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Aquatic Sciences and Engineering aims to contribute to the literature by publishing manuscripts at the highest scientific level on all fields of aquatic sciences. The journal publishes original research and review articles that are prepared in accordance with the ethical guidelines.

The scope of the journal includes but not limited to; aquaculture science, aquaculture diseases, feeds, and genetics, ecological interactions, sustainable systems, fisheries development, fisheries science, fishery hydrography, aquatic ecosystem, fisheries management, fishery biology, wild fisheries, ocean fisheries, biology, taxonomy, stock identification, functional morphology freshwater, brackish and marine environment, marine biology, water conservation and sustainability, inland waters protection and management, seafood technology and safety.

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The Establishment of the Non-indigenous Cyclopoid Copepod *Oithona davisae* in Hamsilos Bay-Sinop, Southern Black Sea, Turkey

Funda Üstün¹ , Tuba Terbiyik Kurt² , Zeynep Hasançavuşoğlu¹ 

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ABSTRACT

Hamsilos Bay, which is located within the first-degree natural protection area, is one of the most important regions on the southern Black Sea coast. Moreover, many fish species utilize this area for breeding and rearing activities. Small copepod *Oithona davisae* has been encountered in the Black Sea for two decades and spread rapidly the overall Black Sea, and become an important component in the pelagic ecosystem of the Black Sea. In this respect, the present study aims to characterize the distribution and abundance of *Oithona davisae* in Hamsilos Bay about environmental variables. Zooplankton samples were collected monthly from seven stations from July 2015 to June 2016 by using a plankton net with a mesh size of 112 µm and a diameter mouth opening of 50 cm. *O. davisae* was observed during the whole sampling period and was the dominant species from July 2015 to December 2015. The highest mean abundance of this species was recorded in September 2015 (3533.33 ind. m⁻³). However, the lowest mean abundance was observed between January 2016 (170.68 ind. m⁻³) and May 2016 (0.24 ind. m⁻³). Copepodites 5-4 were predominant in the *O. davisae* population in all months and all stations. Adult copepods and their copepodites 5-4 were more abundant between July 2015 and December 2015, whereas the abundance of copepodites 3-1 was maximum between September 2015 and December 2015. The abundance of *O. davisae* was highly correlated with chlorophyll- α and physiochemical variables (temperature, dissolved oxygen, and salinity). In conclusion, this species, which is known to be distributed only in the autumn conditions in close areas, extended its temporal distribution and prevailed in the copepod communities during warmer periods. This could be attributed to the acclimatization success of this species in Hamsilos Bay ecosystems.

Keywords: *Oithona davisae*, non-native, invasive species, abundance, developmental stages, physiochemical variables

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INTRODUCTION

Marine ecosystems are influenced by human activities such as implantation, aquacultural facilities, transoceanic shipping, and fisheries activities. These activities lead to the entrance of non-indigenous species and the change in native communities throughout the world (Leidenberger et al., 2015). Non-indigenous species have an impact on native communities, habitats, and ecosystem processes. For this reason, regular monitoring of the introduction of non-native species into the ecosystem and its

ecological effects is important to follow up on the changes that may occur in the environment and respond appropriately in time to the potential effects they can create (Brog, 2009).

The Black Sea, a semi-enclosed inland sea, is connected to the Aegean Sea and the Mediterranean Sea in the southwest through the Turkish Straits System and to the Sea of Azov in the north through the Kerch Strait (Ünlüata et al., 1990). The Black Sea is a very sensitive ecosystem that can be easily affected by alien (non-native) species due to its geographic location (Se-



lifonova, 2011a). It is known that the introduction of non-indigenous species into the Black Sea started in the 19th century and led to great problems in the following decades (Shiganova, 1998). One of the alien species that entered the Black Sea probably with ballast waters (Gubanova et al., 2014) is the cyclopoid copepod *Oithona davisae*. Known to be native to the Seas of Japan and China (Razouls et al., 2023), *Oithona davisae* has successfully settled in many different marine and estuarine ecosystems such as the Barcelona Harbor in Spain (Saiz et al., 2003), Marmara Sea (Doğan & İşinibilir Okyar, 2016), Aegean Sea in Turkey (Terbiyik Kurt & Beşiktepe, 2019), İskenderun Bay in Levantine Sea in Turkey (Terbiyik-Kurt et al., 2022), Sacramento-San Joaquin estuary in California (Ferrari & Orsi, 1984), San Francisco Bay estuary in California (Lougee et al., 2002), Bilbao estuary in Spain (Uriarte et al., 2016). *O. davisae* was first found in 2001 in Sevastopol Bay in the Black Sea. Firstly, it was misidentified as *Oithona brevicornis* (Zagorodnyaya, 2002), but later it was identified as *O. davisae* upon re-examination of zooplankton specimens (Temnykh & Nishida, 2012). The first record of *O. davisae* was in Sinop in 2009 along the Black Sea coast of Turkey (Üstün & Terbiyik Kurt, 2016). Later, *O. davisae* expanded its distribution towards the eastern Black Sea (Trabzon coast) (Yıldız et al., 2017) in 2010 and the western Black Sea (İğneada coast) in 2012-2013 (Üstün et al., 2019). It is seen that *O. davisae* become to have a higher abundance and frequency, thus expanding its range on the Turkish coasts of the Black Sea after reporting by Üstün & Terbiyik Kurt, 2016 on the Sinop coast.

Hamsilos Bay had great significance on the southern Black Sea coast due to its location within the first-degree natural protection area (Anonymous, 2018). Furthermore, Hamsilos Bay is a breeding and rearing area for many fish species. In the present study, the temporal and spatial distribution of the abundance values of *O. davisae* in Hamsilos Bay, a protected area, and its relationship with environmental parameters have been first examined.

MATERIALS AND METHODS

Sampling area

Our study area, Hamsilos Bay, is 11 km from Sinop, a natural harbor located at the northernmost point of Turkey. While it looks like a fjord, it is a small ria formed by the submersion of a valley underwater (Akkan, 1975). The importance of Hamsilos Bay is demonstrated by its status as a natural area of protection, the absence of residential areas, and no dumping of any pollutants in this area, therefore is no stress factor harming the flora and fauna living here. Additionally, as it is an enclosed and protected area in terms of winds (mistral–northbound) and covered with *Zostera* facies in its deep structure (Ersoy Karaçuha, 2006), it provides shelter for various benthic and demersal living beings and makes the Bay a significant reproduction and shelter area for fish (Uygun, 2015).

Sampling and laboratory studies

The study was conducted monthly at seven stations in Hamsilos Bay from July 2015 to June 2016 (Table 1 and Figure 1). Environmental variables were simultaneously obtained after zooplankton samplings. The physical parameter (temperature, dissolved oxygen, and salinity) values of the surface seawater were recorded with the YSI 6600 MDS model multiparameter (YSI Incorporat-

Table 1. Information about sampling stations.

