

Long-term trends in seasonal plankton dynamics in Lake Mead (Nevada-Arizona, USA) and implications for climate change

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Abstract Over the past few decades, Lake Mead (Nevada-Arizona, USA) has experienced multiple ecosystem stressors including drought, increased water demand, and establishment of invasive species (quagga mussels, gizzard shad). Despite these potential stressors, zooplankton and phytoplankton community dynamics in the pelagic regions of Lake Mead have generally followed consistent seasonal patterns. Long-term monitoring results (2007–2015) show that zooplankton and phytoplankton communities remain relatively stable in Lake Mead on an inter-annual basis, but are susceptible to shifts caused by extreme

climate fluctuations and alterations in mixing regimes. A warm winter characterized by low snowpack in 2014/2015 preceded a large bloom of toxic cyanobacteria (*Microcystis aeruginosa* Kützing) in Las Vegas Bay the following summer. Large bloom events are rare in Lake Mead; however, under future climate scenarios, these types of events may become more frequent. Because of the consistency of plankton community dynamics over an extended period of time, Lake Mead offers an ideal system for the study of future climate change impacts. This study aims to characterize the response of plankton communities in Lake Mead to both linear and dynamic environmental changes.

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Introduction

Lake Mead, located along the lower Colorado River (Nevada-Arizona, U.S.A.), is the largest reservoir by volume within the U.S. (Holdren & Turner, 2010). Over the last few decades, Lake Mead has been experiencing ongoing threats to water quality from multiple compounding factors. Since 2000, drought conditions have persisted in reservoirs throughout the southwestern U.S. Average water stage has fallen over 40 m in Lake Mead and inflows have been reduced by

over 3,000 Mm³/year (Moreo & Swancar, 2013). Invasive quagga mussels (*Dreissena bugensis* Andrusov, 1897) were first discovered in Lake Mead in 2007 (LaBounty & Roefer, 2007), but likely became established in the lower Colorado River ca. 2003 (McMahon, 2007). Since introduction, quagga mussels have spread throughout Lake Mead and downstream reservoirs (Wong et al., 2010). Other factors that may be impacting the ecosystem and water quality in Lake Mead include establishment of gizzard shad [*Dorosoma cepedianum* (Lesueur, 1818)] and rapid population growth in the city of Las Vegas and surrounding area, contributing to significantly increased treated wastewater inflow from the Las Vegas Wash (Holdren & Turner, 2010). It is unclear to what extent climate change might impact Lake Mead and other Colorado River reservoirs, although the majority of models predict declines in average annual river discharge over the next 50 years (Rajagopalan et al., 2009).

Decreases in reservoir water levels pose unknown threats to the ecosystem, and the long-term ecological impacts of invasive dreissenid mussels on southwestern reservoirs are still unclear. Drought conditions have the potential to negatively impact overall water quality in lakes and reservoirs via reduced inflows and outflows, warmer surface temperatures and prolonged stratification, increased water residence time, increased nutrient concentrations, and increased frequency of harmful algal blooms (Mosley, 2015). Murdoch et al. (2000) predict that the combined effects of prolonged drought and warming air temperatures will increase the residence time of point-source nutrients and pollutants in lakes and reservoirs as a result of decreased inflow and outflow. In the Great Lakes region of the U.S., dreissenid mussels have caused significant shifts in water quality and ecosystem function by increasing water clarity and reducing phytoplankton populations (Caraco et al., 1997; Fahnenstiel et al., 2010), in turn reducing populations at higher trophic levels (zooplankton and fish) (Bridgeman et al., 1995; MacIsaac, 1996; Pace et al., 1998).

Despite the potential impacts of quagga mussels on plankton ecology in invaded lakes and reservoirs, little evidence exists demonstrating significant changes in either zooplankton or phytoplankton populations in response to the invasion in Lake Mead (Beaver et al., 2010; Wong et al., 2010; Turkett, 2016). Drought impacts on zooplankton appear to be minimal, as long-

term trends in zooplankton community composition follow regular seasonal patterns closely tied to temperature and mixing cues (Beaver et al., 2010). The results of these studies show that the established food web in this system has remained essentially unaltered a decade post-quagga invasion and drought initiation. However, previous studies in many different types of ecosystems have indicated that response of individual species to gradually increasing ecological stressors may not be linear (Scheffer et al., 2001). Many studies have shown that loss of resilience in lake ecosystems can go undetected for several years until a critical threshold of change occurs, which may be induced via abnormal climatic, biological, or hydrological events (Scheffer et al., 2001; Zambrano et al., 2001; Ji et al., 2018). This type of catastrophic change often requires extensive restoration efforts to re-establish biological communities (Jeppesen et al., 2007; Ji et al., 2018). In 2015, warm temperatures reduced winter snowpack and increased thermal stratification during a period of traditional mixing and were followed by a large, toxic bloom of *Microcystis aeruginosa* Kützing in Lake Mead. The compounding effects of low water elevation due to reduced inflows (which in turn reduces wind-driven mixing, as surface waters are more protected by heightened canyon walls), earlier onset of seasonal stratification and increased surface water temperatures provided environmental conditions that have elsewhere led to blooms of *M. aeruginosa* (Jacoby et al., 2000; Cao et al., 2006). Cyanobacterial blooms tend to thrive under warm, highly stratified conditions (Paerl & Huisman, 2008), and can lead to production of a suite of potent toxins that threaten humans and wildlife. Although this type of toxic bloom event has been rare amongst Colorado River reservoirs thus far, it may be a foreshadowing of an increasing frequency of such events to come under warming climate conditions, suggesting that Lake Mead may be vulnerable to major ecological disruptions in the future.

This study examines seasonal and inter-annual dynamics of zooplankton and phytoplankton in open water regions of Lake Mead over a nine-year period following the introduction of quagga mussels (2007–2015), including an abnormally warm year that resulted in a large bloom of toxic cyanobacteria. The relationship between zooplankton biomass and water temperature, as well as phytoplankton biovolume and water temperature, was explored using quadratic

regression and niche centroid analysis in order to determine optimal temperature conditions for dominant species. Canonical correlation analysis (CCA) was then used to determine how fluctuations in water quality variables may affect relative biomass or biovolume of individual taxa. Recently, Beaver et al. (2010) documented a high degree of regularity in seasonal patterns of zooplankton abundance and community composition between 2000 and 2009. Given the long-term duration of unchanging plankton community dynamics under increasing ecological stress, along with the 2015 bloom event that coincided with an extreme climate fluctuation, we suggest that plankton in Lake Mead may be experiencing a threshold, rather than linear response to environmental change. Since inter-annual variability has been relatively low thus far, Lake Mead represents an excellent system for the study of climate change impacts in both the near-term and long-term future. Results underscore the importance of the linkages amongst hydrology, nutrients, phytoplankton, and higher trophic levels in Colorado River reservoirs, and highlight the potential impacts of climate change on these important ecosystems.

Materials and methods

Study sites

Lake Mead, a subtropical monomictic system, is impounded by Hoover Dam, which supplies power generation to a large portion of the southwestern United States. It is an economically important reservoir as it provides municipal water for the City of Las Vegas and the surrounding area, supplying about 90% of municipal water needs in the Las Vegas Valley. Releases from Hoover Dam are critical for water supply to downstream reservoirs such as Lake Mojave (Nevada-Arizona) and Lake Havasu (Arizona-California). The Lake Mead National Recreation Area is one of the largest in the country, with a public launch capacity for over 3,000 boats (Holdren & Turner, 2010). Lake Mead also serves as habitat for several varieties of sportfish, as well as the endangered razorback sucker [*Xyrauchen texanus* (C.C. Abbott, 1860)], native to the Colorado River. Major tributaries of Lake Mead are the Colorado River (97% of inflow), Virgin River, and the Las Vegas Wash. Varied water

inflows from each tributary result in three major, ecologically unique regions of Lake Mead: Las Vegas Bay, Overton Arm, and Colorado River. Las Vegas Bay, the most downstream basin, collects the combined flows from the Overton Arm, the Las Vegas Wash and the main stem of the Colorado River and consists of four sampling stations (LVB sites, Fig. 1). Overton Arm collects the combined drainages of the Virgin and Muddy Rivers and consists of six sampling stations (VR sites). The Colorado River enters Lake Mead at the northern end, nearly 97 km upstream of Hoover Dam, and consists of seven sampling stations (CR sites). Samples from each station in Las Vegas Bay, Overton Arm, and the Colorado River were collected quarterly (winter, spring, summer, fall) from January 2007 through November 2015 by the Bureau of Reclamation Resources Management Office in Boulder City, Nevada.

Sample collection/laboratory analyses

Water quality

Water quality profiles were conducted along the thalweg of the Colorado River (site CR394.0 to CR342.5) from the bottom to the surface of each sampling station using a multi-parameter submersible sonde unit (Manta Multiprobe, Eureka, Austin) capable of measuring depth, dissolved oxygen concentration, pH, electrical conductivity, and temperature. The bottom depth for each station was recorded as the maximum station depth, and temperature data were recorded in 5-m intervals from the bottom depth to 50 m, followed by 2-m intervals from 50 to 20 m, and finally in 1-m intervals from 20 m to the surface. Thermal profiles of Lake Mead were produced using Surfer[®] 12 (Golden Software, LLC). For the purpose of this study, temperature data from the top five meters of each station were used for statistical analyses. Secchi depth was determined using a viewscope and a 20-cm disc. Water samples for total phosphorus determination were collected as a composite of the 0–5 m strata using a pool hose, and at the 20, 100 m (when applicable), and at the maximum depth using a Kemmerer sample bottle. All samples were kept on ice and shipped overnight to Sierra Foothill Laboratory, a certified and accredited laboratory in the state of California, for analysis. Samples to be analyzed for total phosphate as phosphorus (P) were immediately

