton monitoring in the upper estuary. We found marked changes in the
33 seasonality and overall abundance of each study species. *B. longirostris* no longer peaks in
34 abundance in the fall months, *Acanthocyclops* spp. dramatically declined in all months and
35 lost its strong relationship with salinity, and *A. sinensis* adult abundance has become
36 increasingly driven by salinity while juveniles have developed wider seasonal abundance
37 peaks. In this process, we have documented the relationship of each species with salinity and
38 seasonality since the beginning of monitoring or their introduction, thus increasing our
39 understanding of their ecology and importance in the estuary. These results can inform food
40 web models, be paired with fish data to model the contributions of these species toward fish
41 abundance trends, and be mirrored to elucidate other species' trends in future studies.

42 **Keywords**

43 Zooplankton, phenology, salinity, monitoring, generalized additive modeling, copepods,
44 cladocerans, *Bosmina longirostris*, *Acanthocyclops*, *Acartiella sinensis*

45 **Introduction**

46 Zooplankton are critical components of aquatic food webs, connecting primary
47 producers to upper trophic species such as fishes. Most fishes rely on zooplankton food in
48 their larval stages when the starvation risk is highest (Hunter 1981). The seasonality of fish

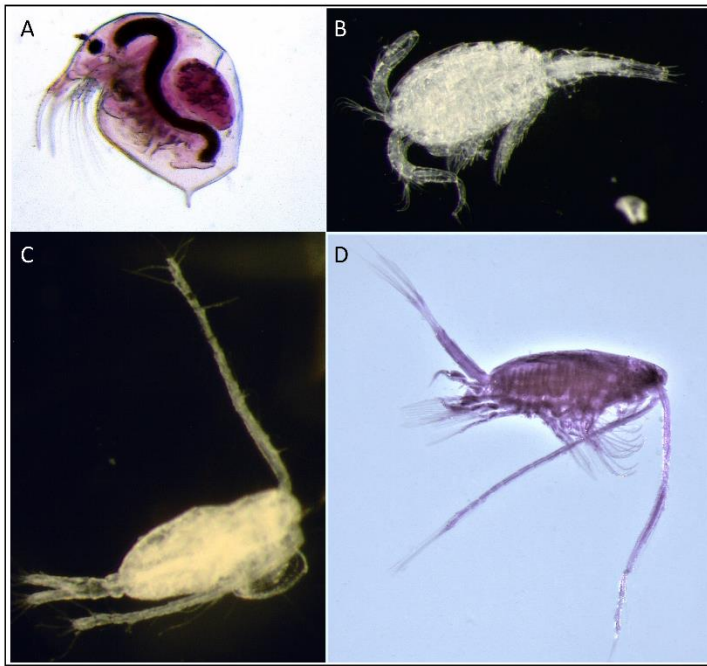
49 reproduction is often timed such that larvae coincide with peaks in the abundance of their
50 zooplankton prey (Cushing 1969; Cushing 1990). However, when the seasonality of
51 zooplankton abundance shifts due to climate change, species introductions, or other factors,
52 larval starvation risk can increase as their peak abundance no longer coincides with peak food
53 availability (Edwards and Richardson 2004; Durant et al. 2007). Thus, an understanding of
54 the patterns in zooplankton abundance is critical to understanding the drivers of fish
55 abundance.

56 The San Francisco Estuary (estuary) is home to several fish species listed under the
57 United States Endangered Species Act and/or the California Endangered Species Act. Most
58 depend on zooplankton prey for part or all of their life cycle. One of these species, Delta
59 Smelt (*Hypomesus transpacificus*) is endemic to the estuary and relies on zooplankton for its
60 full life cycle (Brown et al. 2016). Another fish, Longfin Smelt (*Spirinchus thaleichthys*),
61 also relies on zooplankton throughout its life cycle (P. Chigbu and Sibley 1998; Paulinus
62 Chigbu and Sibley 1998; Barros et al. 2022). A number of fish species, including Delta
63 Smelt and Longfin Smelt, declined sharply in abundance in the early 2000s, during the
64 “pelagic organism decline” (Feyrer et al. 2007; Sommer et al. 2007). This decline is thought
65 to have been caused in part by reduced zooplankton food supply (Winder and Jassby 2011;
66 Brown et al. 2016; Moyle et al. 2016).

67 Zooplankton research in the estuary to date has focused on a few key taxa, most
68 notably the copepods *Pseudodiaptomus forbesi*, *Eurytemora affinis*, and *Limnoithona*
69 *tetraspina*, as they are important food for threatened Delta Smelt and Longfin Smelt. (Bouley
70 and Kimmerer 2006; Kimmerer et al. 2014; Kayfetz and Kimmerer 2017; Kimmerer et al.
71 2018). In floodplains adjacent to the Delta, some cladocerans have also received attention,
72 particularly *Daphnia* spp., due to their importance in the diet of juvenile salmonids (Goertler
73 et al. 2018; Corline et al. 2021).

74 One prior study has investigated changes in zooplankton phenology in the estuary.
75 Merz et al. (2016) used a high-level approach to evaluate changes (from 1972-2014) to the
76 date of maximum abundance of five zooplankton taxa: *E. affinis*, *Pseudodiaptomus* spp.,
77 other calanoids, cyclopoids, and non-copepods. They found that the peak abundances of all
78 these zooplankton groups except the cyclopoids have shifted weeks to months earlier.
79 However, analyses of the date of peak abundance can overlook the presence of multiple
80 seasonal abundance peaks. Furthermore, the coarse taxonomic resolution of this study
81 excluded analyses of some key species. Many members of the zooplankton community
82 remain understudied, despite their role in the pelagic food web

83 In this study, we extensively review prior knowledge and derive new insights from
84 monitoring data on three key but understudied zooplankton taxa. We chose to focus on
85 *Bosmina longirostris*, *Acartiella sinensis*, and *Acanthocyclops* spp. (Fig. 1) due to their
86 importance in estuarine ecology and fish diets as well as the paucity of prior studies on their
87 dynamics in this estuary. We first review past studies from this estuary and other regions
88 where these taxa occur, then describe the importance of each taxa in fish diets by tabulating
89 their records in fish diets, and lastly examine their long-term trends in seasonality and
90 abundance in the estuary by applying Bayesian generalized additive mixed models to an
91 integrated database of zooplankton monitoring data (Bashevkin et al. 2020; Bashevkin et al.
92 in review).



93

94 Figure 1. Photographs (not to scale) of our study species from the San Francisco Estuary. A) *Bosmina*
 95 *longirostris* adult (credit: CDFW Fish Restoration Program), B) *Acanthocyclops* spp. adult (credit: Tricia
 96 Bippus/CDFW), C) *Acartiella sinensis* juvenile (credit: Tricia Bippus/CDFW), D) *Acartiella sinensis* adult
 97 (credit: Michelle Avila/CDFW Fish Restoration Program).

98 Prior research on the study species

99 *Bosmina longirostris*

100 *B. longirostris* (Fig. 1A) is the most abundant cladoceran in the freshwater reaches of
 101 the Sacramento San Joaquin Delta (Bashevkin et al. 2020; Jeffres et al. 2020), where it has
 102 been a major component of the zooplankton community over the past 40 years (Ambler et al.
 103 1985). Like *Daphnia* spp., it is abundant in off-channel habitat in the estuary (Corline et al.
 104 2021), but unlike *Daphnia* spp. it is also abundant in the larger channels of the South Delta
 105 (Bashevkin et al. 2020; Jeffres et al. 2020). *Bosmina* spp. is consumed by fish in the estuary
 106 and elsewhere, including juvenile salmonids and Delta Smelt (Roegner et al. 2015; Goertler
 107 et al. 2018; Slater et al. 2019), but it rarely occurs in diets of larval Longfin Smelt (Jungbluth
 108 et al. 2021; Burris et al. in review), likely due to a mismatch in the salinity preferences of
 109 *Bosmina* spp. and Longfin Smelt. Its small size makes it generally less preferred than larger

110 *Daphnia* spp. in diets of juvenile salmon (Craddock et al. 1976; Holm and Møller 1984), and
111 diets with high percentages of *Bosmina* spp. in juvenile perch result in low growth rates
112 (Romare 2000). Both *Daphnia* spp. and *Bosmina* spp. tend to have lower nutritional quality
113 (as measured by fatty acids) than copepods (Kratina and Winder 2015).

114 *B. longirostris* is a small, filter-feeding cladoceran found in lakes and rivers
115 throughout the world, where it has received little attention due to its small body size
116 (Adamczuk 2016). Despite its small size in comparison to members of the better-studied
117 genus *Daphnia*, *B. longirostris* can still have a large impact on the food web. *B. longirostris*
118 feeds broadly on phytoplankton, protists, and bacteria, ranging from 1-15 μm , and
119 preferentially consumes algae over bacteria, when available (Balcer et al. 1984; Onandia et
120 al. 2015). *B. longirostris* is particularly effective in depressing biomass of small
121 phytoplankton (Carpenter and Kitchell 1984) and is an efficient consumer of ciliates and
122 bacteria (Jürgens et al. 1996), playing a key role in the microbial loop. It can also thrive on
123 many types of cyanobacteria (Tönno et al. 2016). *B. longirostris* is more tolerant of
124 disturbance than many species of *Daphnia* (Hart 2004; Adamczuk 2016), with greater
125 resistance to toxic cyanobacteria (Matveev and Balseiro 1990; Jiang et al. 2013; Jiang et al.
126 2014; Jiang et al. 2017), and a higher salinity tolerance than many other cladocerans
127 (Adamczuk 2016). *Bosmina* spp. also has a higher thermal tolerance than many other
128 freshwater zooplankton (Drenner et al. 1981; Jiang et al. 2014), leading to increasing
129 dominance over other zooplankton as temperatures rise. *B. longirostris* typically lives 20-50
130 days and reproduces parthenogenically, producing 2-6 broods of 2-4 embryos each, though
131 temperature, salinity, and predation pressure may impact life span and reproduction
132 (Adamczuk and Mieczan 2019).

133 *Acanthocyclops* spp.

134 *Acanthocyclops* spp. (Fig. 1B) is a cyclopoid of unknown origin which occurs mostly
135 in freshwater (Orsi and Mecum 1986). Previous studies in the estuary identified the species of
136 *Acanthocyclops* in this region as *Acanthocyclops vernalis*; however, research in other areas
137 discovered that *A. vernalis* is a species complex consisting of three cryptic species;
138 *Acanthocyclops robustus*, *Acanthocyclops americanus*, and *A. vernalis* (Alekseev et al. 2002;
139 Dodson et al. 2003; Miracle et al. 2013; Alekseev 2021). Due to possible morphological
140 misidentifications, it is not known whether all three species in the *A. vernalis* complex are
141 native to the estuary and were present in the past; however, Jungbluth et al. (2021) did find
142 molecular evidence of all these species in larval Longfin Smelt diets recently collected in this
143 region.

144 Before the introduction of the small cyclopoid *L. tetraspina* in 1993 (Orsi and
145 Ohtsuka 1999), *Acanthocyclops* spp. was the most abundant cyclopoid in the estuary (Orsi
146 and Mecum 1986). *Acanthocyclops* spp. is an important component of fish diets in this
147 region, particularly Longfin Smelt (CDFW unpublished data; Hobbs et al. 2006; Burris et al.
148 in review) and larval Pacific Herring (*Clupea pallasii*) (Jungbluth et al. 2021), with
149 *Acanthocyclops* spp. being the most common cyclopoid consumed by larval Longfin Smelt
150 (Jungbluth et al. 2021). *Acanthocyclops* spp. also has a higher nutritional value than *L.*
151 *tetraspina*, and a similar fatty acid composition to that of the calanoid copepods *E. affinis* and
152 *P. forbesi* (Kratina and Winder 2015).

153 While research on the ecology and biology of *Acanthocyclops* spp. is relatively
154 limited in the estuary, the *A. vernalis* complex has been studied extensively in other regions
155 including Europe, Russia, and the Great Lakes. Studies in these areas show that, despite the
156 morphological similarities, *A. vernalis*, *A. robustus*, and *A. americanus* have different
157 ecologies, life cycles, and environmental tolerances (Alekseev et al. 2002; Miracle et al.