Station name	Information of Coordinate	Sampling Depth (m)
A	42°03'38"N – 35°02'31"E	2
B	42°03'36"N – 35°02'36"E	4
C	42°03'40"N – 35°02'36"E	9
D	42°03'45"N – 35°02'40"E	13
E	42°03'52"N – 35°03'14"E	30
F	42°04'5"N – 35°02'59"E	30
G	42°04'12"N – 35°02'45"E	30

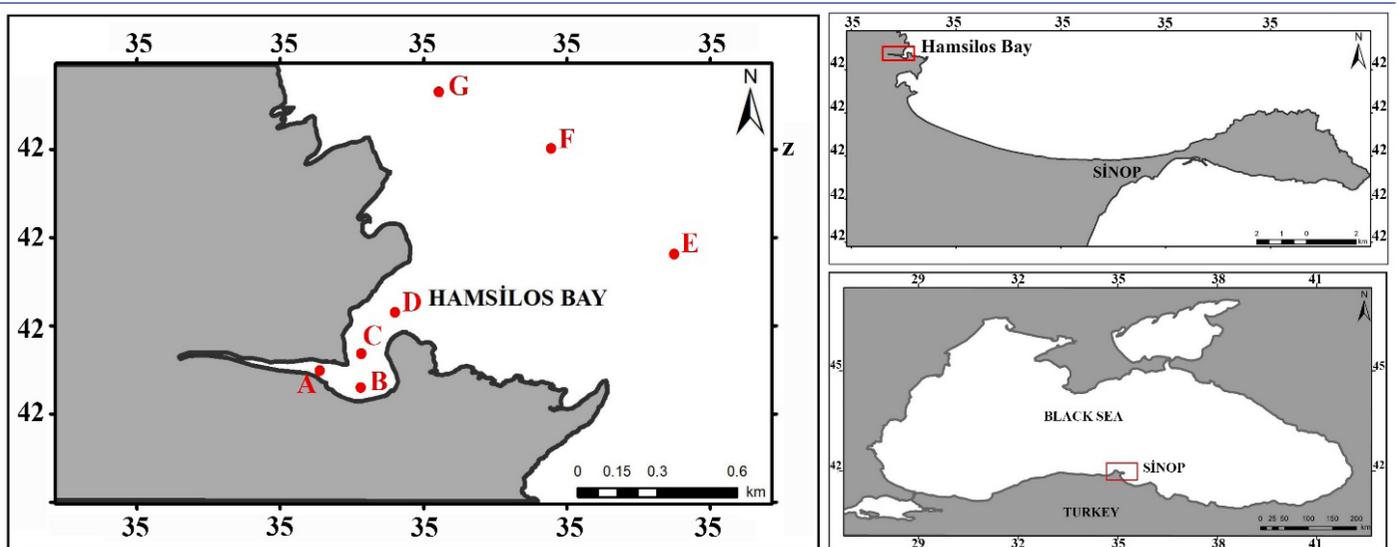


Figure 1. Location of sampling stations.

ed, Ohio, USA). The concentration of chlorophyll-*a* ($\mu\text{g. l}^{-1}$) was estimated according to Parsons et al. (1984).

A plankton net with a mouth opening of a diameter of 50 cm and a mesh size of 112 μm was vertically towed from the bottom to the surface of the water during the daytime. After sampling, the collected material was transferred into a bottle and preserved in a solution of borax-buffered 4% formaldehyde in seawater. *O. davisae* individuals were quantitatively examined using subsamples (two replicates) with a Stempel pipette (1 ml). Subsamples varied from 1/1 to 1/100 (usually 1/25 and 1/50) depending on the abundance of individuals in the samples. The abundance results of *O. davisae* were given in individual (ind. m^{-3}). The consequences were presented as mean \pm standard deviation.

In a zooplankton counting chamber, the development stages of *Oithona davisae* were identified and counted under a stereomicroscope. The present study did not include naupliar stages of this species because the mesh size of our zooplankton was inappropriate (Uye & Sano, 1995). According to Ferrari & Orsi (1984) and Temnykh & Nishida (2012), *O. davisae* was identified.

Analyses of Spearman correlation (SPSS 21, IBM Corp., Armonk, NY, USA) were applied to determine the correlation between the abundance of *O. davisae* and abiotic and biotic variables. The difference in abundance values of developmental stages of *O. davisae* between stations and months was tested by One-way ANOVAs with post hoc Tukey tests (SPSS 21, IBM Corp., Armonk, NY, USA).

RESULTS

Environmental parameters

A detailed description of the hydrological status of the sampling site was given by Üstün (2019) and Üstün & Birinci Özdemir (2019). During the sampling period, a clear seasonal cycle was observed in the mean temperature of surface seawater and fluctuated

from 8.44 ± 0.04 °C (February 2016) to 25.71 ± 0.16 °C (August 2015) in Hamsilos Bay. The surface seawater means salinity ranged between 17.74 ± 0.07 ‰ (September 2015) and 18.96 ± 0.03 ‰ (May 2016). The dissolved oxygen values in surface seawater varied from 7.56 ± 0.36 mg. l^{-1} (July 2015) to 9.88 ± 0.13 mg. l^{-1} (February 2016). The mean concentration of chlorophyll-*a* fluctuated between 0.15 ± 0.10 $\mu\text{g. l}^{-1}$ (March 2016) and 0.71 ± 0.16 $\mu\text{g. l}^{-1}$ (October 2015) in the surface seawater (Figure 2).

Seasonal dynamics of population abundance of *Oithona davisae*

The clear seasonal distribution pattern of *Oithona davisae* abundance was observed in the study area and ANOVA showed that differences between the abundance of *O. davisae* were statistically important in sampling months ($F=14.961$; $p<0.05$), but not at sampling stations ($F=0.195$; $p>0.05$). *O. davisae* had greater mean abundance values during the summer-autumn periods. The highest value for mean abundance was observed in September 2015 (3533.33 ± 1194.89 ind. m^{-3}). A small-scale decrease in mean abundance was observed in the period from September 2015 to December 2015 (1715.05 ± 938.67 ind. m^{-3}). In January 2016, mean abundance values started to decrease sharply and reached 170.68 ± 52.74 ind. m^{-3} . After January 2016, the decreasing trend continued gradually until May 2016 (0.24 ± 0.58 ind. m^{-3}) and reached its lowest level in May 2016 (Figure 3).

O. davisae concentrated in the inshore of the bay and the abundance of this species decreased at stations located outer of the bay. The maximum abundance value of *O. davisae* was noted in September 2015 at station B (5550 ind. m^{-3}) and the minimum abundance value was determined in May 2016 at station D (1.67 ind. m^{-3}). While there were high abundance values at stations inside the bay during July–August–September 2015, the abundance of this species at stations outside of the bay reached high values during October–November–December 2015. *Oithona davisae* was found only at station D in May 2016 (Figure 3, Table 2).