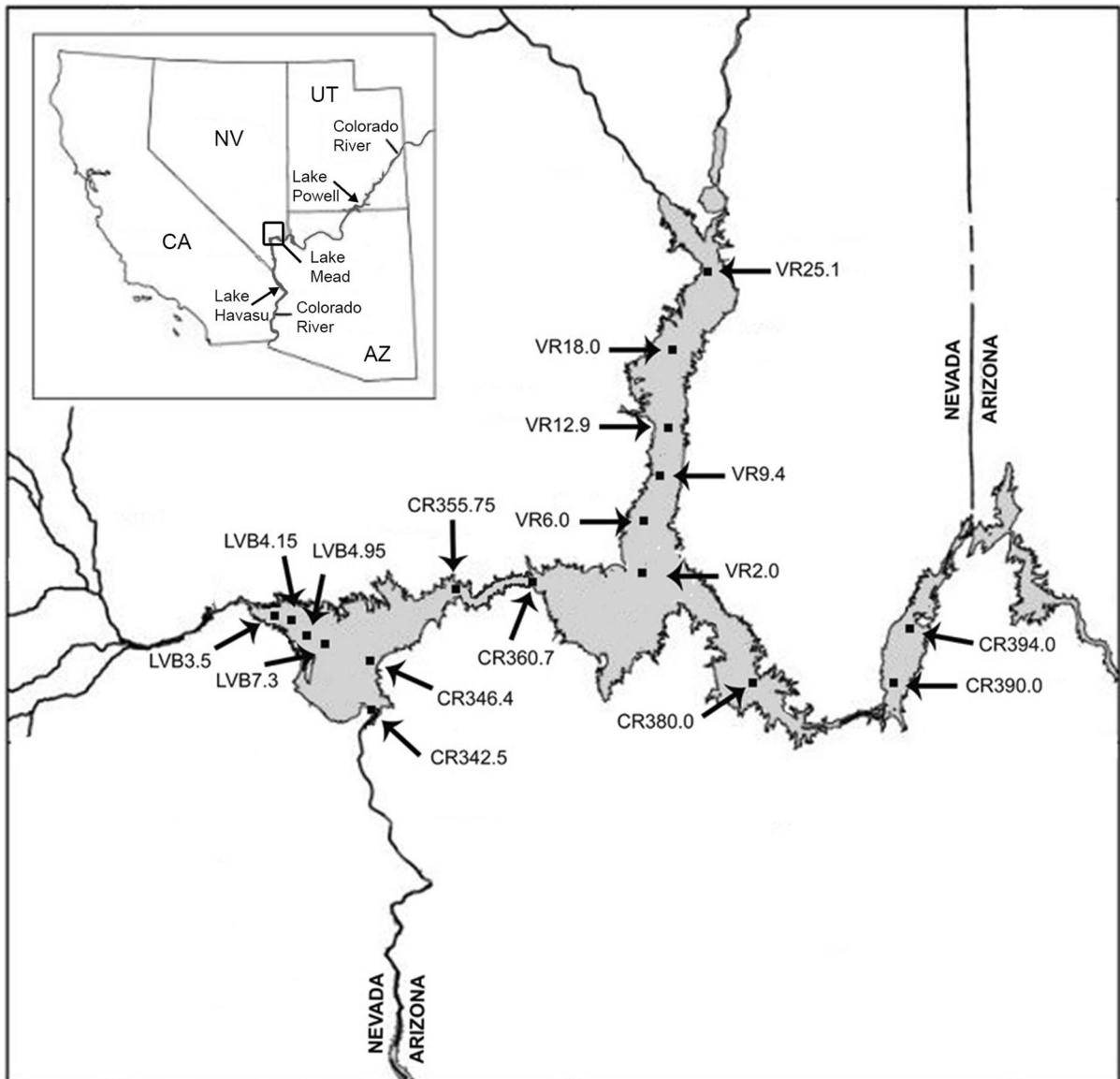


Fig. 1 Locations of sites sampled in Lake Mead. LVB sites represent the Las Vegas Bay, VR (Virgin River) sites represent the Overton Arm, and CR (Colorado River) sites represent the Colorado River. Numbers following site designations represent river miles

acidified in the field and digested and analyzed using EPA method SM4500-P,E, with a reporting limit of 0.002 mg/l as P.

Zooplankton

Water samples collected for zooplankton analysis from 2007 to 2015 were obtained from the top 0–30 m (or identified maximum depth) using a 64- μ m mesh Wisconsin-style plankton net. The net was washed

into 250-ml Nalgene bottles and preserved with 2% Lugol's solution, and promptly shipped on ice to BSA Environmental Services, Inc. Upon arrival at the laboratory, sample volumes were measured and homogenized aliquots examined on Wilovert inverted microscopes at $\times 100$ magnification (up to $\times 1000$ if necessary). All samples were analyzed using the Utermöhl sedimentation technique. A minimum tally of 200 organisms per sample was reached (or the entire sample analyzed), and specimens were identified to

the lowest taxonomic level possible. For each identified taxa, up to 10 individual specimen length and width measurements were recorded (or less for more rare taxa) for the calculation of biomass estimates based on established length/width relationships (Dumont et al., 1975; McCauley, 1984; Lawrence et al., 1987). Zooplankton carbon content was assumed to be 50% of dry mass (Latja & Salonen, 1978).

Phytoplankton

Phytoplankton samples were collected as 0–5 m composites using a pool hose and stored in 250-ml Nalgene bottles. Each sample was immediately preserved with 2% Lugol's solution and iced in the field, then shipped to BSA Environmental Services, Inc. for laboratory analysis. Phytoplankton slides were prepared using standard membrane filtration technique in order to preserve cell structure for examination at high magnification (McNabb, 1960).

Filtered phytoplankton samples were analyzed on Leica DMLB compound microscopes at $\times 630$ magnification for random field and full strip counts, as well as $\times 400$ magnification for half-filter counts to further ensure complete species reporting. As per Lund et al. (1958), a minimum of 400 natural units (colonies, filaments, unicells) were enumerated to the lowest possible taxonomic level from each sample to ensure accuracy within 90% confidence limits. Cell biovolumes, based on measurements of 10 organisms per taxon when possible, were estimated for all identified phytoplankton taxa using formulae for solid geometric shapes that most closely match cell shape (Hillebrand et al., 1999). PCR was performed following microscopic classification to confirm the identification of *Microcystis aeruginosa* (Nübel et al., 1997), a known toxin-producing species (Chorus & Bartram, 1999). Phytoplankton carbon content was estimated as described in Rocha & Duncan (1985). C-phytoplankton was divided by 10^6 in order to achieve comparable units with C-zooplankton ($\mu\text{g/l}$) and carbon ratios (C-zooplankton:C-phytoplankton) were calculated using whole-lake means for each sampling event. Mean carbon ratios were plotted against whole-lake mean surface temperatures, and a regression equation that best fit the data, along with the associated coefficient of determination, was calculated using Microsoft Excel.

Algal toxins

Whole water grab samples for toxin analysis were collected in August, 2015 between 0 and 5 m depth from 19 open water sites throughout the reservoir, as well as two near-shore sites and one beach site. Samples were collected with buffer (TRIZMA, 2-chloroacetamide), held on ice during transport to Texas A&M University Corpus Christi, and processed within 6 h of receipt. 60-ml subsamples were concentrated on SPE C18 cartridges, then eluted with 1 ml 75% MeOH and the elutes were analyzed for the detection of Microcystin-LR using HPLC/MS following EPA method 544.

Temperature intervals and niche centroids

The relationship between water temperature and daphnid biomass was examined using both a quadratic regression model and niche centroid analysis. Spring and summer (May and August 2007–2015) samples were used for the quadratic regression model. Mean biomass values were calculated for both *Daphnia pulex* complex and *Daphnia galeata mendotae* Birge, 1918 for each respective water temperature interval: 16–18, 18–20, 20–22, 22–24, 24–26, 26–28, 28–30, and 30–32°C. All sampling points with temperatures that fell within an interval range were used to calculate biomass means. An equation that best fits the data and the associated coefficient of determination were calculated using Microsoft Excel.

Niche centroid analysis provides a weighted average of an environmental variable (in this case, surface water temperature) in sites at which an individual taxon occurs. Niche centroid temperatures calculated for different taxa can be compared and interpreted as the “optimum” temperature at which that taxon is likely to experience highest biomass or biovolume (infer maximal growth). The relationship between water temperature and biomass/biovolume was quantified using the equation described in ter Braak and Verdonschot (1995):

$$u_k = \sum_{i=1}^n \frac{y_{ik}}{y_{+k}} x_i,$$

where y_{ik} is the density of species k in the i th sample; y_{+k} is the summed biomass/biovolume of species k in all samples, and x_i is the water temperature in the i th sample.

For the niche centroid analysis, *Daphnia pulex* complex and *Daphnia galeata mendotae* biomass were individually analyzed, as well as *Cyclotella* spp., *Rhodomonas* spp., and *Microcystis* spp. biovolume.

Statistical analyses

Zooplankton

34 zooplankton taxa identified from samples collected between January 2007 and November 2015 were used in Canonical Correlation Analyses (CCA) using PRIMER 6 (Clarke & Gorley, 2006). Samples with missing or unavailable water quality data were excluded from the analysis, resulting in 587 individual samples. The zooplankton data set was standardized and a Principal Component Analysis (PCA) was performed, which identified a few statistically significant species that explained a large percentage of the total variance within the data set (*D. pulex* complex and *D. galeata mendotae*). Subsequently, a Bray–Curtis resemblance matrix of the standardized zooplankton biomass data was used in a CCA (using the Canonical Analysis of Principal Coordinates (CAP) function of the PERMANOVA+ add-on in PRIMER 6) with a sample-matched, normalized matrix of 4 water quality variables: water temperature, Secchi depth, station depth, and total phosphorus. Following the CCA, samples were plotted on ordination diagrams using the first two CAP axes, and the biomass values of *D. pulex* complex and *D. galeata mendotae* were superimposed over the CCA plot.

Phytoplankton

Prior to analyses, phytoplankton biovolume data were reduced to include only the taxa that constituted greater than or equal to 5% of the total sample biovolume for at least one sample, resulting in 79 taxa. A PCA on the standardized phytoplankton biovolume data set identified *Rhodomonas* spp. and *Cyclotella* spp. as significant genera contributing to explained variability. Both genera, as well as *Microcystis* spp.,

were then superimposed over the zooplankton CCA ordination diagrams.

Results

Water quality

The Colorado River was the deepest, clearest region of Lake Mead with a mean station depth of 89.4 m and mean Secchi depth of 9.5 m (Table 1). The lowest mean depth and clarity was observed in Las Vegas Bay (41.9 and 6.5 m, respectively). In contrast, Las Vegas Bay was the warmest region with a mean temperature of 20.7°C, and also exhibited the highest mean phosphorus concentration (14.2 µg/l). Temperature and total phosphorus concentrations in Overton Arm and Colorado River displayed narrow differences (20.0, 20.1°C and 7.1, 5.6 µg/l, respectively). Annual mean winter water temperatures from the top 5 m of the water column support the shifting temperature–density pattern observed in the winter for Las Vegas Bay, Overton Arm, and Colorado River regions of Lake Mead (Table 2, Fig. 2). Differences in winter water temperatures from all three basins were determined to be statistically significant ($P < 0.05$, two sample t test assuming equal variance).