158 2013; Alekseev 2021). *A. vernalis* and *A. robustus* inhabit littoral or near-benthic areas and *A.*
159 *vernalis* has a benthic naupliar stage (Alekseev et al. 2002; Miracle et al. 2013) and vertically
160 migrates at night (Evans and Stewart 1977). Both *A. vernalis* and *A. robustus* are also
161 predatory, consuming copepod nauplii, cladocerans, rotifers, and occasionally larval fish
162 (Anderson 1970; Kerfoot 1978; Gliwicz and Stibor 1993; Piasecki 2000). *A. americanus*,
163 however, is pelagic throughout its lifecycle, found in varying salinities from the
164 Mediterranean Sea (Alekseev 2021) to freshwater lakes (Alekseev et al. 2002), and is
165 omnivorous. *A. americanus* nauplii consume primarily algae, with later life stages also
166 consuming filamentous algae and cyanobacteria, in addition to cladocerans, nauplii, and
167 rotifers (Enríquez-García et al. 2013; Sarma et al. 2019). All species can produce about 100
168 eggs per female, develop to sexual maturity in 10-14 days, and live 30-75 days depending on
169 conditions, with *A. americanus* growing and reaching maturity faster than the others
170 (Alekseev 2021). Species in the *A. vernalis* complex likely have different environmental
171 tolerances, as has been shown by studies of their seasonal and spatial variation in abundance
172 in other regions. *A. vernalis* may be more tolerant of colder temperatures than the other
173 species, whereas *A. americanus* could have a higher temperature tolerance based on
174 laboratory experiments and timing of peak abundance in areas outside the estuary (Alekseev
175 2021).

176 *Acartiella sinensis*

177 In the fall of 1993, the non-native calanoid copepod *A. sinensis* (Fig. 1C,D) was first
178 detected in the estuary (Orsi and Ohtsuka 1999). Likely introduced via the ballast water of
179 ships, the large (~1.2 – 1.5 mm in length) predatory calanoid is native to the estuaries and
180 coasts of Southeast Asia (Shen and Lee 1963; Srinui and Ohtsuka 2015). The species has
181 been recorded from estuaries along the East China Sea in salinities around 18-21 ppt (Shen
182 and Lee 1963) to the brackish marshes of Thailand in salinities around 5 ppt (Srinui and

183 Ohtsuka 2015). In the Pearl River estuary of China, *A. sinensis* was the dominant copepod in
184 brackish waters with salinities less than 15 ppt (Tan et al. 2004). Sampling in the Thale-Noi
185 lake of Thailand showed changes in temperature and salinity were the main environmental
186 variables impacting densities of *A. sinensis* in the region (Inpang 2008).

187 Within a year after invasion, *A. sinensis* became the second most common calanoid
188 copepod in the upper estuary, with its highest abundances in the lower salinity zone of Suisun
189 and the West Delta during summer and fall (Hennessy 2018). The predatory *A. sinensis* has
190 been shown to feed on the nauplii and copepodid life stages of other copepods in the estuary,
191 primarily those of the abundant *L. tetraspina* and *P. forbesi* (York et al. 2014; Slaughter et al.
192 2016; Kayfetz and Kimmerer 2017). *A. sinensis* has also become a food item for the
193 endangered Delta Smelt, appearing in their diets in summer and fall (Slater and Baxter 2014;
194 Slater et al. 2019).

195 The invasion of *A. sinensis* was not without consequence. Since the invasion in 1993,
196 the zooplankton assemblage in the low-salinity Suisun area has shifted in trophic
197 composition. Once dominated by herbivorous cladocerans and copepods such as *E. affinis*
198 and *P. forbesi*, the community has become more “top-heavy” with the spread of *A. sinensis*
199 (Kratina et al. 2014; Kratina and Winder 2015). The sustainment of *A. sinensis* in the low-
200 salinity zone and its high predation rate on the nauplii of *P. forbesi* is linked to a shift in the
201 distribution of *P. forbesi* out of the low-salinity zone and upriver into more freshwater
202 habitats (Kayfetz and Kimmerer 2017). This shift in the distribution of the important *P.*
203 *forbesi* could have implications for the majority of planktivorous fishes that feed on the
204 calanoid copepod populations.

205 Materials and Methods

206 Diet data

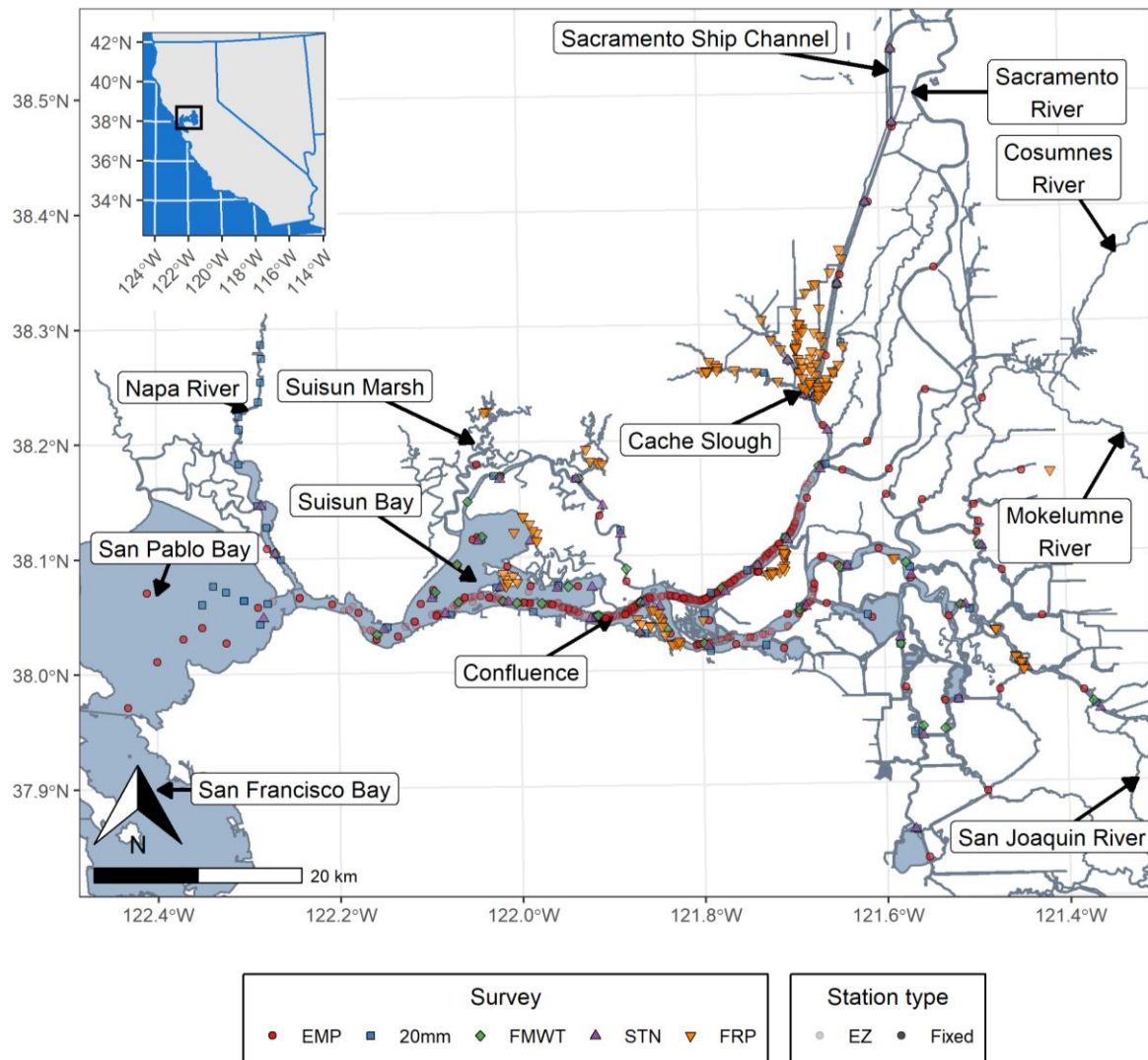
207 To investigate the occurrence of our study species in fish diets, we summarized data
208 from the California Department of Fish and Wildlife (CDFW) Diet and Condition Study
209 (“Diet Study”, <https://wildlife.ca.gov/Conservation/Delta/Special-Studies>). The CDFW Diet
210 Study identifies and enumerates gut contents of young pelagic fishes in the estuary including
211 Delta Smelt, Longfin Smelt, Striped Bass (*Morone saxatilis*), Threadfin Shad (*Dorosoma*
212 *petenense*), American Shad (*Alosa sapidissima*), and other species of interest collected by
213 various Interagency Ecological Program monitoring surveys (Slater and Baxter 2014;
214 Hammock et al. 2017; Slater et al. 2019). Fish were weighed, measured, the guts dissected,
215 and prey items were identified to the lowest possible taxonomic level. In some cases, prey
216 items were heavily digested and therefore could only be identified to higher taxonomic levels
217 than our study taxa. Thus, these prey items were not included in our data. Fishes were chosen
218 for processing based on other study needs, so relative sample size for each fish is not
219 correlated to abundance in the estuary, but provides some information on relative rates of
220 zooplankton consumption for each species.

221 The number of fish processed, number with food in their stomachs, fork length range,
222 and number of fish that had eaten our focal taxa were extracted from the database. To
223 maintain the same spatial coverage as the zooplankton datasets, only fish caught in San Pablo
224 Bay and upstream were included in this analysis.

225 Zooplankton data

226 The data used for these analyses were obtained from an integrated database of five
227 long-term zooplankton monitoring surveys in the upper estuary. These include the
228 Environmental Monitoring Program (EMP), 20-mm Survey (20mm), Fall Midwater Trawl
229 (FMWT), Summer Townet (STN), and Fish Restoration Program monitoring (FRP). These

230 surveys are described in detail in Kayfetz et al. (2020) and Bashevkin et al. (in review).
231 Briefly, EMP samples monthly year-round since 1972, 20mm samples every other week
232 March through July since 1995, STN samples every other week June – August since 2005,
233 FWMT samples monthly September through December since 2011, and FRP samples
234 annually to monthly near tidal marshes (or areas soon to be restored to tidal marshes) March
235 through December since 2015. Each survey samples at a set of fixed stations (Fig. 2). EMP
236 also samples at a set of moving stations (entrapment zone, EZ) at locations where the bottom
237 conductivity is 2 and 6 mS/cm. Many of these surveys collect other parameters such as fish
238 abundance or water quality, but only the time period of zooplankton sampling is described
239 above. Furthermore, sampling locations and frequency have changed over time (Kayfetz et al.
240 2020; Bashevkin et al. in review).



241

242 Figure 2. Map of the study area, depicting all sampling stations. Survey abbreviations are as follows: EMP =
 243 Environmental Monitoring Program, 20mm = 20-mm Survey, FMWT = Fall Midwater Trawl, STN = Summer
 244 Towntnet, FRP = Fish Restoration Program. The EMP survey has some non-fixed entrapment zone (EZ) stations
 245 that move with the salinity field and are depicted here with increased transparency.

246 The data from each survey are integrated by the R package zooper (Bashevkin 2021),
 247 which imports and standardizes the data from each survey. For the purposes of these
 248 analyses, we selected data from the mesozooplankton size-class, which corresponds to
 249 samples from the modified Clarke-Bumpus nets used by each survey, with mesh sizes of 150
 250 – 160 μm . We selected all available data from each of our study species without missing
 251 values in our covariates. Adults and juveniles were analyzed separately since these distinct

252 life stages have different behaviors and abundance drivers and play distinct demographic
253 roles. *A. sinensis* was introduced to the estuary in 1993 (Orsi and Ohtsuka 1999), but adults
254 were first counted in samples in 1994 and juveniles first counted in 2006. Thus, adult data
255 were filtered to the start date of 1994 and juveniles to the start date of 2006. For the other
256 species, only adult data were available at the taxonomic resolution of our analysis. The final
257 dataset had 33,255 samples for *B. longirostris* adults, 32,026 samples for *Acanthocyclops*
258 spp. adults, 17,189 samples for *A. sinensis* adults, and 11,224 samples for *A. sinensis*
259 juveniles.