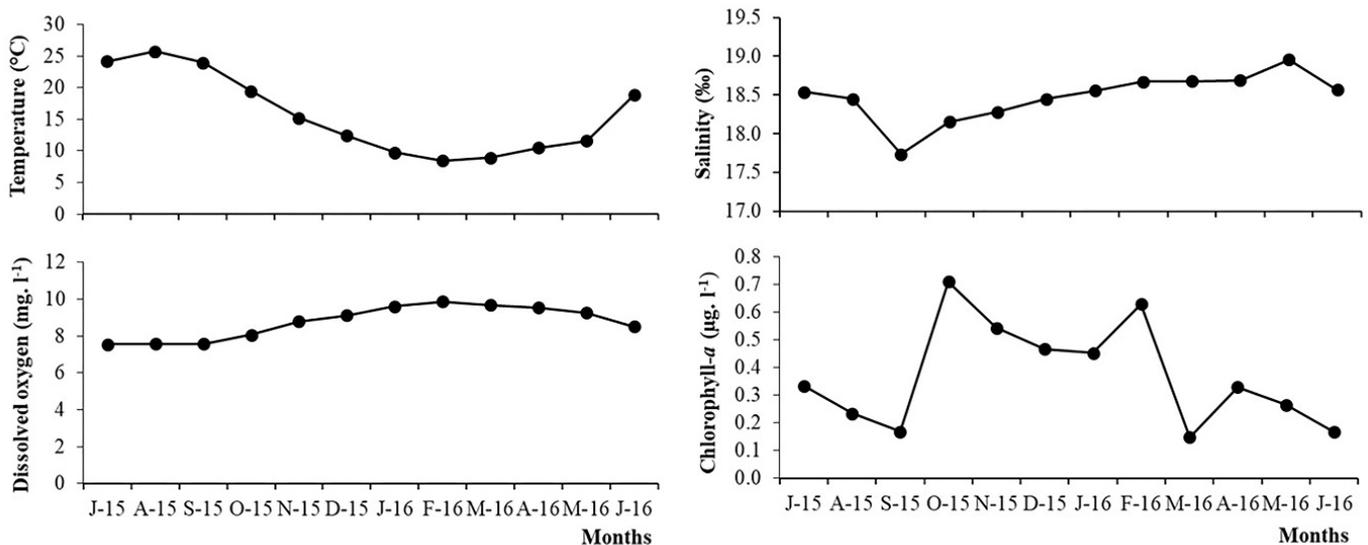


Figure 2. The monthly mean variation in environmental factors on surface seawater in Hamsilos Bay, Sinop.

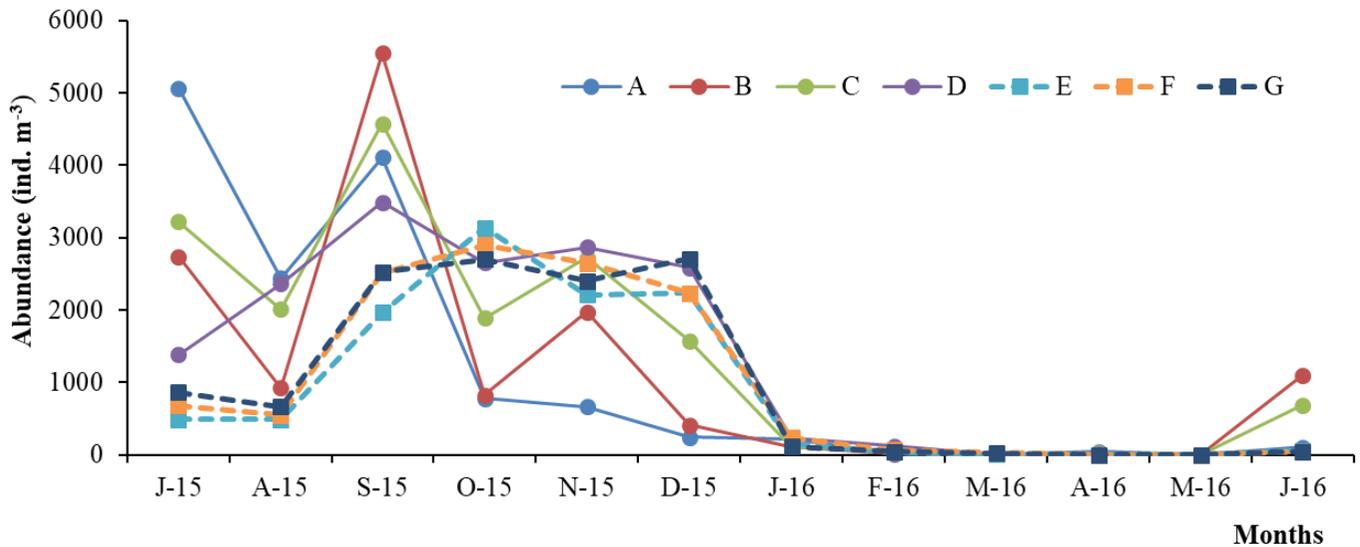


Figure 3. Spatial and temporal distributions in abundance (ind. m⁻³) of *O. davisae* in Hamsilos Bay.

Table 2. Minimum, maximum, and mean of abundance values (ind. m⁻³) of *O. davisae* according to stations in Hamsilos Bay.

Stations	A	B	C	D	E	F	G
Minimum	10 (March 2016)	4 (April 2016)	12.5 (March 2016)	1.67 (May 2016)	4.17 (April 2016)	4.17 (April 2016)	2.5 (April 2016)
Maximum	5062.5 (July 2015)	5550 (September 2015)	4575 (September 2015)	3487.5 (September 2015)	3137.5 (October 2015)	2900 (October 2015)	2712.5 (December 2015)
Mean±STD	1142±1683.7	1569±1568.6	1408±1459.5	1315±1321.7	901±1094.7	994±1145.6	1008±1147.1

Population age structure

The majority of the population of *O. davisae* consisted of copepodite 5–4 (Figure 4). The abundance values of females varied from 0.83 ind. m⁻³ (June 2016; St F and G) to 625 ind. m⁻³ (July 2015; St A). The presence of females was more pronounced in September 2015 and October 2015 (Figure 5). The female abundance values among months differed significantly ($F=11.850$; $p<0.05$). Between stations, there were no statistically significant variations in female abundance ($F=0.217$; $p>0.05$). Abundance values of the male were lowest in January 2016 (2.5 ind. m⁻³; St C) and highest in July 2015 (781.25 ind. m⁻³; St A). Three peak periods (July 2015, September 2015, and December 2015) were observed in males (Figure 5). Male abundance significantly differed between months ($F=5.899$; $p<0.05$). In terms of male abundance, there were no statistical differences between stations ($F=0.600$; $p>0.05$). The mean sex ratio (female/male) values of *O. davisae* varied between 0.56 and 3.79. Female individuals dominated the community from August 2015 to February 2016 (Figure 6).