Thermal profiles

Thermal profiles of the Colorado River thalweg from February 2012–2015 (Fig. 3) display the shifting temperature–density relationship in Lake Mead, which has a water column that has historically been either completely or partially mixed in winter. In February 2012, relative uniformity in temperature at the downstream end of the lake (near station CR342.5) was observed, ranging from about 12.5°C at the surface of the lake to 9.5°C at the lake bed (Fig. 3A). Similar uniformity was observed in February 2013 and February 2014 (Fig. 3B, C). The thermal profile from February 2015 (Fig. 3D) portrayed a strongly stratified water column that has maintained a distinct epilimnion (surface water temperatures approximately 13–16°C above 30 m depth) and hypolimnion (water temperatures approximately 12–13°C below 30 m depth). Thermal profiles from station CR346.4 during winter sampling from 2007 to 2015 also show consistency in winter mixing regimes from 2007 to 2014 (Fig. 4). In

Table 1 Water quality characteristics (\pm SE) for the Las Vegas Bay, Overton Arm, and Colorado River regions of Lake Mead

	Temperature ($^{\circ}$ C)	Secchi depth (m)	Station depth (m)	Total P (μ g/l)
Las Vegas Bay	20.7 (\pm 0.5)	6.5 (\pm 0.4)	41.9 (\pm 2.4)	14.2 (\pm 1.1)
Overton Arm	20.0 (\pm 0.4)	8.0 (\pm 0.3)	55.8 (\pm 2.0)	7.1 (\pm 0.6)
Colorado River	20.1 (\pm 0.4)	9.5 (\pm 0.3)	89.4 (\pm 1.7)	5.6 (\pm 0.3)

Values represent the mean of all data for 2007–2015

Table 2 Annual mean winter water temperatures (0–5 m, $^{\circ}$ C) for the Las Vegas Bay, Overton Arm, and Colorado River regions of Lake Mead

	Las Vegas Bay	Overton Arm	Colorado River
Win-07	12.1	11.3	11.7
Win-08	13.0	11.6	12.0
Win-09	13.7	12.4	12.6
Win-10	13.2	12.2	12.6
Win-11	12.8	11.4	12.0
Win-12	13.2	12.2	12.8
Win-13	12.6	11.8	12.2
Win-14	13.1	12.1	12.7
Win-15	15.5*	13.9*	14.8*

*Significant difference (t test assuming equal variance, $P < 0.01$)

winter 2015, water temperatures from about 40 m depth down to the lake bed were approximately 12° C, consistent with previous years; however water temperatures in the top 40 m of the water column at station CR346.4 ranged from about 13 – 14.5° C, while in the top 3 m of the water column temperatures ranged from 14.9 to 16.3° C. In all previous winters, surface water temperatures did not go above 13.4° C.

Zooplankton and phytoplankton seasonality

Zooplankton biomass was primarily dominated by *D. pulex* complex in all regions of Lake Mead, secondarily by *D. galeata mendotae*. Generally, peak zooplankton populations were in winter and spring (February, May) when water temperatures were 12 – 14 and 18 – 25° C, respectively (Fig. 5). Throughout the study, zooplankton populations were considerably higher in Las Vegas Bay than Overton Arm and Colorado River. Maximum *D. pulex* complex and *D. galeata mendotae* populations occurred in Las Vegas

Bay; *D. pulex* complex in winter 2011 (mean biomass 537.5μ g d.w./l) and *D. galeata mendotae* in spring 2013 (mean biomass 73.8μ g d.w./l). No *D. pulex* complex was observed in Las Vegas Bay in summer 2010, summer 2013, spring, summer, and fall 2014, or summer 2015. *D. galeata mendotae* was not observed in Las Vegas Bay in winter 2008, winter 2009, winter 2014, or spring 2014.

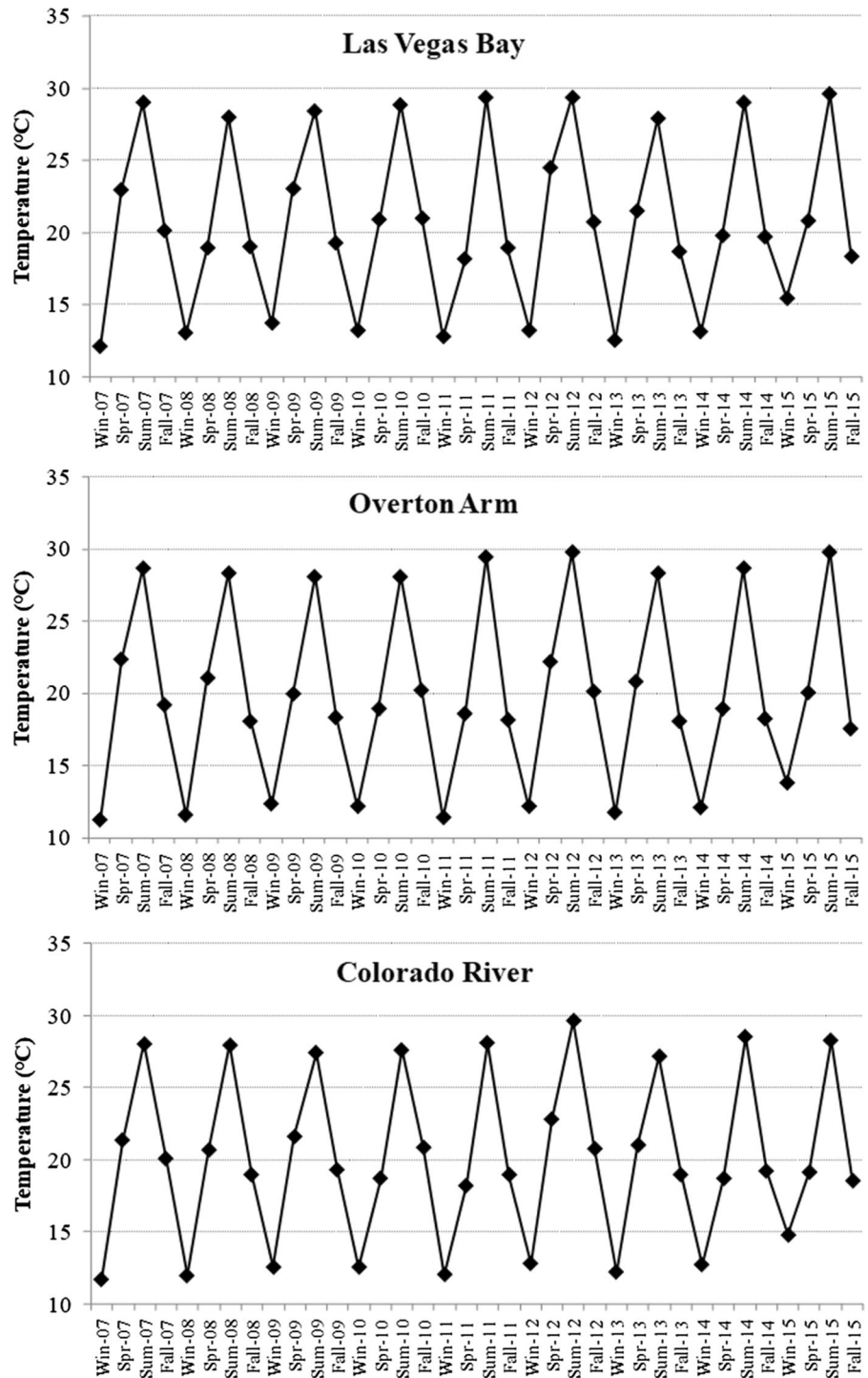
Phytoplankton biovolume in all regions of Lake Mead was divided into four categories: *Cyclotella* spp., *Microcystis* spp., other cyanobacteria, and other phytoplankton. Peak phytoplankton biomass was primarily observed in spring and summer (May, August) when water temperatures were between 18 and 30° C (Fig. 6). Comparable to zooplankton, phytoplankton populations were generally higher in Las Vegas Bay than Overton Arm and Colorado River. Both *Cyclotella* spp. and *Microcystis* spp. mean biovolume peaked within Las Vegas Bay in 2015. *Cyclotella* spp. was highest in summer 2015 ($2.4 \times 10^9 \mu$ m³/l), whereas *Microcystis* spp. was highest in winter 2015 ($6.1 \times 10^8 \mu$ m³/l).

Mean monthly C-zooplankton:C-phytoplankton ratios showed a moderately strong correlation ($R^2 = 0.57$, Fig. 7) to water surface temperatures, which were indicative of lake mixing regimes. Higher Z:P ratios were generally observed at lower surface temperatures, which tend to occur during mixis while lower Z:P ratios were observed at higher temperatures, which typically occur during periods of thermal stratification.

Toxicity

In the spring and summer of 2015, large floating colonies of *Microcystis* spp. were observed in the top 2 m of the water column throughout the reservoir. Microcystin concentrations ranged from non-detect (3 sites in the upper reaches of the Virgin River Arm) to

Fig. 2 Seasonal mean surface water temperatures for Las Vegas Bay stations, Overton Arm stations, and Colorado River stations