260 Exploratory data visualization revealed tighter relationships of each species'
261 abundance with log-transformed salinity than raw salinity values. Thus, salinity was natural
262 log-transformed for analyses. Furthermore, we centered all covariates (including log-
263 transformed salinity) by their mean and standardized them by their standard deviation. Lastly,
264 since many of the sampling stations from the different surveys were nearby one another (Fig.
265 2), we clustered all stations into groups within a 1 km radius.

266 Model structure

267 We fit Bayesian generalized additive mixed models with the R package brms
268 (Bürkner 2017; Bürkner 2018), which uses the Bayesian modeling platform Stan (Stan
269 Development Team 2021), as well as the R package mgcv (Wood 2011) to handle the smooth
270 construction. Models were fit with a hurdle lognormal distribution, using the catch per unit
271 effort (CPUE; organisms m^{-3}) of the specified taxa and life stage as the response variable. A
272 hurdle model was used to account for the large number of 0s in the data.

273 Our overall approach was to model the probability of presence with a smoothed
274 function of salinity, and the non-zero abundances with smooth functions of day of year,
275 salinity, year, and their interactions, while accounting for space with a random intercept for

276 each station. Our combination of a Bayesian method, which propagates uncertainty and
277 handles unbalanced data, with a generalized additive model approach, which accounts for key
278 covariates like salinity and seasonality, allowed us to resolve inconsistencies in the sampling
279 designs, and incorporate both fixed stations and stations that move with the salinity field.

280 The hurdle probability (probability of 0 CPUE) was modeled with a cubic regression
281 spline of salinity with a low basis dimension (k) of 5 since the relationship was expected to
282 have a simple shape. The abundance of *Acartiella sinensis* was so strongly seasonal that we
283 modified the hurdle component to also include seasonality. Thus, for *A. sinensis* we modeled
284 the hurdle probability with a two-dimensional tensor product smooth of salinity (cubic
285 regression spline, $k=5$) and day of year (cyclic cubic regression spline, $k=4$).

286 The non-zero CPUEs were modeled with a three-dimensional tensor product smooth
287 of day of year (cyclic cubic regression spline, $k=13$), salinity (cubic regression spline, $k=5$),
288 and year (cubic regression spline, $k=5$). The basis dimension for day of year was set to 13 to
289 match the monthly nature of these sampling programs (since the smooth is cyclical, a basis
290 dimension of 13 has 12 independent functions). The basis dimension for salinity was set to
291 the low value of 5 because the relationship with salinity was expected to be a simple
292 unimodal shape, and a basis dimension of 5 would still allow much greater complexity than
293 that. The basis dimension for year was set to the low value of 5 because we were interested in
294 evaluating broad long-term patterns, rather than fine-scaled year-to-year abundance trends.
295 Thus, the results of this model represent broad long-term trends, not predicted abundances on
296 specific years. For juvenile *A. sinensis*, the basis dimension for year was reduced to 3 since
297 they have only been counted in samples since 2006 and thus the timeseries is shorter. We also
298 included a random intercept for each station cluster.

299 We fit separate models on each species and life stage (4 total). Models were run on
300 three chains, each for 5,000 iterations including 1,250 used for the warmup that were then
301 discarded. We used weakly informative priors as recommended by the Stan authors (Stan
302 Development Team 2021).

303 Each model was validated and checked prior to use. All models were inspected to
304 ensure adequate sampling by verifying the posterior effective sample size (> 100 per chain)
305 and Rhat values (< 1.05) (McElreath 2015). We further inspected visual plots comparing the
306 model outputs to the raw data to ensure they matched. These plots included the proportion of
307 zero values, the distribution of non-zero CPUE values, and the predicted non-zero CPUE
308 values for each row in the dataset. We also inspected the spatiotemporal variograms for
309 spatiotemporal autocorrelation using the R package gstat (Pebesma 2004; Gräler et al. 2016).
310 We detected some residual autocorrelation, and thus used a conservative significance
311 threshold of $p < 0.01$ to account for any potential impacts.

312 Model predictions

313 We visualized predicted values from the models to explore the abundance patterns of
314 each species and life stage. Predicted values were generated over a range of covariates that
315 included all combinations of 6 evenly spaced time-points per month (from 1972 to 2019) and
316 a series of salinity values selected as quantiles from the raw data (every 0.05 from 0.05 to
317 0.95). Since there were some gaps in the timeseries (e.g., winters were not sampled some
318 years in the 1970s and 1980s), those same gaps are preserved in the model predictions to
319 avoid extrapolation. Model predictions were then plotted along with their 99% credible
320 intervals. To visualize the multidimensional model outputs, we created three plots for each
321 set of model predictions. Each plot had one of the covariates (salinity, day of year, or year) on
322 the x-axis while the other two variables were illustrated with color or separate plots. For the
323 two covariates included as color or separate plots, we chose a subset of the unique values

324 used in generating model predictions in order to reduce plot complexity and aid
325 interpretation. These values were chosen as an evenly spaced subset of the values available.

326 To explore spatial patterns in abundance, we extracted the mean estimated value from
327 each station cluster random intercept. These values were then plotted over a map of the study
328 region.

329 Results

330 Diet results

331 The CDFW Diet Study has processed 11,301 fish caught in San Pablo Bay and
332 upstream through the Delta consisting of nine species; Delta Smelt, Longfin Smelt, age- 0
333 Striped Bass, Threadfin Shad, American Shad, Mississippi Silversides (*Menidia beryllina*),
334 Pacific Herring, Prickly Sculpin (*Cottus asper*), and *Tridentiger* spp. gobies (Table 1). A
335 wide range of fork lengths was processed for all species, except for Pacific Herring, Prickly
336 Sculpin, and *Tridentiger* spp. Gobies which only included larval fish. Time and seasonality of
337 collection varied by fish species with year-round diet information only available for Delta
338 Smelt and Longfin Smelt.

339 Our study zooplankton were eaten by at least one individual of each fish species,
340 except Pacific Herring, which consumed none and is thus not discussed further here (Table
341 1). *Acanthocyclops* spp. were present in the most fish stomachs (n= 885) and *A. sinensis*
342 juveniles were eaten the least (n= 88). Of the fish that consumed *Acanthocyclops* spp., Delta
343 Smelt had the highest consumption (n= 800, 21% of fish with food present), and Striped Bass
344 the least (n= 3 fish, 0.3%). Neither Prickly Sculpin nor *Tridentiger* spp. consumed
345 *Acanthocyclops* spp.

346 *A. sinensis* adults were the second most consumed (n= 561), with at least one
347 individual of each fish species eating adult *A. sinensis*, except Prickly Sculpin. American

348 Shad had the highest consumption of adult *A. sinensis* (n= 20 stomachs, 45%) followed by
 349 Delta Smelt (n= 437, 12%), Mississippi Silversides (n= 15, 8%), *Tridentiger* gobies (n= 23,
 350 5%), Threadfin Shad (n= 17, 3%), Longfin Smelt (n= 48, 2%) and finally Striped Bass (n= 1,
 351 0.1%) (Table 1). *A. sinensis* juveniles were consumed the least of our study species, with only
 352 2% of Delta Smelt individuals (n= 74), and $\leq 1\%$ of individuals from the other fish species
 353 (*Tridentiger*: n= 4, 0.9%; Threadfin Shad: n= 4, 0.7%; Mississippi Silversides: n= 1, 0.5%;
 354 Longfin Smelt: n= 5, 0.2%; Striped Bass and Prickly Sculpin: n= 0) with *A. sinensis* juveniles
 355 present in stomachs.

356 *Bosmina* was present in the guts of all fish species (n= 235) and was the third most
 357 consumed of our study zooplankton. Threadfin Shad ate *Bosmina* the most (n= 53, 9%), Delta
 358 Smelt the second most (n= 163, 4%), and Longfin Smelt ate the least (n= 1, 0.04%).

359 Table 1: Diet results for fish processed by the CDFW Diet and Condition Study and the number of fish that
 360 consumed each zooplankton species. “# fish” represents the number of fish of each species that were processed
 361 for stomach contents. “Season” indicates the seasons in which fish were collected, with “Sum” = summer and
 362 “Win” = winter. Fork lengths are given in mm. Zooplankton species are represented as follows: A =
 363 *Acanthocyclops* spp., AS = *A. sinensis* adults, AS juv = *A. sinensis* juveniles, and B = *B. longirostris*.
 364 Percentages represent the percent of fish with food present in their stomachs that consumed each prey species.

Fish species	# fish	Season	Fish with food present (%)	Min fork length	Max fork length	A (%)	AS (%)	AS juv (%)	B (%)
Delta Smelt	4317	All	3752 (87)	4.3	90	800 (21)	437 (12)	74 (2)	163 (4)
Longfin Smelt	3640	All	2558 (70)	4.4	124	69 (3)	48 (2)	5 (0.2)	1 (0.04)
Striped Bass	1187	Sum to Fall	966 (81)	5.1	142	3 (0.3)	1 (0.1)	0	9 (0.9)

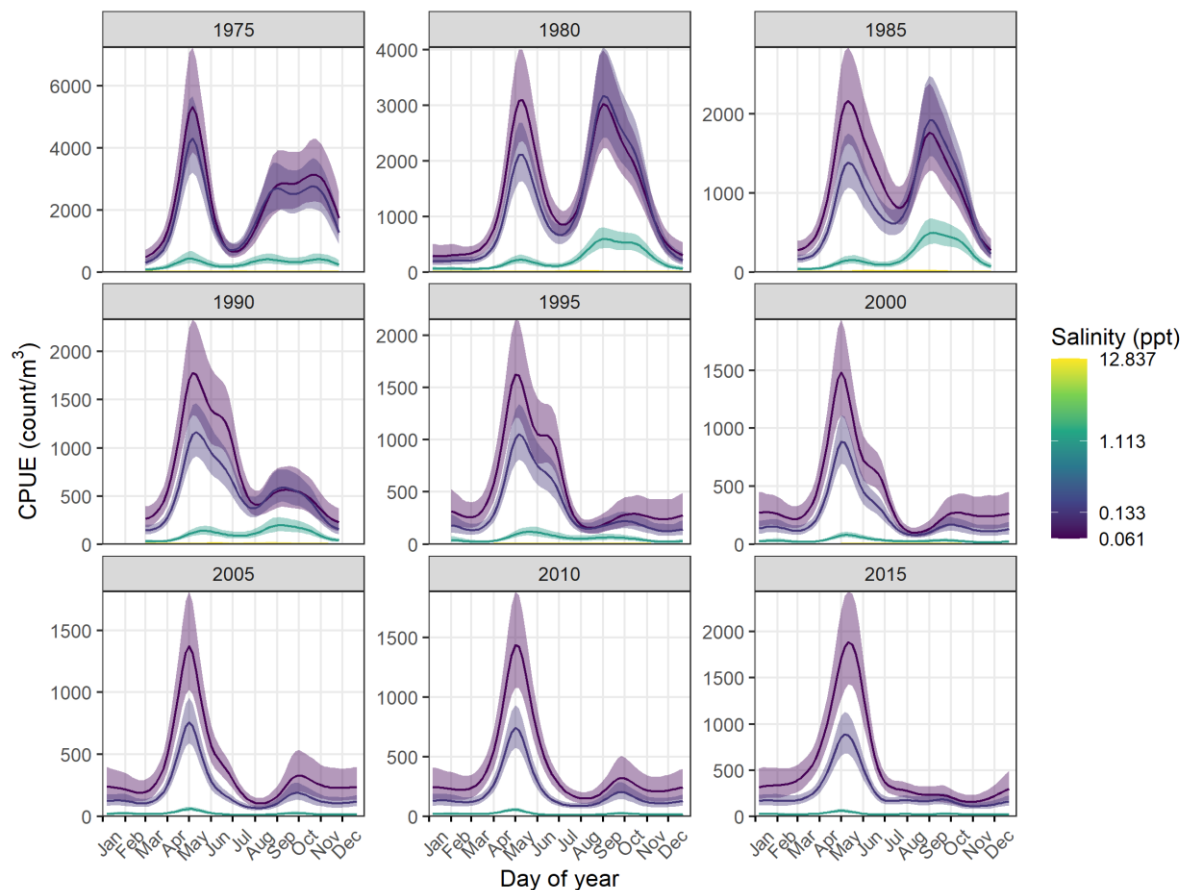
Threadfin	861	Sum to	596 (69)	4.0	98	8 (1.3)	17 (3)	4 (0.7)	53 (9)
Shad		Fall							
American	47	Sum to	44 (94)	19	90	0	20 (45)	2 (4.5)	0
Shad		Fall							
Mississippi	192	Sum to	189 (98)	16.5	95.8	5 (3)	15 (8)	1 (0.5)	6 (3)
Silversides		Fall							
Pacific	389	Win	252 (65)	8.5	35.2	0	0	0	0
Herring									
Prickly	195	Win	82 (42)	5.3	11.2	0	0	0	2 (2)
Sculpin									
<i>Tridentiger</i>	473	Sum	443 (93)	2.7	17	0	23 (5)	4 (0.9)	1 (0.2)
spp. gobies									

365

366 *Bosmina longirostris* adults

367 The abundance of adult *B. longirostris* has regularly peaked in the Spring between
368 April and May. In earlier years, they had another large peak in the late summer to fall
369 between August and October. In some years (1980 and 1985), this fall peak was as large as
370 the spring peak. However, by 1990 the fall peak was greatly reduced to just a small increase,
371 which has since continued to decrease in size, becoming non-existent by 2015 (Fig. 3).