Copepodite 5–4 was present all over the research period (in May 2016 only in station D), with its maximum abundance of 4825 ind. m⁻³ at station B in September 2015 and its minimum abundance of 1.67 ind. m⁻³ at station D in May 2016. Copepodite 5–4 was dominated during the July 2015–December 2015 periods (Figure 5). Abundance values of copepodite 5–4 showed statistically significant differences among months ($F=15.077$; $p<0.05$). There

were no statistically significant differences in copepodite 5–4 abundance among the stations ($F=0.255$; $p>0.05$) (Figure 5).

Copepodite 3–1 abundance values ranged between 1 ind. m⁻³ (February 2016; St B) and 650 ind. m⁻³ (November 2015; St F). It contributed significantly to the *O. davisae* population between September 2015 and December 2015 (Figure 5). Abundance values of copepodite 3–1 showed statistically significant differences among months ($F=13.723$; $p<0.05$). No significant differences in copepodite 3–1 abundance were observed between stations ($F=0.151$; $p>0.05$).

Female, male, and copepodite 3–1 individuals were not encountered in March 2016 and May 2016. In April 2016, female and copepodite 3–1 were determined only at station A, and male at station C (Figure 5). Minimum and maximum mean abundance values were in the range of 0.71 ± 1.75 (April 2016) to 317.26 ± 48.40 (September 2015) ind. m⁻³ in female, 0.24 ± 0.58 (May 2016) to 2818.75 ± 1163.20 (September 2015) ind. m⁻³ in copepodite 5–4, 1.43 ± 3.5 (April 2016) to 441.55 ± 176.88 (November 2015) ind. m⁻³ in copepodite 3–1 (Figure 5).

The abundance of developmental stages of *O. davisae* was well correlated with trophic elements and physiochemical variables. It was found that the abundance and distribution of developmental stages of *O. davisae* correlated negatively with salinity and

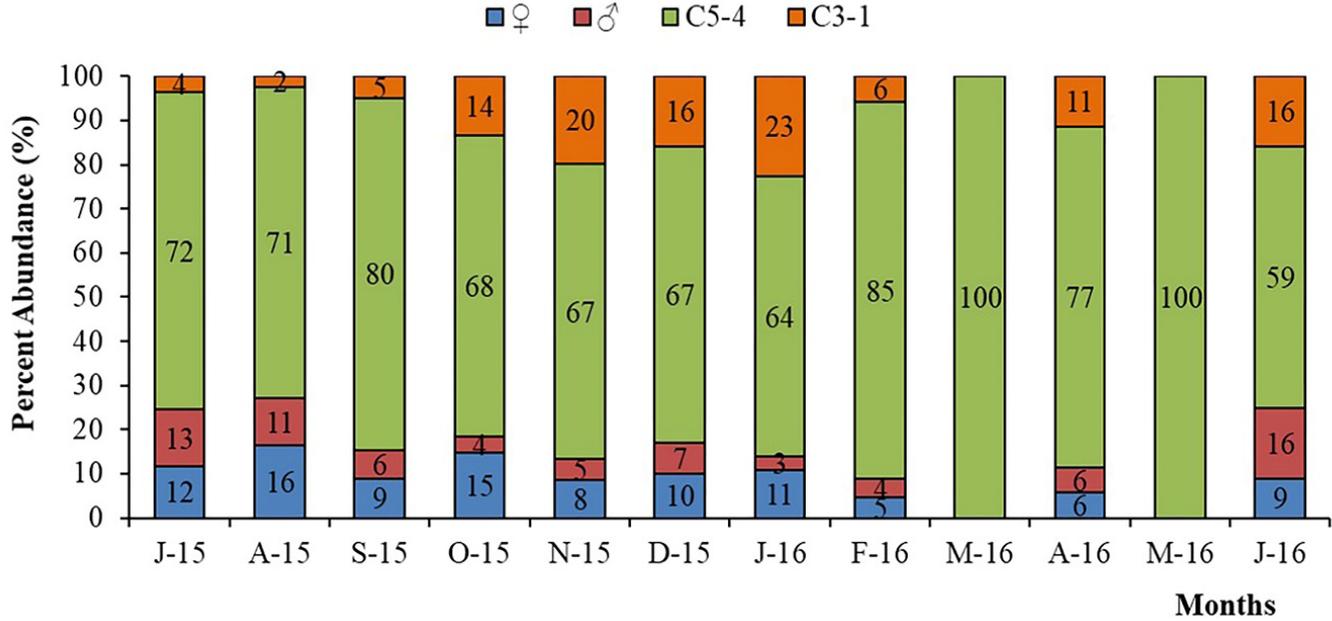


Figure 4. The percentage mean abundance contributions of different developmental stages of *O. davisae* in Hamsilos Bay.

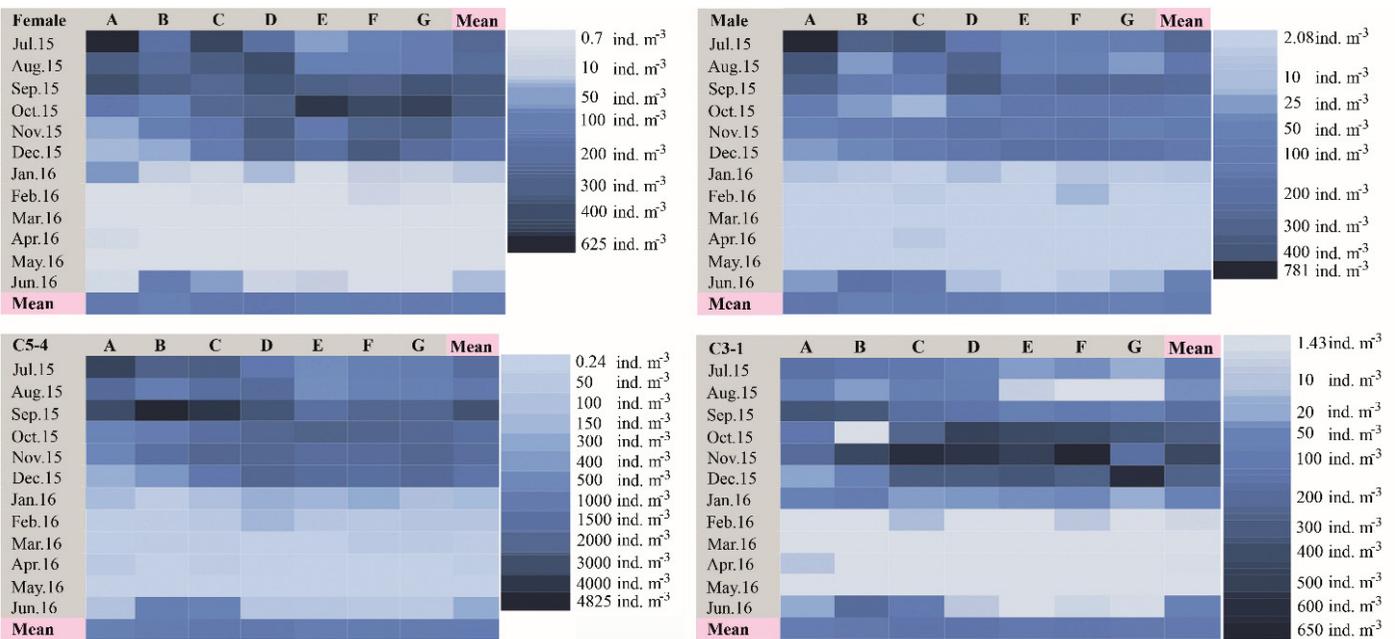


Figure 5. Temporal changes in mean abundance distributions of different developmental stages of *O. davisae* in Hamsilos Bay.

dissolved oxygen, but showed a positive correlation with chlorophyll-a and temperature. No correlation was found between the abundance of males and chlorophyll-a (Table 3).