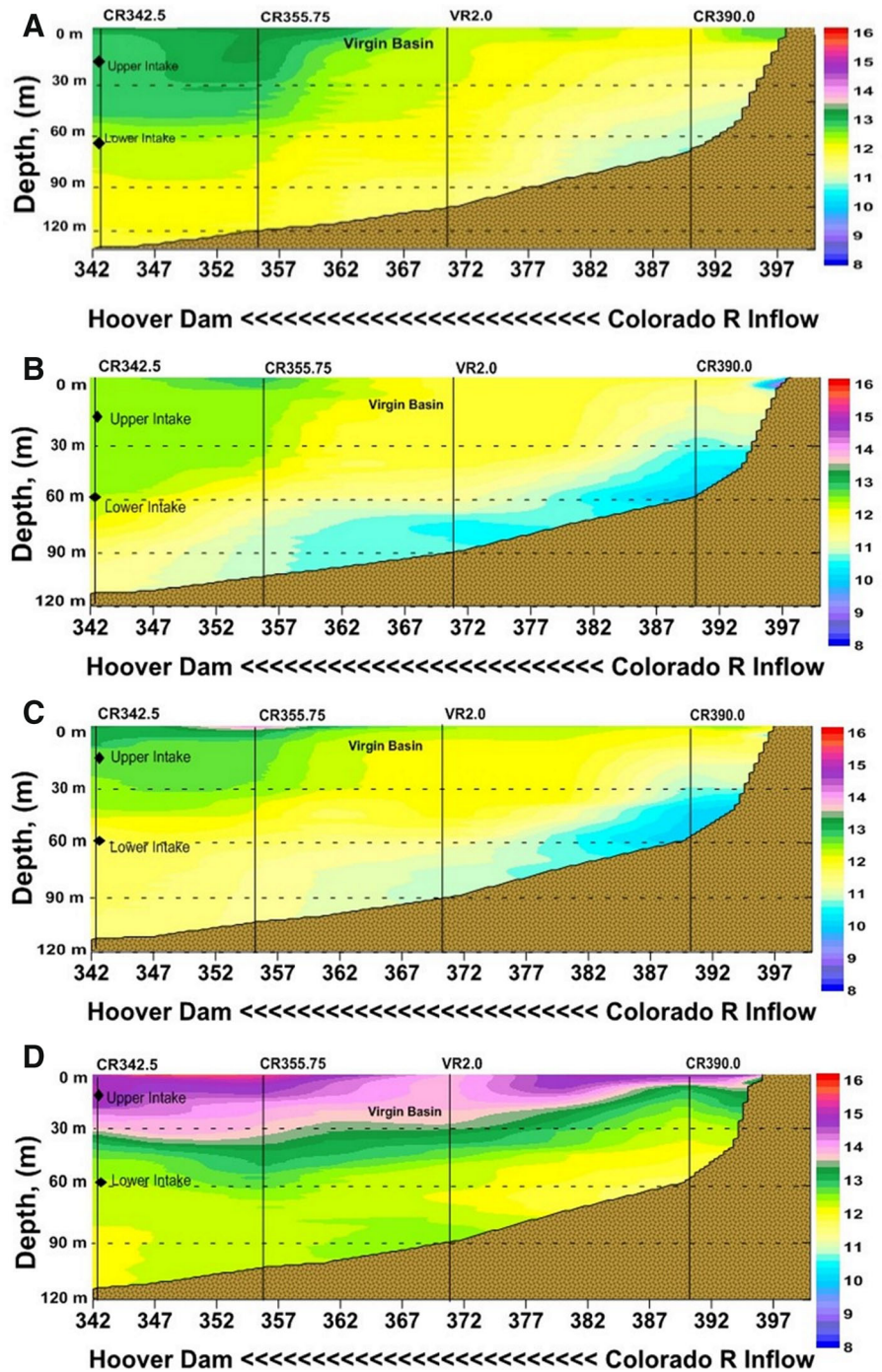


4,696 (Hemenway Harbor) $\mu\text{g/l}$ (Fig. 8). Concentrations were highest in the three near-shore sites. In open water sites, mean microcystin concentration was 1.48 $\mu\text{g/l}$.

Temperature intervals and niche centroids

A relationship between spring/summer daphnid biomass and water temperature was revealed by both the

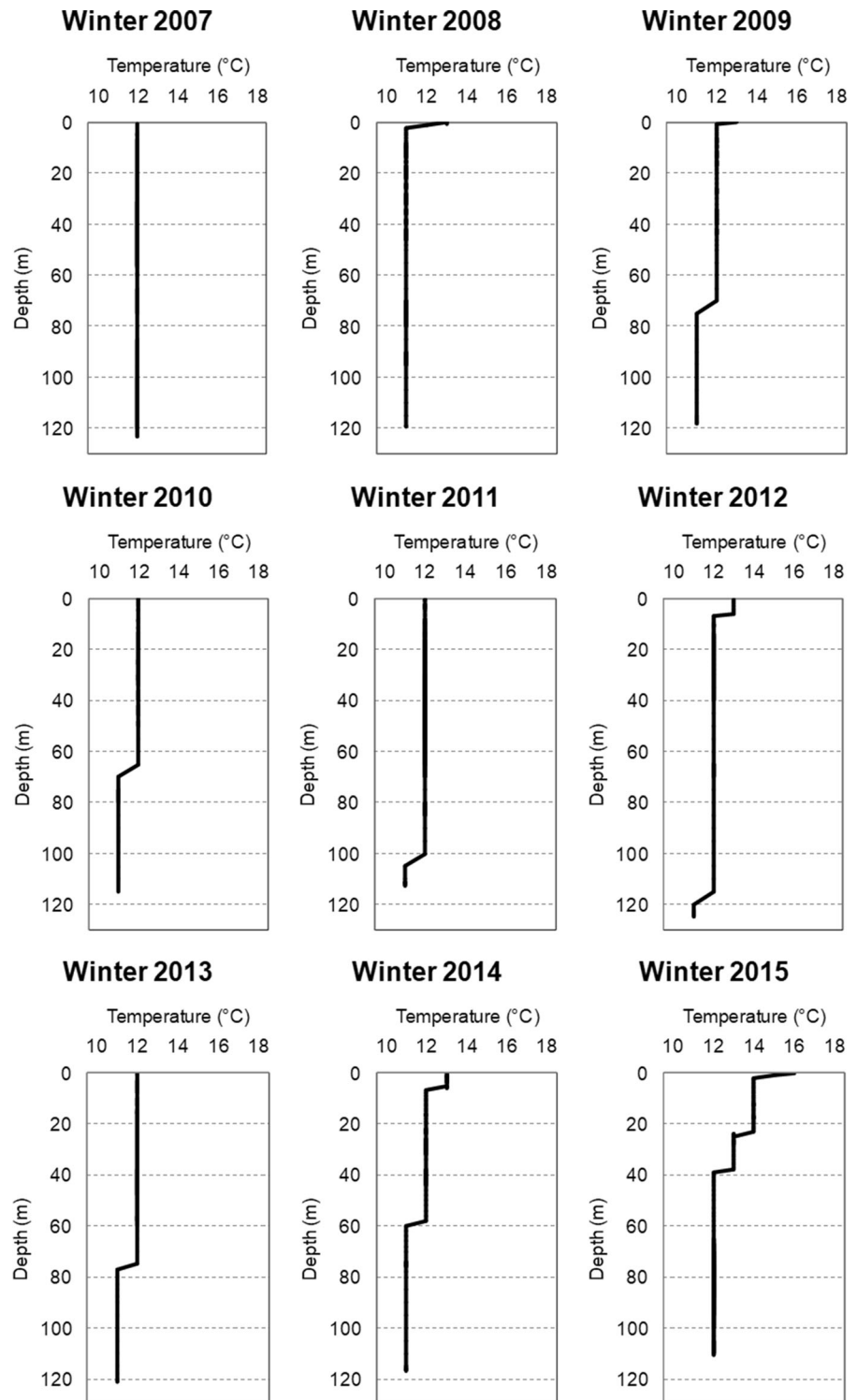
Fig. 3 Thermal profiles of the Colorado River thalweg in Lake Mead in **A** February 2012, **B** February 2013, **C** February 2014, and **D** February 2015



temperature interval regression and the niche centroid analysis (Fig. 9, Table 3). The temperature interval regression models for *D. pulex* complex and *D. galeata mendotae* showed differing temperature preferences for the two species. The regression for *D.*

pulex complex ($R^2 = 0.48$) showed peak biomass between 16 and 18°C, with declines in biomass past about 24°C. The regression for *D. galeata mendotae* ($R^2 = 0.32$) showed a trend of rising biomass until about 24–26°C, and then declines in biomass at higher

Fig. 4 Thermal profiles of station CR364.4 during winter (January/February) sampling events from 2007 to 2015



temperatures. Very high biomass (137.75 $\mu\text{g d.w./l}$) of *D. galeata mendotae* was observed at station LVB3.5

in spring of 2012, producing a very high average biomass for that species at the 24–26°C interval