372 *B. longirostris* adults are most abundant in freshwater, and abundance decreases as
373 salinity increases (Fig. S1). In the moderate salinity of 1.113, the fall peak was larger than the
374 spring peak in earlier years. However, as with the lower salinities, the fall peak was greatly
375 reduced by 1995, shrinking smaller than the spring peak even as the overall abundance in this
376 salinity continued to decrease over time (Fig. 3, S1).



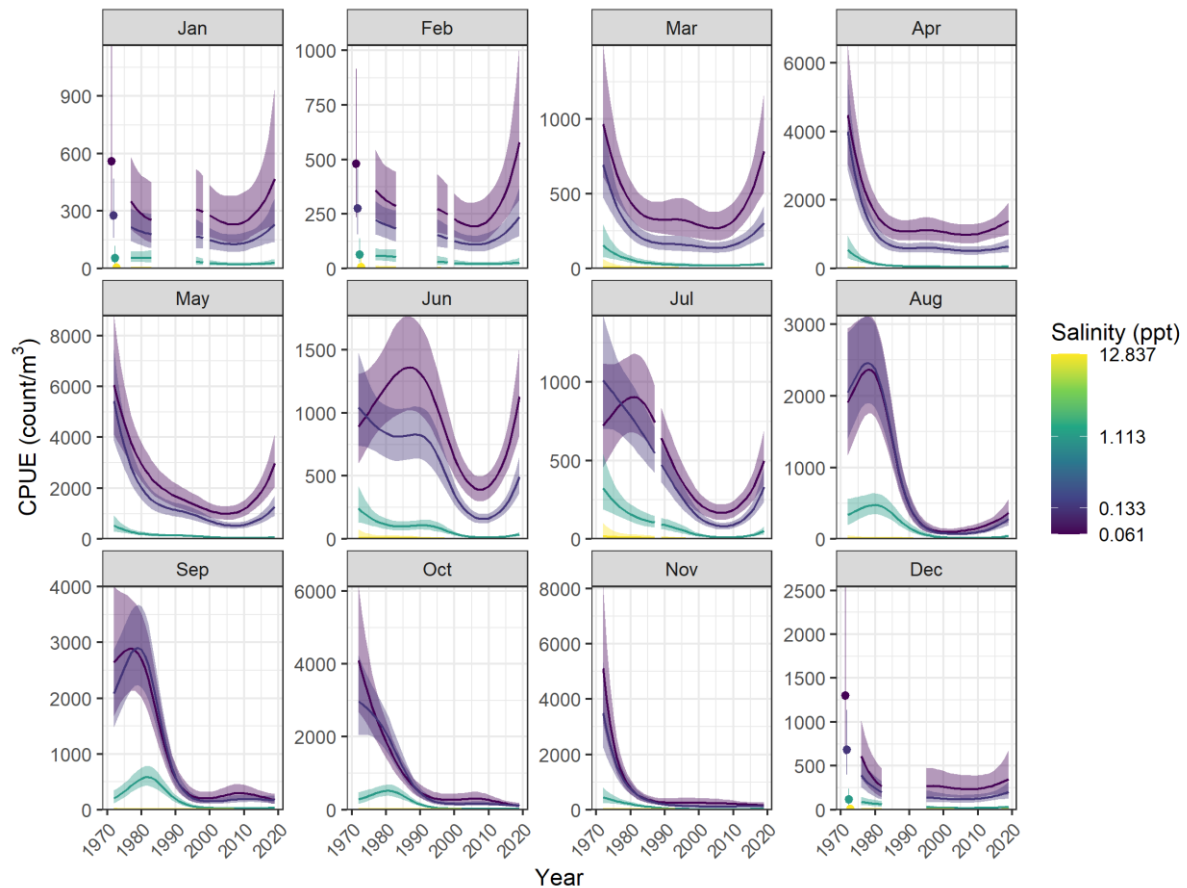
377

378 Figure 3. Seasonal patterns in *Bosmina longirostris* adult abundance with 99% credible intervals. Each plot
 379 represents predictions from different position on the smoothed yearly trend and may not represent exact
 380 conditions for that particular year. Predictions for different salinity values are represented by line and shading
 381 color, which is on the log scale. The y-axis limits differ among plots to facilitate comparison of seasonal trends.
 382 Absolute trends in abundance over time are represented in Fig. 4. Missing values (e.g., the months of Jan, Feb,
 383 and Dec in 1975) represent gaps in the raw data.

384 The abundance of adult *B. longirostris* has declined over time in most months. This is
 385 most apparent in August through November, corresponding to the loss of their former fall
 386 peak. In some months, their abundance has mostly decreased over time except in recent years
 387 which have a slight uptick. This recent uptick appears in January through March and May
 388 through July but is most significant in May through July (Fig. 4).

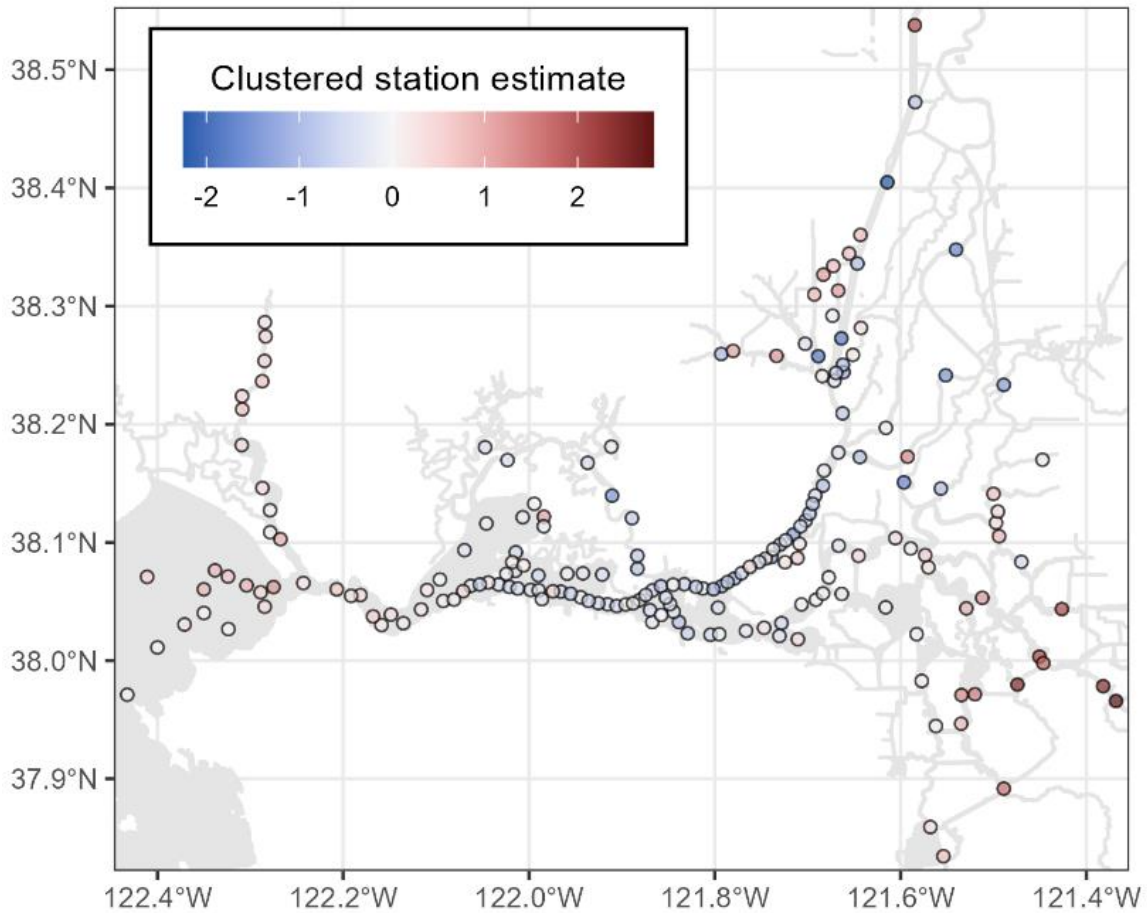
389 Controlling for all other factors (salinity, year, month), *B. longirostris* adults had the
 390 highest abundance in the Southeastern Delta. Other areas around the boundaries of the

391 sampled area also had higher abundance such as San Pablo Bay, Napa River, and parts of the
 392 Cache Slough Complex and Sacramento Deep Water Ship Channel. The Sacramento River
 393 corridor between Cache Slough and Suisun Bay generally had low abundance, as did parts of
 394 the Northeastern Delta (Fig. 5).



395

396 Figure 4. Yearly patterns in *Bosmina longirostris* adult abundance with 99% credible intervals. Each plot
 397 represents the pattern for a separate month. Predictions for different salinity values are represented by line and
 398 shading color, which is on the log scale. The y-axis limits differ among plots to facilitate comparison of long-
 399 term trends. Seasonal trends in abundance are represented in Fig. 3. Missing values represent gaps in the raw
 400 data.



401

402 Figure 5. Estimated values of the clustered station random intercepts for *Bosmina longirostris* adults. Stations
 403 within 1 km were clustered into groups. Point color indicates whether each station cluster has higher or lower *B.*
 404 *longirostris* adult abundance, after controlling for the other covariates (day of year, salinity, and year).

405 *Acanthocyclops* spp. adults

406 The abundance of *Acanthocyclops* spp. adults has peaked regularly in the spring from
 407 April to May in all years. This peak is apparent in the two lower salinity levels (0.062 and
 408 0.137) but generally not in the two higher salinities (1.402 and 13.017). In the second highest
 409 salinity of 1.402, abundance peaked in the winter from February through March in most
 410 years, although limited sampling in these months in some years may have masked the signal
 411 (Fig. 6).