DISCUSSION

Seven oithonid species were found in the Black Sea until now (Razouls et al., 2023). Most of these recorded oithonid species were known as questionable and had insufficient resources (Gu-

banova et al., 2014). Although *Oithona similis* are commonly distributed overall Black Sea (Gubanova et al., 2014), *Oithona davisae* expanded its distribution and became one of the most important components in the coastal community in the Black Sea (Gubanova et al., 2014), after the first observation of this species in Sevastopol Bay in the early 2000s (Zagorodnyaya, 2002). The abundance of this species in Sevastopol Bay already exceeded 40,000 ind. m⁻³ in October 2005 (Gubanova & Altukhov, 2007) and reached almost 100,000 ind. m⁻³ in October 2016 (Svetlichny

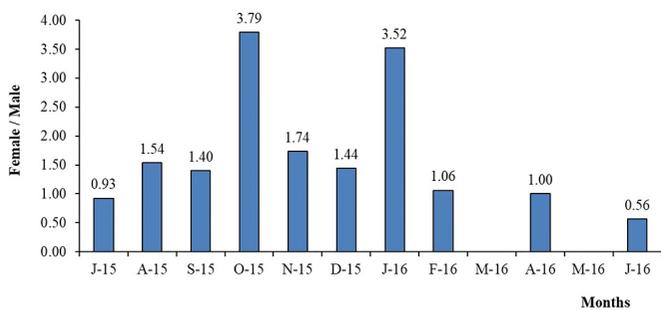


Figure 6. Mean sex ratio (male/female) values of *O. davisae* in Hamsilos Bay.

production style (Svetlichny et al., 2016), food source (principally feeds on ciliates and heterotrophic dinoflagellates) (Uchima, 1988; Saiz et al., 2003; Khanaychenko et al., 2018), locomotor activity and energy metabolism (body mass density, sinking, and swimming speeds) (Svetlichny et al., 2016).

Oithona davisae is a long-lived species and demonstrates considerable variation in population size depending on temperature changes affecting food resources and the development of planktonic predators (Svetlichny et al., 2016). It is either absent (Takahashi & Uchiyama, 2007) or very scarce (Uye & Sano, 1995) during the winter–spring seasons among the planktons of the coastal waters of East Asia, its original habitat. The abundance of this cy-

Table 3. Correlation between the abundance of *O. davisae* and developmental stages and environmental variables.

	<i>O. davisae</i>	Female	Male	Copepodit 5–4	Copepodit 3–1
Chlorophyll-a	.253 (*)	.232 (*)	.151	.246 (*)	.331 (**)
Temperature (°C)	.669 (**)	.752 (**)	.754 (**)	.685 (**)	.488 (**)
Salinity (‰)	-.771 (**)	-.795(**)	-.730(**)	-.810 (**)	-.734 (**)
Dissolved oxygen (DO mg. l⁻¹)	-.710 (**)	-.763 (**)	-.770(**)	-.738 (**)	-.509 (**)

*Correlation is significant at the 0.05 level (2-tailed); **Correlation is significant at the 0.01 level (2-tailed)

et al., 2018). Only eight years after the first appearance in the Black Sea, the evidence of the distribution of *Oithona davisae* was first observed on Turkey's coast in the Black Sea (Üstün & Terbiyik Kurt, 2016). It is seen that the presence of *O. davisae* is observed in the plankton every month and its abundance value increased when compared with the present study conducted in 2015–2016 in Sinop in respect of the species that were found to be present only in September–December period in Sinop in 2009. It was observed in other studies (Yıldız et al., 2017; Üstün et al., 2019; Yıldız & Feyzioğlu, 2020) carried out on the Turkish coasts of the Black Sea that the species reached values of maximum in the July–December (summer–autumn) period and demonstrated the lowest values in the January–May (winter–spring) period as is the case with the present study. Similar results were also revealed in the studies carried out on the coasts of Sevastopol Bay, Novorossiysk Bay, Romania, Bulgaria, and Georgia (Gubanov & Altukhov, 2007; Selifonova, 2011b; Timofte & Tabarcea, 2012; Mihneva & Stefanova, 2013; Seregin & Popova, 2016; Shvelidze, 2016; Svetlichny et al., 2018). Although the different plankton net sizes and sampling periods and depths do not allow us to compare the abundance values obtained from different studies, the studies demonstrated that the species has expanded its abundance and both temporal and spatial distribution since it was first found in the Black Sea.

In 2009, *O. davisae* was only distributed autumn period (Üstün & Terbiyik Kurt, 2016), however, it seems that *O. davisae* expanded its temporal distribution and was observed the following year on the Sinop coast. *O. davisae* to successfully inhabit and spread to new marine areas most probably due to having bioecological features, such as thermophilic (Ferrari & Orsi, 1984; Uye & Sano, 1995; Mihneva & Stefanova, 2013; this study), euryhaline (Svetlichny & Hubareva, 2014; Hubareva & Svetlichny, 2016), re-

clopoid copepod is low in the Black Sea during winter and spring and increases from summer to autumn with the increasing temperature (Gubanov & Altukhov, 2007; Selifonova, 2011b; Altukhov et al., 2014; Seregin & Popova, 2016). Capable of surviving the cold seasons in the Black Sea, fertilized females reproduce the next generation under favorable spring conditions (Svetlichny et al., 2016).

In the coastal waters of southeastern Japan, the original habitat of *O. davisae*, the species has been found to inhabit in seasonal temperature variations between 8.9 and 28.2 °C (Uye & Sano, 1995). The present study observed the highest abundance values between July 2015 and September–December 2015. The water temperature varied between 12–24 °C during this period. The lowest abundance values were recorded between February (8.4 °C–54 ind. m⁻³) and May (11.6 °C–0.24 ind. m⁻³). The results of this study agree with findings acquired in the Black Sea previously (Mihneva & Stefanova, 2013; Üstün & Terbiyik Kurt, 2016; Yıldız et al., 2017; Yıldız & Feyzioğlu, 2020). It was observed in the Black Sea that an intense increase in the population size of this species started when the surface water temperature reached 18–20 °C (Seregin & Popova, 2016). The analysis of seasonal variations of *O. davisae* recorded in samples collected from the coastal areas of Sinop suggested that temperature has a remarkable influence on the development time and abundance of the species. This study detected a strong positive correlation between temperature and *O. davisae* abundance. A negative correlation was calculated between *O. davisae* abundance values and surface seawater temperature in the seasonal study managed in İğneada (Üstün et al., 2019).