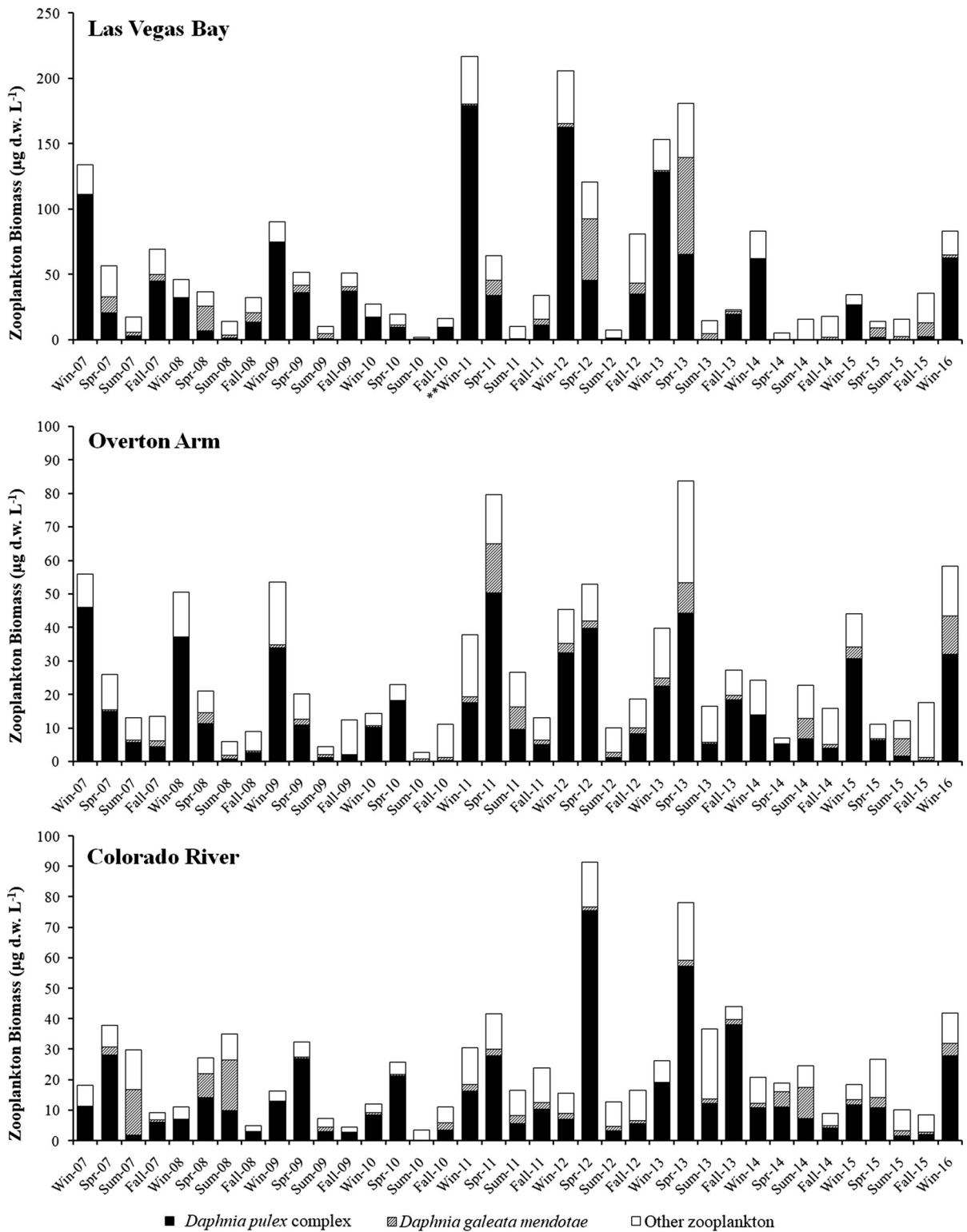
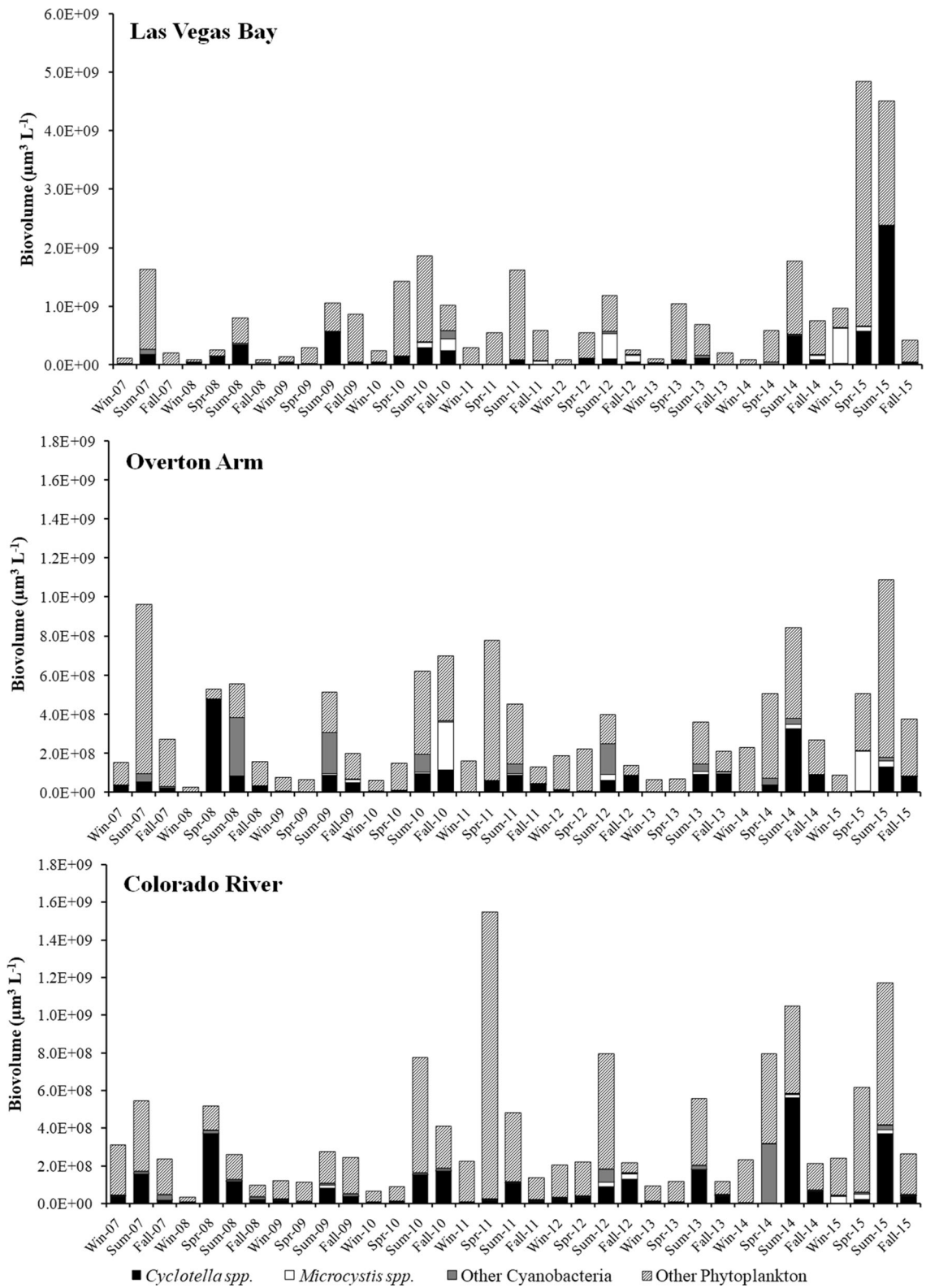


Fig. 5 Quarterly zooplankton biomass means for Las Vegas Bay ($n = 134$), Overton Arm ($n = 208$), and Colorado River ($n = 245$). **Indicates that values are three times greater than shown



◀ **Fig. 6** Quarterly *Cyclotella* spp., *Microcystis* spp., other cyanobacteria, and other phytoplankton biovolume means for Las Vegas Bay ($n = 134$), Overton Arm ($n = 208$), and Colorado River ($n = 245$)

(mean = 10.29 $\mu\text{g d.w./l}$ at 24–26°C, $R^2 = 0.58$ when outlier is excluded). Despite the influence of the outlier value for *D. galeata mendotae*, the results seen from temperature interval regressions imply a seasonal succession pattern in which *D. pulex* complex is dominant early on in the season when water temperatures are cooler, followed by a gradual rise in *D.*

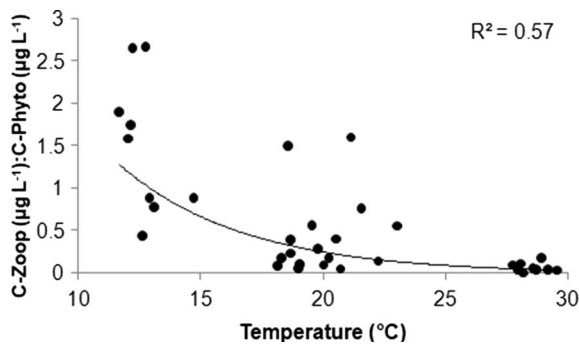
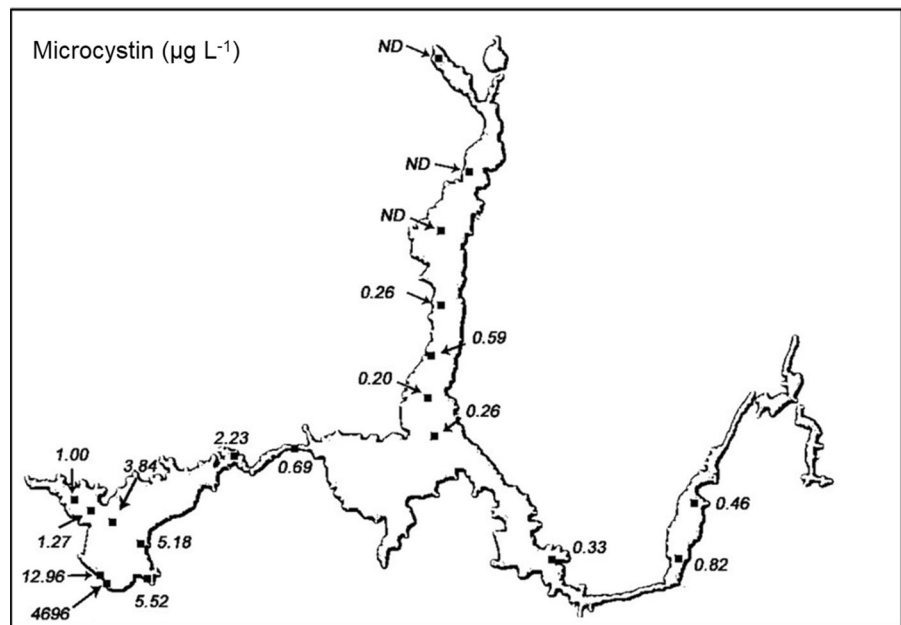


Fig. 7 C-zooplankton:C-phytoplankton ratio (whole-lake means) versus surface (0–5 m) water temperature (whole-lake means) for each sampling event ($n = 36$)

Fig. 8 Map of Lake Mead showing microcystin concentrations at various sampling stations in August, 2015



galeata mendotae biomass as temperatures warm and *D. pulex* complex biomass declines. The niche centroid analysis displayed similar results, suggesting that highest *D. pulex* complex biomass would be found at 16.7°C and highest *D. galeata mendotae* biomass at 22.3°C.

Canonical correlation analysis

Canonical correlation analysis (CCA) compared 34 zooplankton taxa and 4 water quality variables, producing two canonical axes with strong correlations (Table 4). Axes one and two explained 61 and 12% of the total variance, respectively. Water temperature exhibited a strong positive correlation with the first axis (0.600) and a strong negative correlation with the second axis (− 0.640). Secchi depth displayed a strong negative correlation with the first axis (− 0.637), and however was weakly correlated with the second axis. A moderate negative correlation with the first axis and a strong negative correlation with the second axis was observed in regard to station depth (− 0.449 and − 0.547, respectively). Total phosphorus was weakly correlated with the first axis and displayed a strong positive correlation with the second axis (0.535).

For each of the 587 samples analyzed in the CCA, *D. pulex* complex and *D. galeata mendotae* biomass

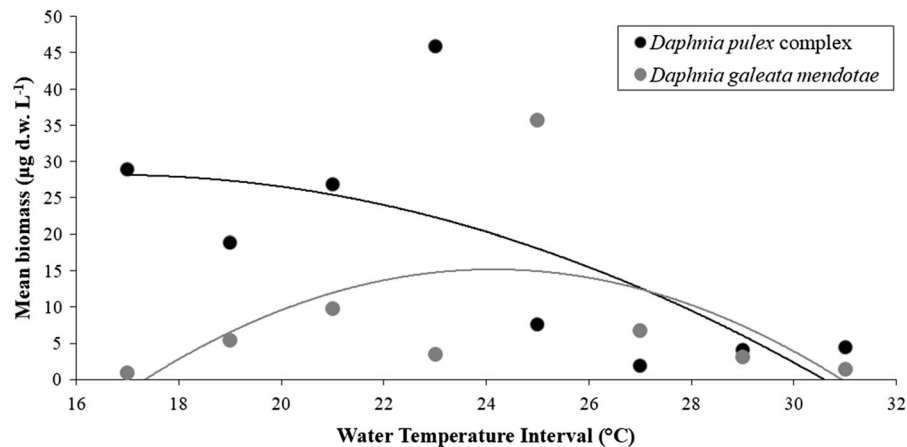


Fig. 9 Mean annual spring and summer (May, Aug) *Daphnia pulex* complex biomass versus surface (0–5 m) water temperature interval (black, $y = -0.145x^2 + 4.836x - 12.04$, $R^2 = 0.48$) and *Daphnia galeata mendotae* biomass versus

surface (0–5 m) water temperature interval (gray, $y = -0.329x^2 + 15.88x - 176.5$, $R^2 = 0.32$) for all three regions of Lake Mead ($n = 276$)

Table 3 Locations of niche centroids (ter Braak & Verdonschot, 1995) showing the relationship between water temperature (°C) and *Daphnia pulex* complex biomass, *Daphnia galeata mendotae* biomass, *Rhodomonas* spp. biovolume, *Cyclotella* spp. biovolume, and *Microcystis* spp. biovolume

	Niche centroid temperature (°C)
<i>Daphnia pulex</i> complex	16.7
<i>Daphnia galeata mendotae</i>	22.3
<i>Rhodomonas</i> spp.	21.5
<i>Cyclotella</i> spp.	25.0
<i>Microcystis</i> spp.	22.4

Table 4 Correlations of water quality variables with their first and second canonical axes in relation to zooplankton biomass ($n = 587$)

	CAP1	CAP2
Temperature (°C)	0.600	– 0.640
Secchi depth (m)	– 0.637	– 0.065
Station depth (m)	– 0.449	– 0.547
Total phosphorus (µg/l)	0.182	0.535

were superimposed onto the ordination diagrams to illustrate correlations with the water quality variables (Fig. 10A, B). *D. pulex* complex was negatively correlated with water temperature indicating a preference for cooler water, and was positively correlated

with water clarity (Secchi depth), which was highest in the Colorado River. *D. galeata mendotae* was negatively correlated with water clarity and positively correlated with temperature, indicating a preference for more turbid, warmer water.