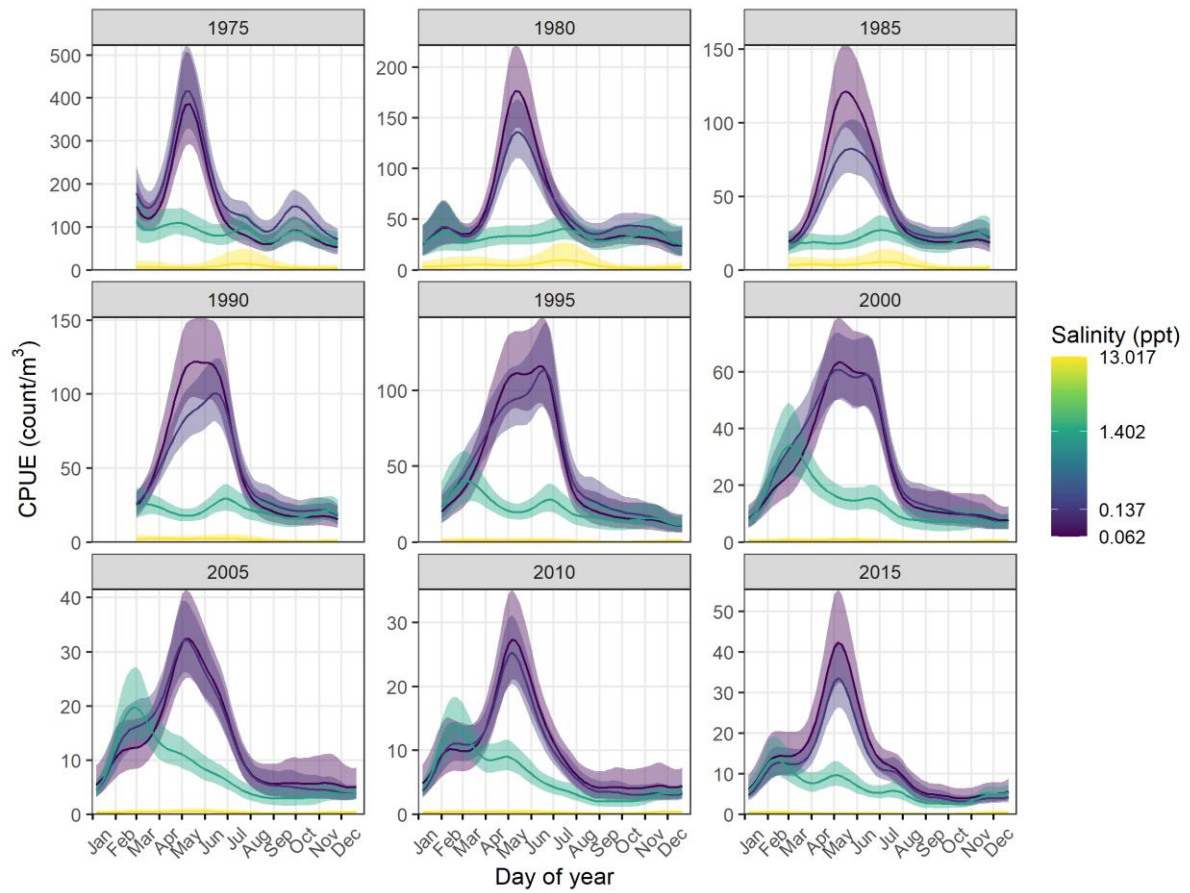
412 *Acanthocyclops* spp. were most abundant in the lower salinities, peaking around 0.3.

413 Abundance decreased on either side of that peak, but fell much lower in the highest salinities

414 (Fig. S2). Over time, the relationship of *Acanthocyclops* spp. with salinity leveled out such
415 that they became equally abundant in all salinity levels in most months (Fig. S2).

416 The abundance of *Acanthocyclops* spp. has dramatically declined over time in all
417 months. There was a slight uptick in the 1980s to 1990s in most months, but populations
418 crashed again after this period (Fig. 7). Overall trends in January and February appear mostly
419 flat with generally low abundance in all years but the missing data in those months may be
420 obscuring patterns.

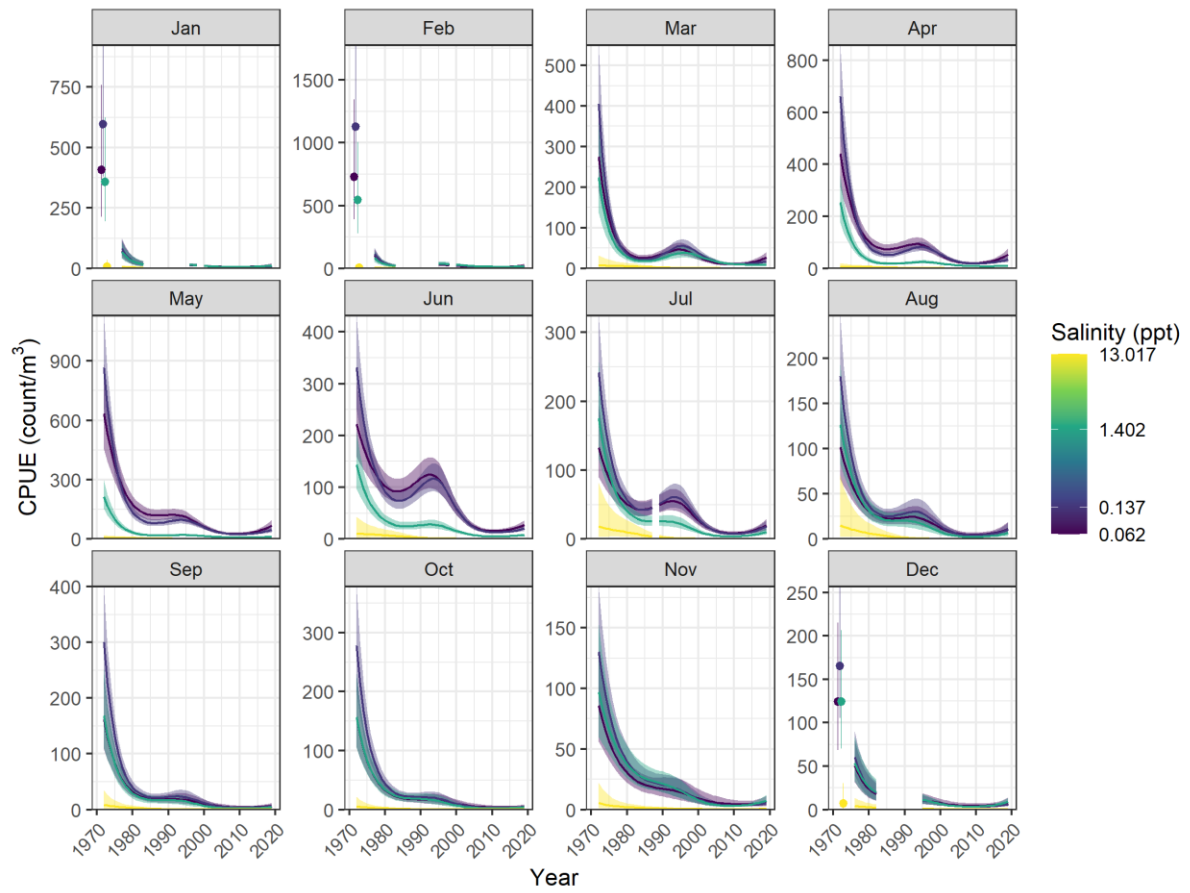
421 The highest *Acanthocyclops* spp. abundance after controlling for the other covariates
422 was in Cache Slough, Suisun Marsh, the Southeastern Delta, Carquinez Strait, San Pablo
423 Bay, and the Napa River. The lower Sacramento and San Joaquin rivers through Suisun Bay
424 generally had lower abundances, as did most of the western-most areas (Fig. 8).



425

426 Figure 6. Seasonal patterns in *Acanthocyclops* spp. adult abundance with 99% credible intervals. See Fig. 3 for a

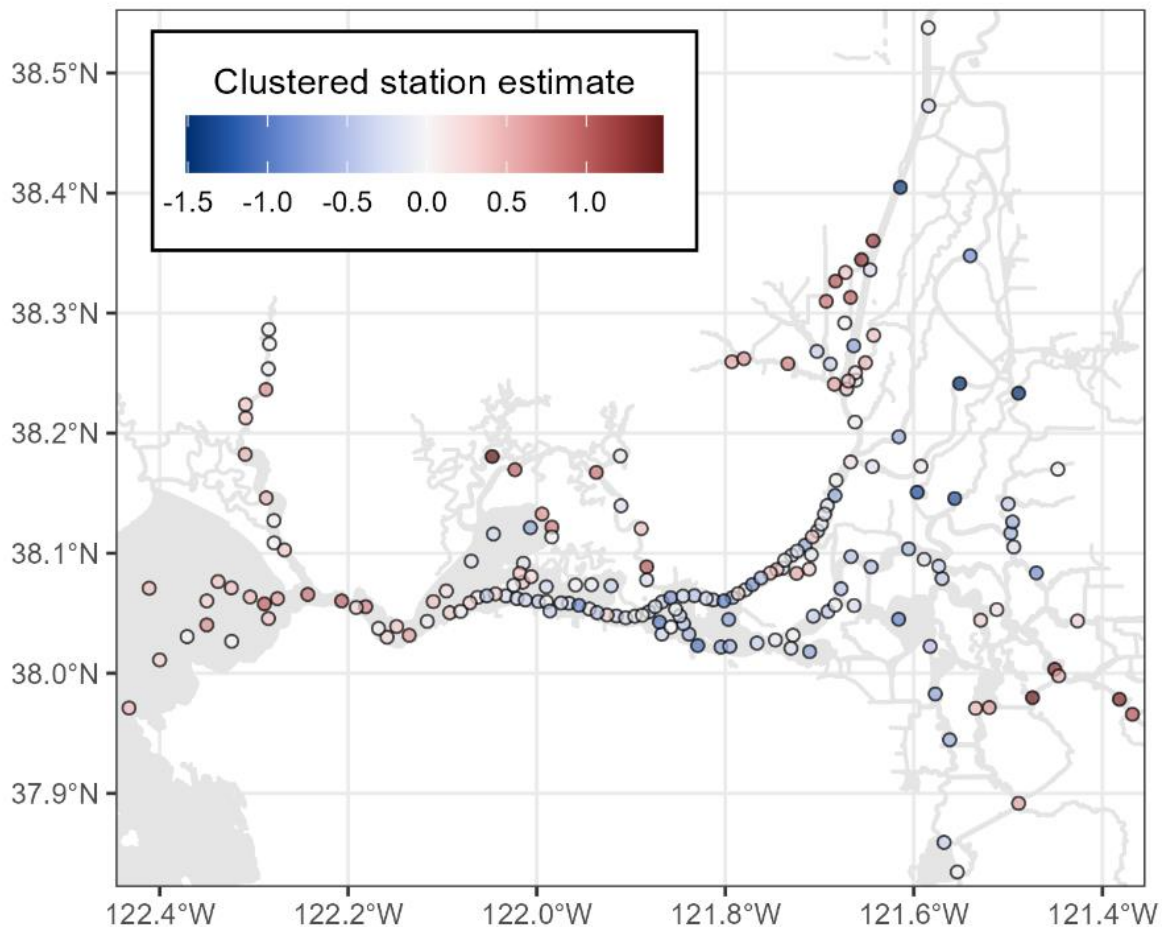
427 full description.



428

429 Figure 7. Yearly patterns in *Acanthocyclops* spp. adult abundance with 99% credible intervals. See Fig. 4 for a

430 full description.



431

432 Figure 8. Estimated values of the clustered station random intercepts for *Acanthocyclops* spp. adults. See Fig. 5
 433 for a full description.

434 *Acartiella sinensis* adults

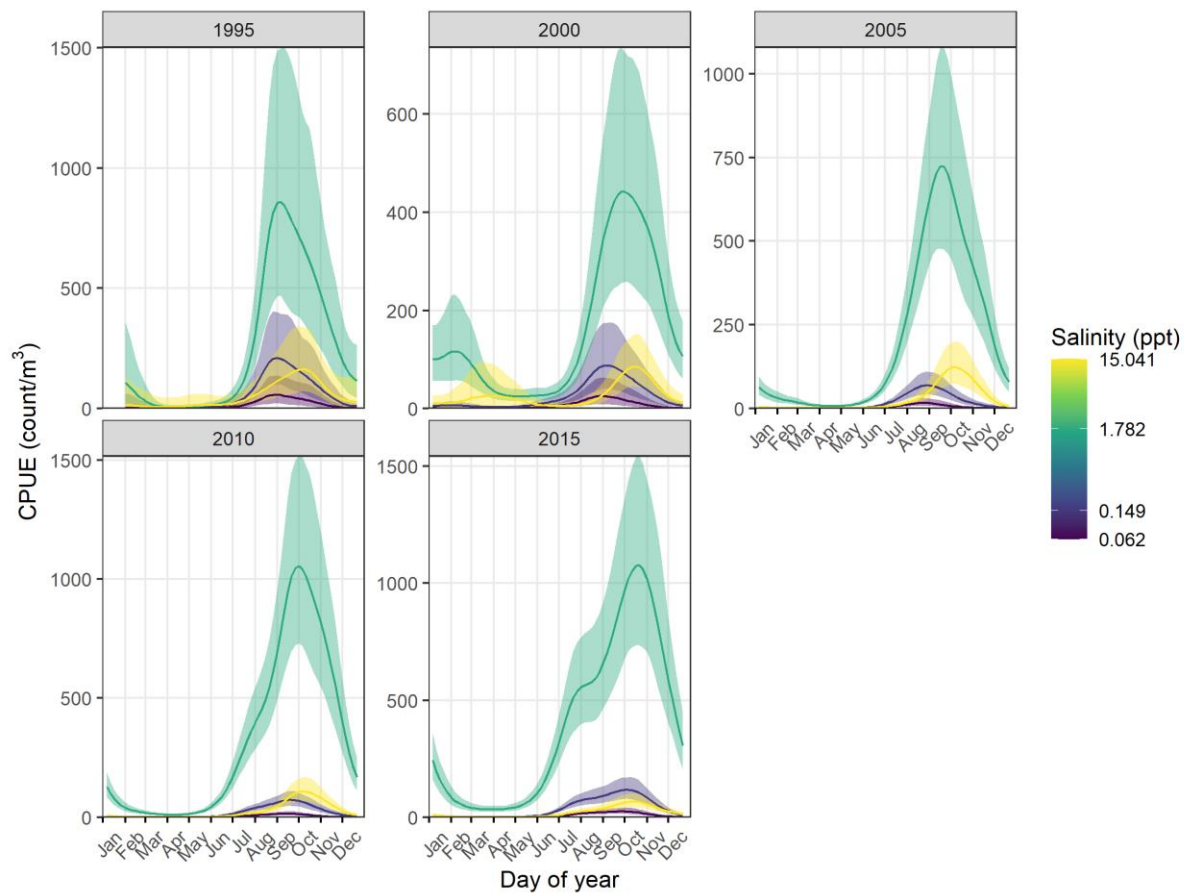
435 *A. sinensis* had the strongest seasonality of the three species investigated. Adults peak
 436 in the fall from August through December with the highest levels in September and October.
 437 Abundances then dip close to zero from February through May (Fig. 9).