Exotic species that can successfully colonize new habitats are considered to have a wide salinity tolerance (Lee et al., 2003). *O.*

davisae has been detected in the salinity range of 28.6–32.3‰ in the Sea of Japan (Uye & Sano, 1995), in the salinity range of 12–19‰ in the San Francisco estuary (Ferrari & Orsi, 1984) and in the salinity range of 15.8–17.2‰ in the western Black Sea (Mihneva & Stefanova, 2013). In their experimental studies, Svetlichny & Hubareva (2014) and Hubareva & Svetlichny (2016) have shown that *O. davisae* can adapt well to high salinity and the species can live in the salinity range of 3–55‰ by keeping its body mass density constant and that *O. davisae* is (partly, 12–40‰) osmoregulatory species. The salinity ranged between 17.74–18.54‰ (mean: $18.27 \pm 0.27\%$) during the period when the values of the maximum abundance of the species were observed in our study. This study determined a negative correlation between salinity and abundance values of *O. davisae*. A positive correlation was calculated between the abundance values of *O. davisae* and surface seawater salinity in the seasonal study conducted in İçneada (Üstün et al., 2019). This contradictory report could be related to the joint and complex effect of the environmental variables.

In its original habitat, Asian waters, there is a congruence between the months with the highest abundance of the species and the months with the highest concentration of chlorophyll-*a* (Uye & Sano, 1995). It was revealed in our study that chlorophyll-*a* values ranged between 0.17–0.71 $\mu\text{g. l}^{-1}$ (mean: $0.41 \pm 0.19 \mu\text{g. l}^{-1}$) during the period when the highest abundance values of the species were observed. A positive correlation was detected between abundance values of *Oithona* and chlorophyll-*a* in this study, however, such a case was not detected in the study conducted in Trabzon (Yıldız et al., 2017).

CONCLUSION

O. davisae has a high tolerance to different environmental conditions and physiological structures (osmoregulatory systems, high egg production rate, high metabolic activity, etc.) which facilitate to survival and reproduction of *Oithona davisae* in the Black Sea. In our study, it seems that *O. davisae* has expanded its spatial and temporal distribution along the Sinop coast in the Black Sea and has been inhabited by increasing its population, and biotic and abiotic factors affected the temporal distribution of this species. This study suggests that *O. davisae* has been successful in inhabiting and establishing a constant population in the southern Black Sea.

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Effect of Pozzolanic Cement Exposure in Nile Tilapia (*Oreochromis niloticus*)

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ABSTRACT

Cement used in structures such as bridges, dams, and retaining walls built on the aquatic ecosystem harms different organisms. The current study reveals the effect of pozzolanic cement (PC) on Nile Tilapia (*Oreochromis niloticus*). The LC₅₀ value for *O. niloticus* exposed to 6 different PC concentrations for 24 hours was calculated as 306.66 mg L⁻¹. Increasing PC concentration significantly increased the pH of water (<10) at the end of the study (p<0.05). The increase in pH level showed a positive correlation with the increase in fish mortality (p<0.05). In addition, PC negatively affected the hematological parameters of fish. Thus, the current study reveals the negative effects of PC on *O. niloticus* in the acute period. Future studies should focus on developing cement with environmentally friendly materials that will not affect the pH level of water.

Keywords: Aquatic toxicology, hematology, LC₅₀, pozzolanic cement

INTRODUCTION

The construction industry is indispensable in that it meets the many visible and essential needs of humanity, such as new constructions, restorations, and post-earthquake reconstruction. Cement, as the basic material of the construction industry, is the second most produced material after steel, with a production of around 4.1 billion tons worldwide (Farfan et al., 2019). Since cement is the main material in the construction of many structures such as bridges, tunnels, dams, and retaining walls, the negative effects of possible effluents on natural water resources are inevitable (Kurtoglu et al., 2016). Even the cement wastes from construction sites that are far from the aquatic system eventually contaminate the aquatic ecosystems through rainwater and sewage systems. However, there are limited studies on the effect of cement in natural waters and habitats (Er & Kayış, 2022).

of aquatic organisms. In order to ensure the continuity of biodiversity, it is necessary to protect organisms in water resources against anthropogenic pollutants and to investigate their resistance (Bashir et al., 2020; Gadzała-Kopciuch et al., 2004). Possible pollutants in the aquatic environment affecting water quality parameters such as pH, turbidity and dissolved oxygen are the primary factors that affect the living standards of organisms (Austin, 1998). Cement, which is often used in the construction industry, poses a danger to aquatic organisms because it has the potential to increase the pH of water. In a study where cement increased the pH level above 10.5 it was reported that the mortality rate of rainbow trout (*Oncorhynchus mykiss*) individuals was around 60% (Er et al., 2019). In general, a pH value between 9 and 10 can be harmful to fish, while pH values above 10 can be lethal to the rest (European Inland Fisheries Advisory, 1969).

Natural water resources, which are generally the final destination of all pollutants, are the habitats

Physical and chemical changes in the aquatic environment cause measurable physiological

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changes in fish (Fazio et al., 2013). This is because fish are in contact with the aquatic environment, which significantly affects blood homeostasis. Biomarkers are defined as a change in biological response, ranging from molecular to cellular and from physiological responses to behavioral changes (Depledge et al., 1995). As an indicator of water quality, the hematological profile in fish has been recently evaluated as a rapid analysis (Akram et al., 2021; Minaz et al., 2022a). In particular, automatic hematological assay is a significant diagnostic alternative to conventional methods today (Fazio, 2019). Blood parameters provide useful information about various body processes and are of great importance in assessing the harmful effects of anthropogenic pollution on aquatic environments (Guerrera et al., 2021).

The effect of cement as an anthropogenic pollutant on aquatic organisms has been discussed in several studies (Adamu et al., 2008; Er et al., 2019; Er & Kayış, 2022). Unpublished case studies of cement-related mortalities in aquaculture facilities are known. In this context, the impact of cement on Nile Tilapia (*Oreochromis niloticus*), one of the most cultured species in the world, is very significant. The current study focuses on the toxic effects of pozzolanic cement (PC), which is frequently used due to its qualities of permanent durability, easy workability and high impermeability. In our study the hematological response of *O. niloticus* against the potential sub-lethal concentration at the end of the acute period (24 hours) was also investigated.

MATERIALS AND METHODS

The toxic material (Pozzolanic Cement)

In the present study, the toxic effect of a commercial pozzolanic cement (PC) was evaluated (CEM IV/B (P) 32,5 R, Oyak Çimento, Türkiye). Pozzolan materials do not have individual binding properties. However, they have high binding abilities in aqueous ambient with calcium hydroxide. The PC in the current study includes 65-89% natural clinker (30% clay and 70% limestone) and 11-35% pozzolan materials. Therefore, the potential toxic material in the available study has a content of Silicon dioxide (SiO_2), calcium oxide (CaO), aluminum oxide (Al_2O_3), and iron oxide (Fe_2O_3).