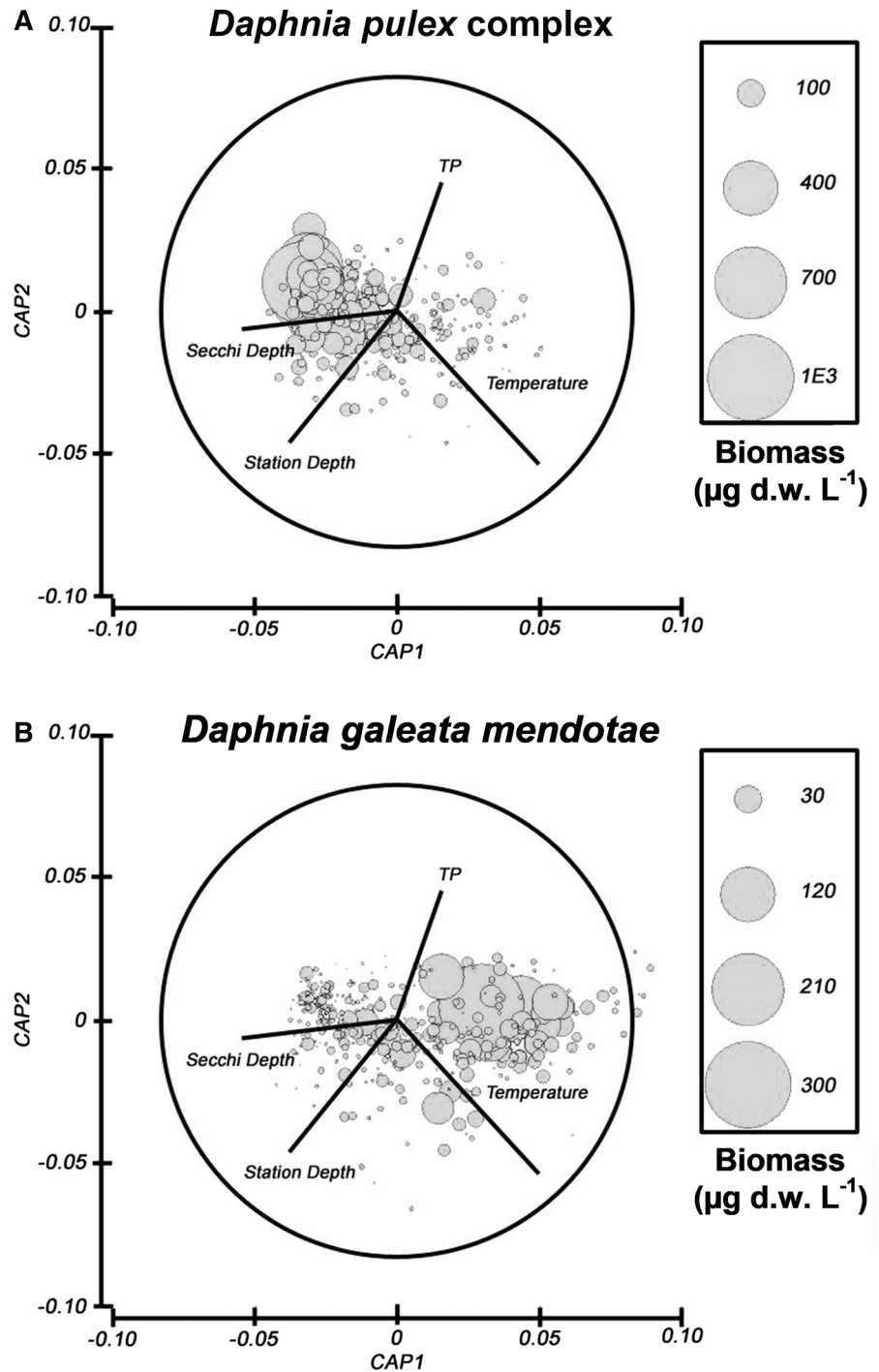
As with the zooplankton biomass, *Cyclotella* spp., *Rhodomonas* spp. and *Microcystis* spp. biovolumes were overlaid onto the zooplankton CCA ordination diagrams to illustrate correlations with the water quality variables (Fig. 11A–C). All three genera displayed similar patterns: *Cyclotella* spp., *Rhodomonas* spp., and *Microcystis* spp. biovolumes were correlated with warm water temperatures and greater water clarity (Secchi depth), which were reflected by the niche centroid temperature optima (25.0, 21.5, and 22.4°C, respectively).

Discussion

Consistency in plankton communities

Previous studies of the zooplankton communities (e.g., Burke 1977; Paulson & Baker 1983; Wilde 1984; Axler et al., 1988; Wilde & Paulson, 1988; Sollberger & Paulson, 1992; Loomis, 2009) and phytoplankton communities (Everett, 1972; Koenig et al., 1972; Staker et al., 1974; Baker & Paulson, 1980; Paulson et al., 1980; Evans & Paulson, 1983; Paulson & Baker, 1983; Prentki & Paulson, 1983;

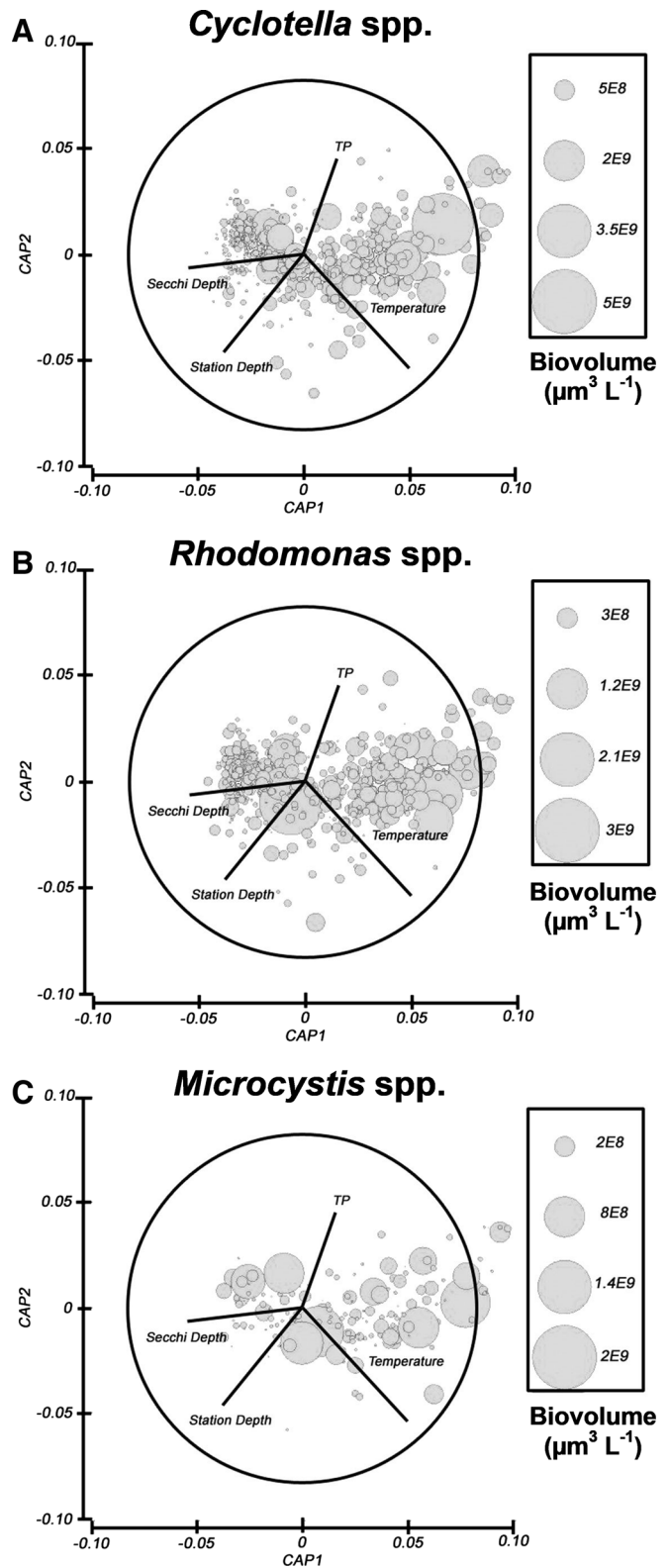
Fig. 10 Zooplankton CCA ordination bubble diagrams for **A** *Daphnia pulex* complex and **B** *Daphnia galeata mendotae* ($n = 587$). Individual bubbles represent the biomass in one sample. Bubble size represents the magnitude of biomass



Janik, 1984) of Lake Mead have been limited in scope and duration and were largely conducted before the current drought conditions, invasion of quagga mussels and establishment of gizzard shad. Consequently, their value for assessing water quality under current

environmental conditions is limited. However, studies examining zooplankton communities prior to current environmental disturbances (drought, invasive species) show that both *D. pulex* complex and *D. galeata mendotae* dominated the crustacean zooplankton

Fig. 11 Phytoplankton CCA ordination bubble diagrams for **A** *Cyclotella* spp., **B** *Rhodomonas* spp., and **C** *Microcystis* spp. ($n = 587$). Individual bubbles represent the biomass in one sample. Bubble size represents the magnitude of biomass



community (Burke, 1977), and exhibited similar seasonal succession patterns to what was observed in this study (dominance of *D. pulex* complex in winter followed by *D. galeata mendotae* in spring, Wilde, 1984). The most recent long-term analysis of zooplankton community dynamics in Lake Mead (Beaver et al., 2010) found that biomass of the dominant zooplankton taxa (*Daphnia* spp.) generally peaked in the winter/early-spring and was lowest during summer. Results from this study show that seasonal patterns in *Daphnia* dominance have continued through the current study period, as similar patterns were observed across all three basins. The results of these two studies cover, together, a 15-year period (2000–2015) showing little change in *Daphnia* dynamics and minimal shifts in community composition.