438 Adult *A. sinensis* were most abundant in moderate salinities between about 1 and 4.
 439 The effect of salinity on abundance has increased over time, especially in May through July
 440 where the peak was greatly reduced in earlier years (Fig. S3).

441 Unlike the other two species, *A. sinensis* adults did not exhibit any overall long-term
 442 decreases in abundance. However, the timeseries is shorter, only starting in 1994. In most

443 months, the most recent abundance is similar to the earliest abundance level, but abundance
 444 did increase over time in March through July. The abundance peaked in the 2010s in most
 445 months and some months also had an earlier peak around 2000 (Fig. 10).

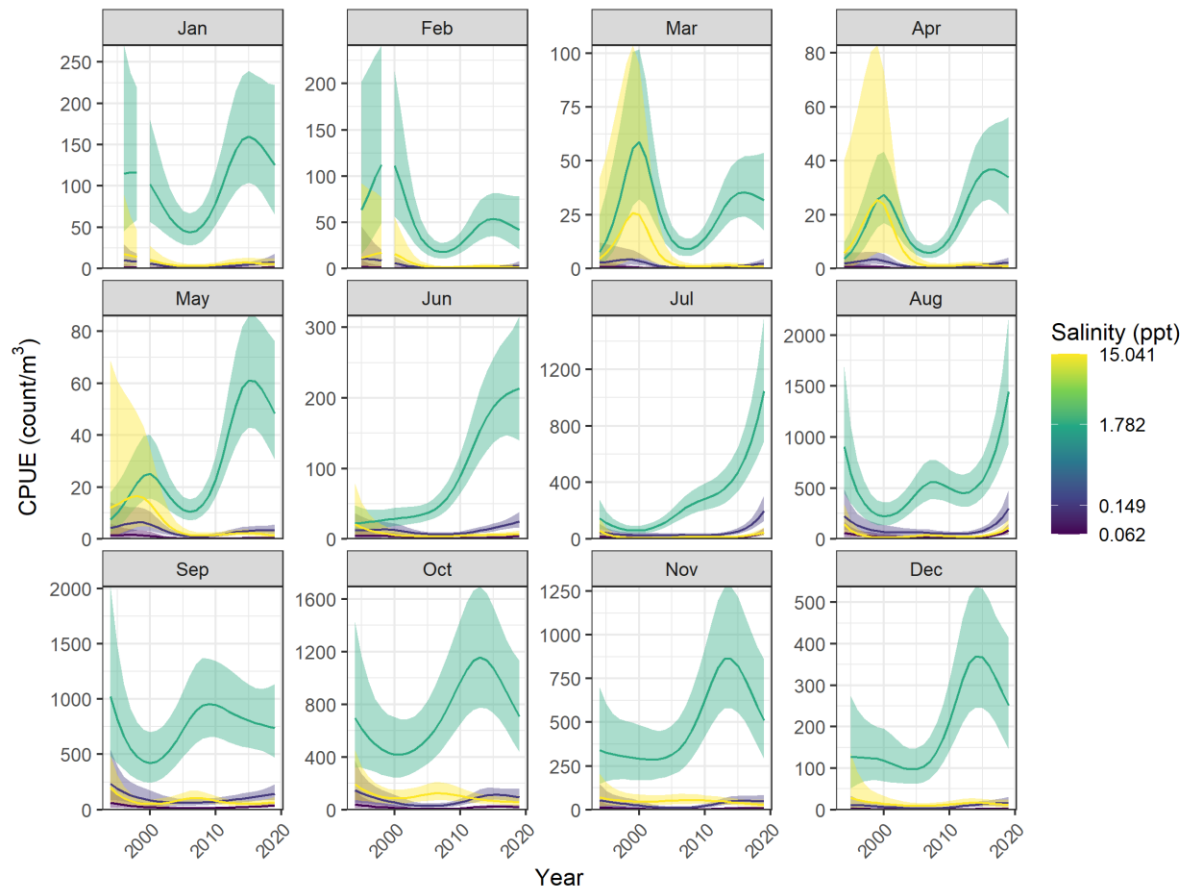
446 Spatially, *A. sinensis* adults had the highest abundance (controlling for all other
 447 covariates) along the corridor from the lower Sacramento River just below Cache Slough all
 448 the way through to Carquinez Strait. The Southeastern and Northern Delta had the lowest
 449 abundances (Fig. 11).



450

451 Figure 9. Seasonal patterns in *Acartiella sinensis* adult abundance with 99% credible intervals. See Fig. 3 for a

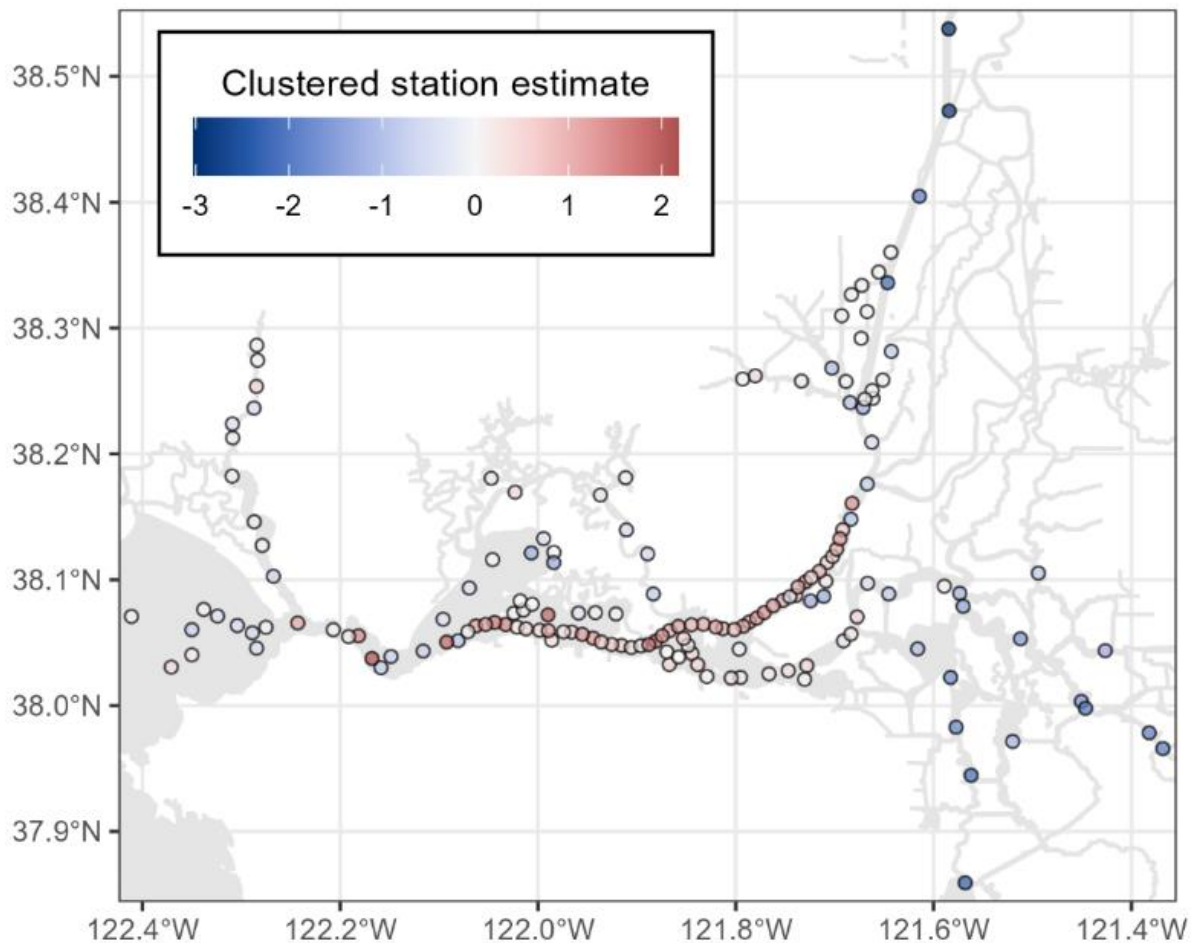
452 full description.



453

454 Figure 10. Yearly patterns in *Acartia sinensis* adult abundance with 99% credible intervals. See Fig. 4 for a

455 full description.



456

457 Figure 11. Estimated values of the clustered station random intercepts for *Acartia sinensis* adults. See Fig. 5
 458 for a full description.

459 *Acartia sinensis* juveniles

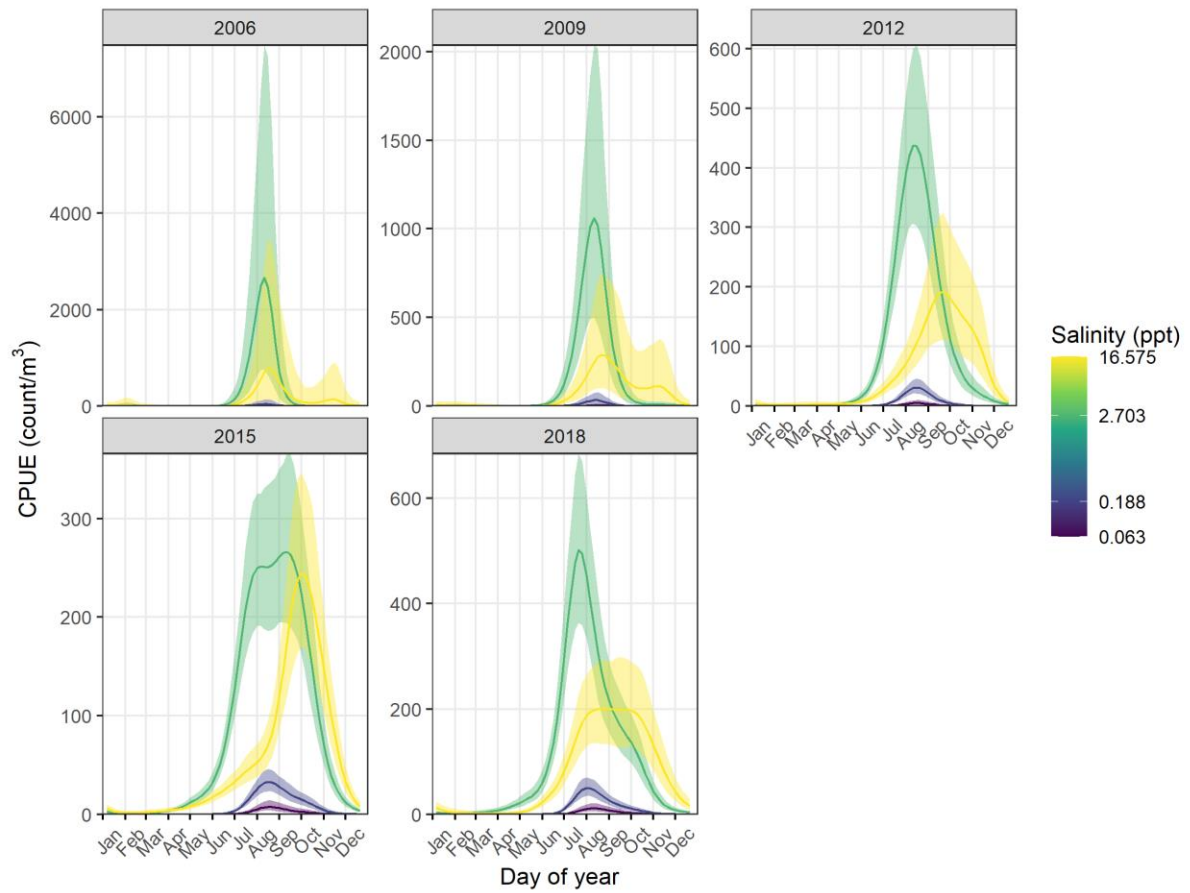
460 Like the adults, juvenile *A. sinensis* also had strong seasonal abundance patterns,
 461 peaking over just a few months and then subsiding to close to zero abundance. Peaks
 462 occurred in the summer from July through September, but the width of the seasonal peak
 463 grew over time. In 2006 they were abundant for just 2 months (July and August) while in
 464 2015 and 2018 they were abundant from April through November (Fig. 12).