Experimental design, fish, and water quality

This study took place in the toxicology laboratory, Recep Tayyip Erdoğan University, Rize, Türkiye. Nile Tilapia fish (50.4 ± 4.4 g) were purchased from the Aquaculture Application and Research Center. The system design was triplicate configured with a 50 L aquarium and ten fish in each aquarium. For adaptation, the fish were placed in the aquariums ten days before the study. The study was carried out using rested tap water and in accordance with a 12-hour dark/light photoperiod. Aeration was distributed to the aquariums with air stones from a central pump system. Since PC is not a water-soluble material, it tends to aggregate at the bottom of the aquarium. In order to prevent this accumulation and keep the water homogeneous, a mixing system that provides water circulation was placed at the bottom of the aquarium. The fish were not fed in the acute period in order to provide stable water quality. All experimental studies were checked and approved by the Ethical Local Committee of the Recep Tayyip Erdoğan University (Decision No: 2021/14).

Water quality parameters were measured at the beginning of the study to ensure their suitability for the living standards of the fish. Temperature ($^{\circ}\text{C}$), pH and dissolved oxygen (mg L^{-1}) were measured with a portable multi-parameter (Hach, HQ40D 58258-00). Accordingly, water quality parameters were recorded as 17.1 ± 0.6 $^{\circ}\text{C}$, 6.51 ± 0.2 and 7.65 ± 1.4 mg L^{-1} for temperature, pH and dissolved oxygen, respectively.

Acute toxicity

Within the scope of the current study, the lethal concentration that killed 50% of the target population was determined. In this context, the fish were exposed to six different PC concentrations (100 mg L^{-1} , 200 mg L^{-1} , 300 mg L^{-1} , 400 mg L^{-1} , 500 mg L^{-1} and 600 mg L^{-1}) for 24 hours (Tablo 1). No PC was added to the control group but all other manipulative conditions were the same as for the PC groups. LC_{10} , LC_{50} and LC_{90} for 24 h values were determined as 255 mg L^{-1} , 306 mg L^{-1} , and 339 mg L^{-1} respectively.

Hematological analysis

Fish blood indicators were analyzed according to the automatic hematological assay method (Minaz et al., 2022a). Before the hematological analysis, five fish were randomly chosen from each group and transferred to an anesthesia tank that contained 60 mg L^{-1} clove oil. Afterward, 2.5 ml syringes were used to take blood samples from the caudal vein of the fish. Blood samples (0.3-0.5 mL) were fastish transferred to EDTA K3 tubes for the prevention of congelation. Finally, the following indicators were investigated using automatic hematological assay (Prokan-6800VET): leukocyte (WBC), erythrocyte (RBC), lymphocyte (LYM), monocyte (MID), granulocyte (GRAN), hemoglobin (HGB), hematocrit (HCT), mean corpuscular volume (MCV) mean corpuscular hemoglobin (MCH), and mean corpuscular hemoglobin concentration (MCHC). The blood counter device was calibrated with blood from healthy fish prior to the study.

Statistical analyses

In the current study, all data were shown in mean \pm standard deviation. The Kolmogorov-Smirnov test was used to determine normality distribution in the data set. According to the normality analysis, it was observed that all data sets showed normal distribution. Therefore, parametric tests were applied to determine statistical differences between the groups. Significant differences among groups were evaluated by a one-way ANOVA test. Also, the Tukey test was used to detect the differences in the groups. The Pearson Correlation test was performed to examine the statistical relationship between PC concentration, mortality rate and pH values. At the end of acute period, the Probit test was used to detect the LC_{50} value. P-values of less than 0.05 were considered statistically significant. All datasets were analyzed by SPSS 25 software package for Windows.

RESULTS AND DISCUSSION

Cement is the basic building block of many reinforced concrete structures in aquatic environments. When construction plans using cement are made to increase human welfare in an aquatic environment, most researchers typically focus on the fact that construction areas restrict the vital needs of organisms in the ecosystem, such as nutrition, reproduction and migration (Kocabaş et

Table 1. Lethal concentration of PC in *O. niloticus*.

Concentration (mg L ⁻¹)	Total fish (n)	Number of dead fish (mortality%)
100	30	0 (0%)
200	30	0 (0%)
300	30	12 (40%)
400	30	27 (90%)
500	30	30 (100%)
600	30	30 (100%)
LC ₁₀ (mg L ⁻¹)		255.08
LC ₅₀ (mg L ⁻¹)		306.66
LC ₉₀ (mg L ⁻¹)		339.40

al., 2013). Although many studies have been conducted on the effects of reinforced concrete structures on aquatic environments, the effect of cement on organisms has been ignored. Therefore, the current study focused on the effect of possible cement leakage from reinforced concrete structures on *O. niloticus* at the end of acute period.

Relationship between mortality rate and pH

The relationship between the mortality rate and pH are shown in Figure 1. Accordingly, while no death was observed up to 200 mg L⁻¹ PC concentration at the end of 24 hours, death was observed in 40% at 300 mg L⁻¹ PC, 90% at 400 mg L⁻¹ PC, and in all individuals at 500 and 600 mg L⁻¹. Additionally, higher pH levels (pH level of up to almost 11) were observed with increasing PC concentration, while the pH was around 6.5 in the control group. At the LC₅₀ level (306 mg L⁻¹), the pH level was estimated to be around 10.18. There was a significantly positive relationship between PC concentration, mortality rate and pH (Table 2, p<0.05).

Cement content tends to increase the pH level of water (Lee & Hur, 2005). However, studies have shown that this increase in pH level disappears within 24 hours in static water and returns to normal levels (Law et al., 2013). Therefore, a 24-hour LC₅₀ value

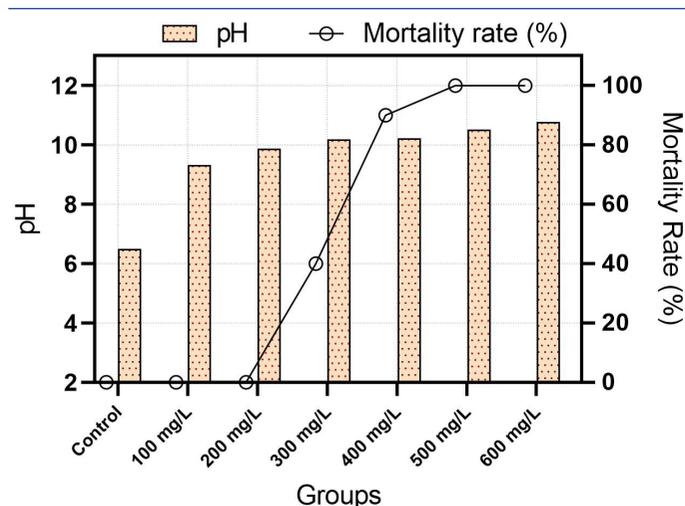


Figure 1. Mortality rate and pH depending on PC concentration.