Phytoplankton composition and seasonality in this study also showed similar trends to what has been observed in the past (Staker et al., 1974; LaBounty & Burns, 2005; Turkett, 2016), with highest biovolume occurring in Boulder Basin (Las Vegas Bay sites and downstream CR sites) and annual biovolume peaks during the summer. The seasonal pattern observed for phytoplankton, as well as niche centroid analysis performed for three major taxa, indicate that phytoplankton biovolume is strongly tied to temperature cues and water clarity. Although not explored in this report, previous studies have determined that Lake Mead is severely phosphorus-limited (Paulson & Baker, 1983; LaBounty & Burns, 2005) and that past phytoplankton blooms have occurred in response to sudden, increased nutrient inputs (LaBounty & Burns, 2005). Alignment of zooplankton and phytoplankton data display an inverse correlation in terms of seasonal shifts in total biomass/biovolume, suggesting that grazing may help limit phytoplankton growth in the winter, when *Daphnia* are at their most abundant. Alternatively, zooplankton growth might be limited by phytoplankton, as phytoplankton biovolume in Lake Mead is generally much lower than in other U.S. reservoirs (Turkett, 2016).

A recent study of seasonal food web dynamics in Lake Taupō, New Zealand (Stewart et al., 2017) indicated that in large, deep oligotrophic monomictic lakes, the control of the food web may alternate between two states; during the stratified period, primary production may be limited by nutrient availability (bottom-up control), while during mixing,

primary production may be more strongly controlled by grazing (top-down control). In Lake Mead, zooplankton:phytoplankton carbon ratios also vary seasonally, with highest ratios observed during mixing periods (winter) most years. This suggests that mixis is critical in terms of structuring plankton communities and setting boundaries on primary production. Long-term trends in chlorophyll from Beaver et al. (2010) showed a decline in Las Vegas Bay, but no change between 2000 and 2007 in other reservoir basins. Over the course of this study, no significant increases or decreases in total phytoplankton biovolume were observed. Species composition also remained relatively consistent on a seasonal and inter-annual basis, save for a few isolated instances of high *Microcystis* spp. biovolume, the highest of which occurred in Las Vegas Bay in 2015 following abnormally strong thermal stratification in the winter of that year.

Detailed taxonomic analysis revealed a seasonal succession within the daphnid community, showing that the larger *D. pulex* complex is dominant during winter, while the smaller *D. galeata mendotae* makes more significant contributions to total biomass during spring and summer. Co-occurrence of *D. pulex* complex and *D. galeata mendotae* was common in a recent survey of reservoir systems in the western U.S. (Beaver et al., 2014), implying that zooplankton community dynamics observed in Lake Mead are likely to be comparable elsewhere in the region. CCA performed in this study revealed that surface temperature was a strong driver of species variation within the zooplankton community. Both niche centroid analysis and temperature interval regressions showed a lower temperature optimum for *D. pulex* complex than for *D. galeata mendotae*. These results indicate that seasonal shifts between the two daphnid species are closely tied to changes in surface water temperatures. In Lake Mead surface temperatures range from about 11 to 31°C over the course of the year, with annual thermal stratification during summer months (Holdren & Turner, 2010). Habitat partitioning via diel vertical migration in deep lakes with hypolimnetic refugia is thought to be a response to increased zooplanktivory (Jeppesen et al., 2003), which generally is higher during the summer. In both Lake Mead and Lake Powell (the second largest U.S. reservoir, located upstream from Lake Mead on the Colorado River), large-bodied *D. pulex* complex are nearly absent from the epilimnion during the summer, with

largest populations located in cooler strata below 10 m at open water sites, while small-bodied *D. galeata mendotae* remain in the upper 10 m and are usually found in warm, productive bays influenced by nutrient inflows (Beaver, unpublished data). Similar habitat partitioning between daphnid species with differing thermal tolerances and vertical migration tendencies has been observed in other studies (Stich & Lampert, 1981; Leibold, 1991; Seda et al., 2007).

Zooplankton form a critical link in the Lake Mead food web for both forage fish (threadfin shad, Loomis et al., 2011) and sportfish such as largemouth bass (Wilde & Paulson, 1988) and striped bass (Wilde & Paulson, 1989a), as well as the native, endangered razorback sucker (Marsh & Langhorst, 1988). Because of their contribution to total zooplankton biomass, *Daphnia* are the pivotal intermediaries in energy transfer to higher trophic levels in Lake Mead and other southwestern U.S. reservoirs. Threadfin shad spawning coincides with daphnid peaks and continues through June in lower Colorado River reservoirs (Wilde & Paulson, 1988, 1989b; Mueller & Horn, 2004; Beaver et al., 2010; Loomis et al., 2011). Annual productivity of sportfish is strongly related to interactions among nutrients, zooplankton and phytoplankton in reservoirs throughout North America (Stockner et al., 2000). The lack of significant changes in both the zooplankton and phytoplankton communities in Lake Mead implies that, despite climatic and ecological disturbances, major forage and sportfish communities (and thus recreational fishing activities) are not currently threatened by disruptions in the food web at lower trophic levels.

Effects of invasive species and drought

Empirical data from the Great Lakes region indicate that dreissenid mussels have both direct and indirect effects on zooplankton populations via consumption of small zooplankters (rotifers and small cladocerans) and competition for food (phytoplankton) (MacIsaac, 1996). Competition between the two groups has the potential to decouple established trophic linkages among nutrients, phytoplankton, zooplankton, and planktivorous fish leading to increased water clarity, reduced turbidity, and altered phytoplankton and zooplankton composition (e.g., Fahnenstiel et al., 2010; Kerfoot et al., 2010; Kissman et al., 2010). Results from the CCA show that *D. pulex* complex

exhibits higher biomass under increased water clarity in Lake Mead. As such, it is possible that filtration by quagga mussels might benefit *D. pulex* complex by improving water clarity and expanding the extent of preferable habitat for the large cladoceran. On the other hand, increased water clarity has improved the reaction distance for size-selective visual predators which most negatively impacts large-bodied daphnids, such as *D. pulex* complex (Vinyard & O'Brien, 1976; Miner & Stein, 1993).

A minor decrease in chlorophyll and an increase in water clarity in Boulder Basin (including Las Vegas Bay sites and station CR346.4) post-quagga invasion compared to pre-quagga invasion has been documented (Wong et al., 2010, 2014); however, changes in these parameters were found to be closely associated with declines in phosphorus (human-mediated efforts) and cannot be directly attributed to quagga filtering. A similar decrease in chlorophyll was observed by Beaver et al. (2010); however, this decline was also concurrent with large reductions in phosphorus inputs. Thus, it is difficult to say whether quagga mussels have contributed to declines in phytoplankton, as would be expected following invasion in smaller lakes or reservoirs. The apparently low impact that dreissenid mussels have had on plankton communities in Lake Mead is likely due to specificities in basin morphology, as Lake Mead is much deeper and has a longer water residence time (1–3 years, LaBounty & Burns, 2005) than most eastern U.S. systems that have been impacted by quaggas (Bridgeman et al., 1995; Fanslow et al., 1995; Caraco et al., 1997; Turkett, 2016). Dreissenid filtration rate is dependent on particle concentration, and is accelerated when particle concentration is low (Sprung & Rose, 1988). Filtration speed is also impacted by particle size, temperature, population size, and quality of food available (i.e., ingestion vs. production of pseudofeces) (Turkett, 2016 and references therein). Because of the comparatively large volume of water in Lake Mead, expected ecosystem effects of quagga filtration (i.e., reduction in zooplankton and phytoplankton) may be delayed.

Juvenile (0-year) gizzard shad feed primarily on zooplankton, but may feed on detritus or phytoplankton as they grow (Dettmers & Stein, 1992). Previous studies have shown that gizzard shad have the capacity to nearly eliminate crustacean zooplankton from lakes, if densities are high enough (Dettmers & Stein,

1992, 1996). The expansion and proliferation of gizzard shad throughout the lower Colorado River has been well documented; however, few studies have specifically examined the impacts of invasion on Colorado River reservoir ecosystems (Webber & Jones, 2013). Introductions of gizzard shad and quagga mussels into Lake Mead likely occurred within 4–5 years of each other. Both species are known to consume phytoplankton as a food source; however, the presence of quagga mussels appears to have had minimal impact on slowing the increase in gizzard shad populations in Lake Mead (Wong & Gerstenberger, 2011). While quagga mussels are known for improvement of water clarity in lakes after becoming established (Limburg et al., 2010), gizzard shad often have the opposite effect. Benthic foraging by gizzard shad can disturb sediment in lakes, which may cause increases in total nutrients, reduced water clarity and increases in phytoplankton abundance (Schaus & Vanni, 2000). Thus far, however, neither species has had significant impacts on water clarity or overall phytoplankton abundance in Lake Mead. Both quagga mussels and gizzard shad have expanded in Lake Mead since the time of their introduction; however, declines in zooplankton have not been observed, despite the potential for both species to decimate populations. As with quagga mussels, most previous studies of the impacts of gizzard shad on plankton communities were conducted in smaller, shallower lakes (e.g., Pierce et al., 1981; Dettmers & Stein, 1996; Schaus et al., 2010); it is likely that the depth of Lake Mead restricts interactions between the benthos and the epilimnion to the shallow inflow areas.

Limited studies exploring the effects of drought on plankton communities have shown either an increase in total zooplankton abundance (Olds et al., 2014), a shift from dominance of cladocerans to copepods (Havens et al., 2007), a greater contribution from *Daphnia* spp. (Beaver et al., 2013) or no significant shift in phytoplankton or zooplankton populations (McGowan et al., 2005). Long-term trends in lake water surface temperature worldwide show that the majority of lakes, regardless of geographical location, have warmed over the last few decades and that surface water warming is closely tied to rising air temperatures; however, the degree and character of surface water warming are highly variable and dependent on specific lake morphology (Butcher et al., 2015; Kraemer et al., 2015; O'Reilly et al.,

2015). Accordingly, deeper lakes and reservoirs such as Lake Mead may exhibit slower ecosystem response to the effects of drought and increasing air temperatures.