465 *A. sinensis* juveniles were abundant in higher salinities > 4 but declined at the very
 466 highest salinities close to 16. Their abundance in lower salinities increased over time but
 467 always remained lower than the higher salinities (Fig. S4). In most years, the seasonal

468 abundance peak was 1-2 months later at the highest salinity of ~16 than the other salinity
469 levels (Fig. 12).

470 While the timeseries was much shorter (2006 – 2020) for *A. sinensis* juveniles than
471 any of the other species and life stages investigated, we did detect some long-term trends in a
472 few months. The trends were most apparent in the second highest salinity of 2.703 where
473 they were most abundant. Abundance increased over time in April through June and
474 decreased over time in August. This corresponds to the widening of the seasonal peak over
475 time. The other months generally did not have any significant long-term trends (Fig. 13).

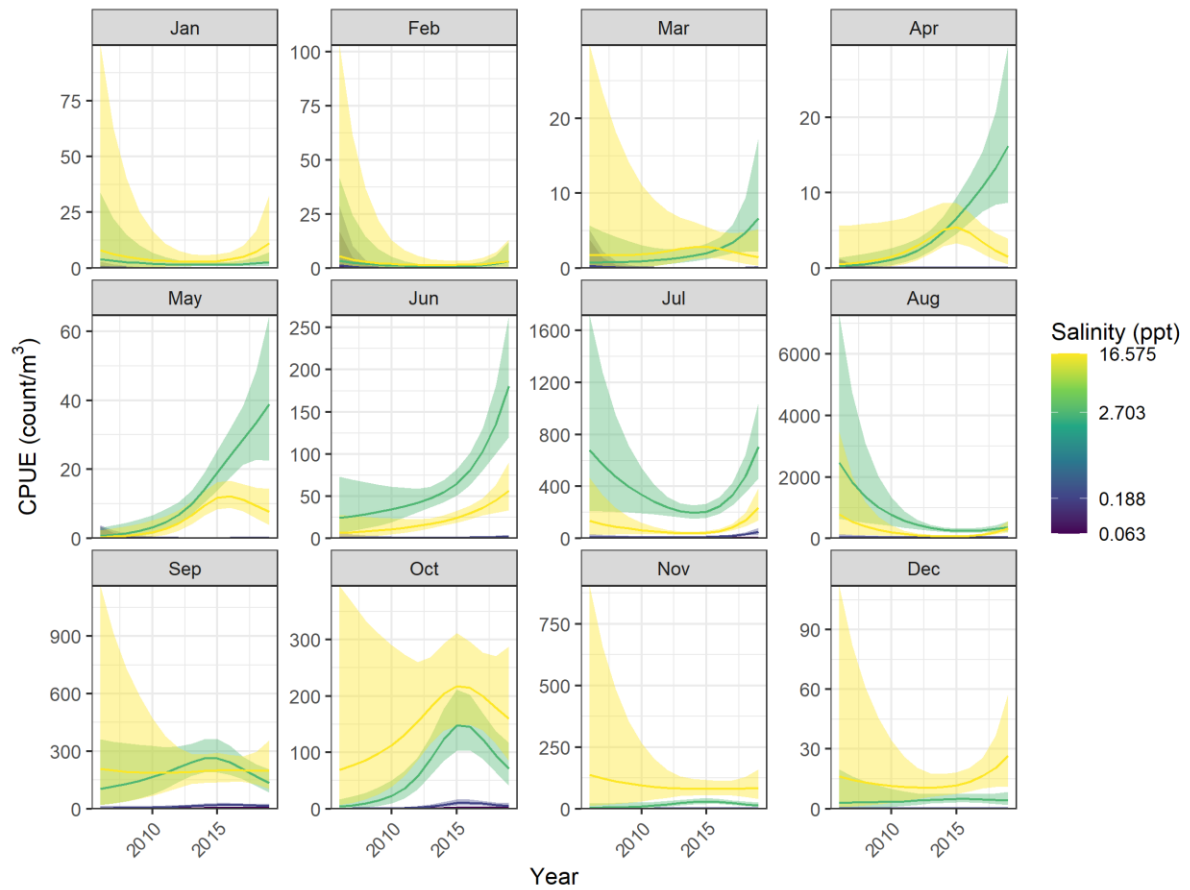
476 The spatial pattern of *A. sinensis* juveniles was less clear than the other species and
477 life stages. However, they were generally most abundant along the San Joaquin River
478 corridor in the Southern Delta and in some Suisun Bay stations. They were least abundant in
479 the lower Sacramento River between Cache Slough and the Confluence, as well as in the
480 Napa River and Eastern Suisun Marsh (Fig. 14).



481

482 Figure 12. Seasonal patterns in *Acartia sinensis* juvenile abundance with 99% credible intervals. See Fig. 3

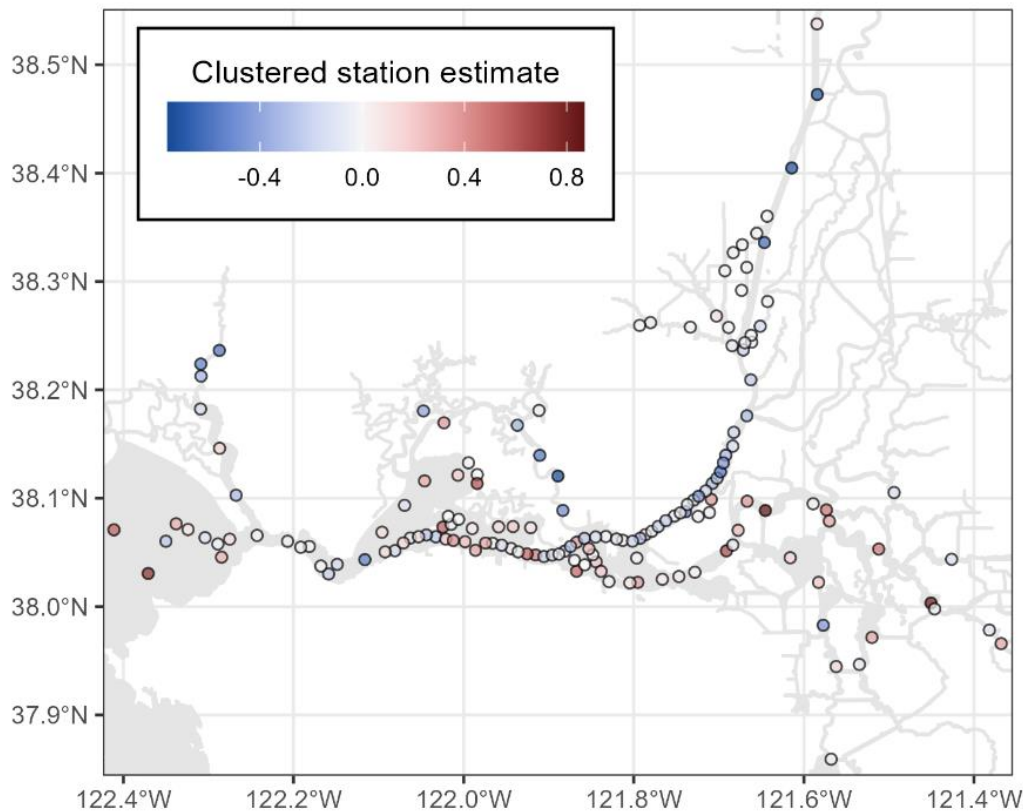
483 for a full description.



484

485 Figure 13. Yearly patterns in *Acartia sinensis* juvenile abundance with 99% credible intervals. See Fig. 4 for a

486 full description.



487

488 Figure 14. Estimated values of the clustered station random intercepts for *Acartia sinensis* juveniles. See Fig.
 489 5 for a full description.

490 Discussion

491 We found marked changes in the seasonality and overall abundance of three key
 492 zooplankton species in the upper estuary. *B. longirostris* no longer peaks in abundance in the
 493 fall months, *Acanthocyclops* spp. dramatically declined in all months and lost its strong
 494 relationship with salinity, and *A. sinensis* adult abundance has become increasingly driven by
 495 salinity while juveniles have developed wider abundance peaks. In this process, we have
 496 documented the relationship of each species with salinity and seasonality back to the
 497 beginning of monitoring or their introduction, increasing our understanding of their ecology
 498 and importance in the estuary.

499 Importance in fish diets

500 Of our study species, *Acanthocyclops* spp. was consumed by the most fish,
501 particularly Delta Smelt and Longfin Smelt. This is consistent with other studies that show
502 the importance of *Acanthocyclops* spp. in Longfin Smelt (Hobbs et al. 2006; Jungbluth et al.
503 2021), Delta Smelt (Lott 1998; Hammock et al. 2017), and larval Pacific Herring (Jungbluth
504 et al. 2021) diets. *Acanthocyclops* spp. has a similar fatty acid composition to the calanoid
505 copepod *E. affinis* (Kratina and Winder 2015), the preferred prey of Delta Smelt (Slater and
506 Baxter 2014), and Longfin Smelt (Barros et al. 2022; Burris et al. in review), and presents an
507 additional prey resource during the critical life stages of these fishes.

508 Adult *A. sinensis* were consumed the second most of our study species, but juveniles
509 were consumed the least. Since its introduction, *A. sinensis* has become important prey for
510 many fishes in the estuary during fall, especially Delta Smelt (Slater and Baxter 2014;
511 Hammock et al. 2017; Slater et al. 2019). Because *A. sinensis* is predatory, it has a lower
512 nutritional value (i.e. higher Carbon to Nitrogen ratio) than herbivorous copepods such as *P.*
513 *forbesi* (Kratina and Winder 2015), another common prey of Delta Smelt. *A. sinensis* predaes
514 and limits the availability of *P. forbesi* in the low salinity zone (Kayfetz and Kimmerer
515 2017), replacing *P. forbesi* as a prey source for juvenile and young-of-year Delta Smelt. *A.*
516 *sinensis* is not common in Longfin Smelt or Striped Bass diets, likely due to the peak
517 seasonal abundance occurring when both fish species are larger and have transitioned to
518 larger prey such as mysids and shrimp (Feyrer et al. 2003; Barros et al. 2022, CDFW
519 unpublished data).

520 *B. longirostris* were consumed the second least of our study species. Threadfin Shad
521 was the most frequent consumer of *B. longirostris* compared to the other fish species,
522 although the amount was still low despite their spatial overlap. Unlike most of the other
523 fishes examined, Threadfin Shad switch their feeding strategy from filter feeding to benthic

524 organisms and detritus based on prey availability (Ingram and Ziebell 1983) and this could
525 explain the low number of individuals that consumed *B. longirostris*. *B. longirostris*, and
526 other cladocerans, have a lower nutritional value than copepods (Kratina and Winder 2015).
527 However, cladocerans are still an important prey resource for fishes in the freshwater regions
528 in the estuary such as salmonids (Goertler et al. 2018; Aha et al. 2021), Mississippi
529 Silversides, and Threadfin Shad (Whitley and Bollens 2014), or during times when copepods
530 are less abundant such as for Delta Smelt in winter and spring (Hammock et al. 2017; Slater
531 et al. 2019).

532 Based on previous research (Lott 1998; Hobbs et al. 2006; Hammock et al. 2017;
533 Jungbluth et al. 2021), it is likely that the presence of our study species in fish diets is higher
534 than our data indicates. This could be due to damage during digestion hindering identification
535 of prey to the taxonomic level of our study taxa, or that most fish samples were limited to
536 specific seasons and may not have been collected at a time that our zooplankton species were
537 present in the environment. Thus, our results likely represent undercounts of the actual
538 abundance of our study taxa in fish diets.

539 Seasonal, salinity-related, and geographic abundance patterns

540 Currently, *B. longirostris* and *Acanthocyclops* spp. adults peak in the spring while *A.*
541 *sinensis* adults peak in the fall and juveniles peak in the summer. The spring peaks line up
542 with the spawning of Delta Smelt while summer and fall peaks provide critical food for Delta
543 Smelt juveniles and young-of-the-year (Slater and Baxter 2014; Slater et al. 2019). The
544 spring peaks also correspond to the outmigration of juvenile salmon and could provide
545 important food necessary to increase growth rates and reduce oceanic predation risks
546 (Herbold et al. 2018; Phillis et al. 2018; Zeug et al. 2019).

547 *B. longirostris* and *Acanthocyclops* spp. are also both most abundant in the lowest
548 salinities, although *Acanthocyclops* spp. has a broader salinity range likely due to the
549 different tolerances of the species in the complex. Freshwater habitat, especially in the spring,
550 is where spawning for most native fish species occurs, including Delta Smelt and Sacramento
551 Splittail (Moyle 2002). *A. sinensis* peaks in more saline water corresponding to the low
552 salinity zone (generally defined as 0.5 to 6 ppt), which is a key habitat for rearing Delta Smelt
553 (Sommer and Mejia 2013). The juveniles have a smaller salinity range and are abundant in
554 more saline waters than the adults. The brackish and low salinity habitats are important
555 rearing habitats for many native fishes that evolved in highly variable conditions, giving
556 native fishes an advantage over non-native fishes (Moyle et al. 2010). Both *Acanthocyclops*
557 spp. and *A. sinensis* are found in the diets of Longfin Smelt, which spawn and rear at slightly
558 higher salinities than many other native fishes (Hobbs et al. 2006; Grimaldo et al. 2017;
559 Jungbluth et al. 2021).

560 Adding to the pattern of similarity, *B. longirostris* and *Acanthocyclops* spp. had
561 similar geographic patterns, with the highest abundance (controlling for other covariates) in
562 the Cache Slough Complex and the southeastern and eastern boundaries of the study region.
563 *B. longirostris* especially seemed to peak in areas of high residence time such as the
564 northernmost location on the Sacramento Ship Channel and areas in the East Delta (Vroom et
565 al. 2017; Lenocho et al. 2021). The Sacramento Ship channel is an important last refuge for
566 Delta Smelt and other fishes (Young et al. 2021). *Acanthocyclops* spp. had very high
567 geographic peaks in Suisun Marsh and Cache Slough, both areas with remnant and restored
568 tidal wetlands that are important habitats for native fishes (Kimmerer et al. 2018; Colombano
569 et al. 2020). They both also had generally low abundance from Suisun Bay upstream through
570 the lower Sacramento and San Joaquin Rivers, which was the region of highest abundance for
571 *A. sinensis* adults. *A. sinensis* adults had low abundance in the southeastern and northern

572 regions. *A. sinensis* juveniles had some similarities to the adults but were much more
573 abundant along the San Joaquin River corridor and had low abundance along the Sacramento
574 River. They also had areas of high abundance in San Pablo Bay.

575 Long term changes

576 While *B. longirostris* and *Acanthocyclops* spp. have experienced overall declines in
577 abundance over time, *A. sinensis* has mostly increased, although over a shorter time period.
578 The declines of *B. longirostris* and *Acanthocyclops* correspond to noted regime shifts and
579 overall plankton declines across many species (Winder and Jassby 2011). The increase of *A.*
580 *sinensis* could be related to expansion following its recent invasion as it fills niches left by
581 declining species.

582 The change to the seasonal pattern of *B. longirostris* abundance was puzzling. Prior to
583 1990, *Bosmina* experienced two peaks, one in the spring and a second peak in the fall.
584 Something happened in the late 1980s to cause a crash in *B. longirostris* abundance during
585 the fall. This fall peak may have been an important food source for juvenile Delta Smelt.
586 While we may never be able to tell exactly why this peak disappeared, we discuss three
587 theories here: changes to water management, invasive clams, and invasive copepods.
588 Operation of the State Water Project and Central Valley project can significantly change
589 flows in the San Joaquin River and the channels of the South Delta, where *Bosmina* is most
590 abundant (Jassby et al. 2002; Jassby 2005). Exports from these water projects cause a
591 decrease in residence time in the South Delta, particularly during the fall peak of *B.*
592 *longirostris* (Hammock et al. 2019). Decreased residence time limits phytoplankton
593 production, as well as directly exporting phytoplankton and zooplankton (Jassby et al. 2002;
594 Hammock et al. 2019). However, this pattern first became apparent during increases to
595 exports in the late 1970s (Hammock et al. 2019), well before the disappearance of the fall

596 peak of *B. longirostris*, so the export explanation is unlikely to be the main factor driving the
597 decrease, though it could be part of the story.

598 Invasive species may be a more likely explanation for the change in the seasonal
599 peaks of *B. longirostris*. The entire zooplankton community experienced a decline in
600 abundance in the western portions of the estuary due, in part, to the introduction of the
601 overbite clam (*Potamocorbula amurensis*) in 1986 (Kimmerer and Orsi 1996). The overbite
602 clam has an extremely high filtration rate, greater than any native benthic grazer, and caused
603 a crash in chlorophyll-a in Suisun Marsh and Suisun Bay. Zooplankton suffered from loss of
604 food resources as well as directly through grazing on zooplankton nauplii (Kimmerer and
605 Thompson 2014; Kimmerer and Lougee 2015). Grazing rates of overbite clams peak in
606 summer and fall as biomass increases and higher water temperatures cause increases in
607 metabolic rates (Crauder et al. 2016), so they would be well positioned to reduce the *B.*
608 *longirostris* fall peak without impacting the spring peak. Could this have caused a reduction
609 in fall *B. longirostris*? We feel this is unlikely to be the main cause, because *B. longirostris*
610 chiefly occurs in the freshwater reaches of the Delta, whereas the overbite clam occurs
611 mainly in areas with salinity > 2ppt (Crauder et al. 2016). While clams may not be the
612 obvious answer, there was another important species introduction in the late 1980s with
613 greater potential to directly impact *B. longirostris*. The calanoid copepod *P. forbesi* was
614 introduced in 1987 and quickly became the most abundant calanoid in the system (Orsi and
615 Walter 1991). *P. forbesi* peaks in abundance from July through October, overlapping with the
616 historic peak in *B. longirostris* abundance, and they occur in high abundances in the South
617 Delta where *B. longirostris* is common (Kayfetz and Kimmerer 2017). *P. forbesi* may be
618 competing with *B. longirostris* for food resources during the fall when previously most other
619 zooplankton peaked earlier in the year.

620 The dramatic decline of *Acanthocyclops* spp. abundance may be related to the
621 invasion of the cyclopoid *Limnoithona tetraspina* in 1993 (Orsi and Ohtsuka 1999). Before
622 this invasion, *Acanthocyclops* spp. was the most abundant cyclopoid in the region (Orsi and
623 Mecum 1986). After 1994, *L. tetraspina* quickly dominated the copepod community
624 (Hennessy 2018) with *Acanthocyclops* spp. averaging ~1% of the abundance of *L. tetraspina*
625 (Bashevkin et al. 2020). The invasion of *L. tetraspina* could have facilitated the decline of
626 *Acanthocyclops* spp. either through direct competition or through facilitation of a common
627 predator, *A. sinensis*, as it likely did for *P. forbesi* (Kayfetz and Kimmerer 2017).

628 *Acanthocyclops* spp. also exhibited a reduction in their relationship to salinity over
629 time. This is likely due to the differing salinity tolerances of the species within the complex
630 and changes in the relative abundances of those species over time. Unfortunately, we do not
631 have data on the abundances of each species in the complex so we cannot untangle the
632 individual patterns. *A. vernalis* has been described as native to the estuary (Orsi and Mecum
633 1986; Kratina and Winder 2015). However, since the presence of the *A. vernalis* species
634 complex in the estuary has only recently been confirmed (Jungbluth et al 2021), it is
635 unknown which species could be native, or if some were introduced during the study period.
636 The strong relationship with salinity in early years (Fig. S2) may be indicative of one or more
637 species with lower salinity tolerances initially dominating, then later an invasion and
638 expansion of the higher salinity species such as *A. americanus* which has been found in high
639 salinity areas including the Mediterranean Sea (Alekseev 2021). The invasion of the
640 predatory *A. sinensis* in more saline areas, as well as the shift in phytoplankton communities
641 caused by invasion of *Potamocorbula amurensis* (Lucas et al. 2016), could also have
642 contributed to this shift by taking over from salinity as the prime limiting abundance factor.

643 Adult *A. sinensis* were increasingly driven by salinity over time. In the earlier years,
644 they were present in all salinities in our study and even equally abundant in high salinities of

645 15 and low salinities of 0.15. They also exhibited a unique winter-spring abundance peak in
646 the two highest salinities that disappeared by 2005. They invaded in 1993 (Orsi and Ohtsuka
647 1999), thus this pattern could reflect them settling into their ecological niche over time.
648 Another predatory copepod, *Tortanus dextrilobatus*, was introduced in 1994 and is found in
649 higher salinities around 13- 19 ppt (Orsi and Ohtsuka 1999; Bollens et al. 2002). *T.*
650 *dextrilobatus* typically peaks in summer (Bollens et al. 2014); however, when first introduced
651 it also peaked in March and April (Hooff and Bollens 2004). Potential competition from *T.*
652 *dextrilobatus* could thus have contributed to adult *A. sinensis* losing its spring peak in those
653 higher salinities.

654 Interestingly, *A. sinensis* juveniles had increasingly wide seasonal abundance peaks
655 over time, driven in part by differing timing of the two highest salinity bins. Abundance
656 peaks were regularly 1-2 months later in the highest salinity (16.575) than in any of the lower
657 salinities, which all peaked around the same time. The abundance peak of the highest salinity
658 also grew relatively larger compared to the lower salinities over time, which led to an overall
659 widening of the seasonal abundance peak for *A. sinensis* juveniles. However, the width of the
660 abundance peak in each lower salinity level also seemed to widen over time. This
661 demonstrates shifting phenology of *A. sinensis*, which could be caused by changes in the
662 timing and location of reproduction, predation, or feeding. The zooplankton community has
663 undergone many shifts over the history of this dataset (Orsi and Ohtsuka 1999; Winder and
664 Jassby 2011), with *A. sinensis* potentially having its own impacts on lower trophic level
665 zooplankton (Kayfetz and Kimmerer 2017). Since *A. sinensis* is predatory it could be
666 following the abundance shifts of other species, resulting in changes to the location and
667 timing of reproduction, thus impacting both adult and juvenile abundances. Alternatively, this
668 shifted phenology could be caused by changes to the salinity field in recent years which have

669 resulted in higher salinity intrusions further into the estuary due to increased and persisting
670 droughts (Ghalambor et al. 2021).

671 Conclusions

672 Many of the fishes in the estuary rely on zooplankton for at least part of their life
673 cycle. Changes in prey resources can affect higher trophic levels by reducing the amount of
674 available food or shifting the timing of peak abundance, thereby creating a mismatch between
675 critical fish life stages and their prey. We found long-term shifts in all three of our study
676 species. These shifts included changes in seasonality, salinity preference, and long-term
677 abundance. Further studies investigating these patterns in additional species would be
678 important to understand the past dynamics of zooplankton in the estuary. These results
679 increase our understanding of the zooplankton community, which could inform the
680 development of food web models and be matched to trends in fish abundance to examine the
681 direct influence of declining zooplankton species on managed species.

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