Occurrence of toxic cyanobacteria

In the past, Lake Mead has experienced periods of high productivity (due to elevated phosphorus before 2002) and blooms of non-toxic species (LaBounty & Burns, 2005). Since about 2000, *Microcystis* spp. has been observed in Lake Mead, often appearing in high abundances in net tow samples (Beaver, unpublished data). However, the 2015 bloom of *M. aeruginosa* was the first event, to our knowledge, in which harmful toxins were detected in Lake Mead. Although toxin levels were lower in pelagic zones of the reservoir, all basins experienced some degree of toxicity. Traditionally, cyanobacterial blooms have been associated with high phosphorus inputs, contradicting what was observed in Lake Mead; however, there is growing evidence that non-N₂-fixing cyanobacteria (such as *Microcystis* spp.) can outcompete other cyanobacterial taxa under low-P conditions, as long as nitrogen concentrations are sufficiently high (Paerl & Otten, 2013). Since total nitrogen was not analyzed in this study, it is difficult to say if this was the case. Our analyses show that *Microcystis* spp. had a thermal optimum during the study period of ~ 22°C. Previous studies have found that *Microcystis* spp. can thrive at temperatures between 15 and 30°C (Chorus & Bartram, 1999) and does best under highly stratified conditions in which it can regulate buoyancy to occupy the surface layer of the water column. During the 2015 bloom event, large colonies of *M. aeruginosa* were observed several meters below the water's surface. Previous long-term studies on the German lake Müggelsee show that toxic cyanobacteria can dominate phytoplankton communities once thermal stratification thresholds are reached, regardless of ambient air temperatures (Wagner & Adrian, 2009; Huber et al., 2012), and enclosure experiments show that cyanobacterial blooms can actually help to prolong stratified conditions (Kumagai et al., 2000). Overall toxicity of *Microcystis* spp. has also been linked to increased temperatures (Davis et al., 2009).

In Lake Mead, the likelihood of toxic bloom reoccurrence may be exacerbated by populations of quagga mussels. In dreissenid-invaded systems in the

Great Lakes region of the United States, invasion has often been followed by establishment of annual *Microcystis* spp. blooms (Lavrentyev et al., 1995; Vanderploeg et al., 2002). Selective filtration by quagga favors *Microcystis* spp., as mussels preferentially discard that species in favor of diatoms and other phytoplankton. As a result, microcystin concentrations have been shown to be higher in lakes with established populations of dreissenid mussels (Knoll et al., 2008). Likewise, invasion of gizzard shad may increase likelihood of phytoplankton blooms if zooplankton predators are significantly decreased. The potential thermal overlap between dominant *Daphnia* spp. and *Microcystis* spp. indicates that these taxa could co-occur. *Daphnia* are susceptible to inhibitory impacts to food ingestion (Rohrback et al., 1999) and death from microcystin poisoning (Rohrback et al., 2005). Recurrences of toxic blooms could be detrimental to the dominant zooplankton of Lake Mead and could cause cascading reductions in higher trophic levels.

The 2015 *M. aeruginosa* bloom followed high water temperatures and earlier than normal stratification in Lake Mead resulting from a combination of low winter precipitation and warm spring and summer air temperatures. Similar conditions were observed during that time period across most western states in the U.S. (Mote et al., 2016) and were likely related to higher than average off-shore sea surface temperatures that persisted from the winter of 2013/2014 through the spring/summer of 2015 (Bond et al., 2015). Varying ecological consequences from across the western U.S. and eastern Pacific coastline were reported following the 2014/2015 warm weather anomaly (Cavole et al., 2016), including multiple instances of organisms appearing outside of their expected range. There are many examples of sea surface temperature anomalies causing droughts and disrupting plankton composition in lakes and reservoirs particularly in shallow productive systems. Prolonged droughts create favorable conditions for cyanobacteria by increasing water residence time, thermal stability, and surface temperature, (e.g., Bouvy et al., 2000; Reichwaldt & Ghadouani, 2012; Brasil et al., 2015) while creating conditions less favorable for cladocerans (Havens et al., 2017; Ji et al., 2017). The fact that the summer of 2015 was ubiquitously warm in western U.S. states leads to the conclusion that biological consequences during that time period (including the 2015 *M. aeruginosa* bloom

in Lake Mead) stem primarily from anomalous ocean conditions that have since dissipated. However, it remains unclear whether this type of scenario will be more common in the future as drought in the southwestern U.S. persists.

Non-linear responses to change

Ecological theory suggests that biological responses to environmental stress may not exhibit linear reactions to gradual change; instead, gradual degradation in environmental quality may cause a system to slowly lose resiliency in non-quantifiable ways until a threshold is reached, inducing drastic changes in population dynamics (Scheffer et al., 2001). Lake Mead has certainly experienced gradual environmental change over the last 15 years, as drought has steadily decreased reservoir water levels and invasive species have spread and multiplied (Fig. 12). The 2015 *M. aeruginosa* bloom in Lake Mead may be an example of a non-linear response to environmental change. Up until 2015, surface water temperatures largely remained consistent on an inter-annual basis. Likewise, phytoplankton seasonal dynamics and community composition have been relatively stable, indicating that the effects of drought on zooplankton and phytoplankton have been minimal thus far. The unusually warm weather in 2015 (and associated effects on snowpack and thermal stratification) demonstrated that phytoplankton composition can change dramatically when water temperatures rise a few degrees.

The morphology of Lake Mead (specifically in terms of size and depth relative to many North American reservoirs) has likely been a factor contributing to the resilience of this ecosystem; however, given predictions about climate warming and continued proliferation of quagga mussels and gizzard shad, it is probable that environmental stress will continue to grow in scope and that impacts to the food web will become more severe. Lake Havasu, located approximately 250 km downstream from Lake Mead, also experienced a large bloom of *M. aeruginosa* in the summer of 2015. Although the 2015 bloom appears to be an isolated incident thus far in Lake Mead, Lake Havasu has been experiencing annual recurrences of high *M. aeruginosa* densities in the past several years (H. McMaster, unpublished data). This may be due in part to the differences in morphology and hydrology

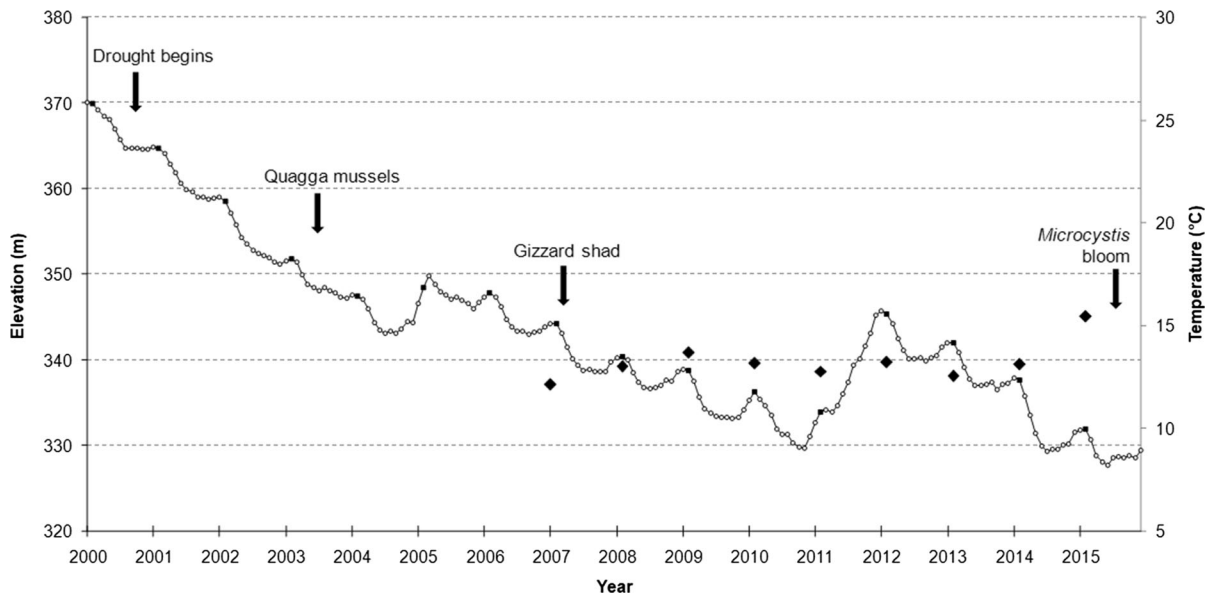


Fig. 12 Monthly mean surface water elevation for Lake Mead (open circles) from 2000 to 2015. Black squares represent mean surface water elevation in February of each year. Black

diamonds represent mean winter surface water temperature (0–5 m) in Las Vegas Bay (2007–2015). Major disturbances affecting Lake Mead are denoted with black arrows

between the two reservoirs, as Lake Havasu is shallower and warmer on average than Lake Mead. Both Lake Mead and Lake Havasu contain a reservoir of *M. aeruginosa* cells that are present in the phytoplankton community nearly every year. The fact that Lake Havasu experiences higher densities of *M. aeruginosa* annually than Lake Mead suggests that the smaller, shallower reservoir may have already crossed an environmental threshold to where high abundances of cyanobacteria are seen during the stratified period every summer. This indicates that Lake Mead may be closer to an irreversible ecological tipping point, where similar toxic bloom events could occur more frequently in a climatically warmer future.

Conclusions

Lake Mead has not demonstrated significant impacts to established trophic level relationships since the beginning of the drought and subsequent invasion by invasive species in the southwestern U.S. Despite multiple ongoing environmental stressors such as drought, invasion of dreissenid mussels, and the establishment of gizzard shad, temporal and spatial patterns of zooplankton and phytoplankton in Lake

Mead remained comparable to historic trends during the study period.

The 2015 bloom of *M. aeruginosa* in Lake Mead was likely a result of anomalous weather conditions (i.e., decreased snowpack and above average air temperatures); however, these conditions may become more commonplace as the climate warms. The compounding effects of climate warming, persistent drought, increasing water demand, and expansion of invasive species have unknown consequences for the future health of the Lake Mead ecosystem. Because seasonal patterns have been relatively consistent on an inter-annual basis, shifts in zooplankton and phytoplankton species composition and phenology could be early signs of changing environmental conditions, indicative that Lake Mead might be experiencing a threshold response to environmental disturbances. Continuation of current monitoring efforts will shed light on the 2015 *M. aeruginosa* bloom in Lake Mead and will reveal whether the combination of reduced flow and warmer surface temperatures are indeed predictable catalysts for the occurrence of toxic blooms in southwestern U.S. reservoirs.

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