Table 2. Correlation table between PC concentration, mortality rate, and pH.

		Concentration	Mortality Rate	pH
Concentration	Pearson Correlation	1		
	Sig. (2-tailed)			
Mortality Rate	Pearson Correlation	0.938**	1	
	Sig. (2-tailed)	0.002		
pH	Pearson Correlation	0.825*	0.759*	1
	Sig. (2-tailed)	0.022	0.045	

* Correlation is significant at the 0.05 level (2-tailed);
 ** Correlation is significant at the 0.01 level (2-tailed)

was determined in the current study. In a study conducted to determine the toxic effects of portland cement on tilapia (7.2 g), the LC₅₀ value was found as 41.21 mg L⁻¹ (Adamu et al., 2008). In the current study, this value was calculated as 307 mg L⁻¹. The difference between these two studies involving the same species may be due to the type of cement or the difference in fish size (34.8 g). In another study, the 96-hour LC₅₀ value in rainbow trout exposed to different concentrations of PC was 440 mg L⁻¹ (Kurtoglu et al., 2016). The reason for the high 96-hour LC₅₀ value in the previous study may be related to the fact that the temperature was lower than in our study and more suitable for rainbow trout. It would be expected that the 24-hour LC₅₀ value for rainbow trout, which is a more sensitive fish than tilapia, would be much higher.

In an aquatic ecosystem, pH is vital to organisms and can affect the solubility and toxicity of chemicals in the water. Although most aquatic organisms live in the pH range of 6.5-9, there may also be organisms living outside this range. However, high or low pH levels have negative effects on growth and reproduction. Cement tends to increase pH levels, resulting in toxicity to aquatic organisms. For instance, the increase in pH level caused a decrease in the EC₅₀ (median-effective concentration) value for *Daphnia magna* (Rodrigues et al., 2020). In Atlantic salmon eggs, a low pH level (pH 2.78) caused the death of 98% of all individuals after 18 hours, while a high pH level (12.23) caused all individuals to die within 15 minutes (Foldvik et al., 2022). The toxicity ability of high pH level is much more effective. In the current study, the pH level showed a positive relationship with increasing concentration in the first 24 hours. However, pH levels in all groups returned to levels similar to the control group after 24 hours. In our previous study, a different group was designed whose pH level was reduced with HCL to reveal the toxic driving force of cement (Er & Kayis, 2022). At the end of the study, it was observed that there was no death in the group in which the pH level was manipulated. Therefore, it has been proven that the

main driving force in cement toxicity studies is pH. The lethal effect of high pH level on organisms will contribute to the management of the contamination process both in nature and in aquaculture ambients. However, today's common belief is that in cases of cement-related contamination, losses should be determined by sampling fish rather than the characteristics of the water. Additionally, there is a deficiency regarding the time period for providing water samples.

Hematological evaluation

For hematological studies, 100 mg L⁻¹ PC (T100) and 200 mg L⁻¹ PC (T200) concentration groups, which were 1/3 and 2/3 times the LC₅₀ value and in which all fish survived, were evaluated (Figure 2). Accordingly, significant differences were observed between groups for all hematological parameters except MCV. Significant decreases were observed between all groups in WBC values depending on increasing concentration ($p < 0.01$). PC groups showed significantly higher RBC, HGB, and HCT values compared to the control ($p < 0.01$). For LYM value, the T200 group was significantly lower than the other groups ($p < 0.05$). Finally, significantly lower values were observed in the PC groups for MID, GRAN, MCH, and MCHC values ($p < 0.01$).

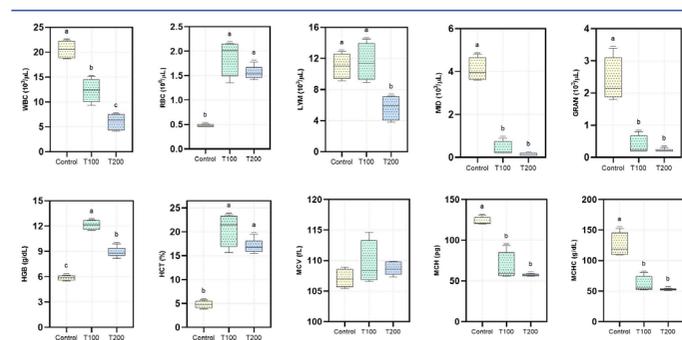


Figure 2. Hematological parameters of *O. niloticus* exposed to different PC concentrations.

One of the most important and rapid indicators for aquatic organisms encountering any toxic pollutant is hematological outputs (Ak et al., 2023; Fazio et al., 2017). This is because blood represents fish health depending on exposure to environmental stressors (Gabriel et al., 2011; Minaz et al., 2022a). Blood parameters may differ from normal levels in aquatic organisms during exposure to pollutants, transportation and anesthesia (Minaz et al., 2022b). In the current study, significantly decreased WBC values were noted in the increasing PC groups. WBCs are circulatory cells that assist in physiological responses and constitute different cell types such as lymphocytes and monocytes. Therefore, like WBC, lower MID and LYM in PC groups is an expected phenomenon. Although previous studies have rarely shown a tendency for WBC to increase in response to stressors, WBC tends to decrease in fish against many stressors (Esmaili, 2021). Under stress conditions, the immune system is affected positively or negatively by many neuroendocrine elements (Khansari et al., 1990). The decrease in WBC, LYM and MID may be directly related to the decrease in cell production of the hematopoietic or-

gans or in total leukocytes of the immune defense (Alam et al., 2021). It has previously been reported that exposure to herbicides as toxic substances reduces WBC values for the same fish species (Abdel-Warith et al., 2021). In general, exposure to toxic substances results in decreased leukocyte functions, subsequent inhibition of lymphocyte responses, apoptosis and cell death (Guzzi et al., 2012). RBC absorbs oxygen in the gills and carries it to the tissues. RBC in fish varies between $0.4 \times 10^6 \text{ mm}^{-3}$ and $5.2 \times 10^6 \text{ mm}^{-3}$ (Esmaili, 2021). In the current study, this value remained within the general range. The immune system of fish may increase RBC under stress or due to infection. This is a defense mechanism to fight infection or toxic substances and it speeds up recovery. Like WBC, HGB is at higher levels in PC-exposed groups. The increase of HGB and HCT in fish may occur in order to increase the body's oxygen carrying capacity and respond to oxygen needs. However, this increase may vary under a particular fish species or in certain environmental conditions.

CONCLUSION

The release of cement, the basic building block of reinforced concrete structures, into the aquatic ecosystem has become an inevitable issue. Cement effluent entering the aquatic environment as a result of many developments such as agricultural activities and energy systems poses a danger to both the environmental environment and all organisms. In this context, the current study revealed the effects of pozzolanic cement exposure on Nile tilapia (*Oreochromis niloticus*). PC, which has a vital importance in the aquatic ecosystem due to its tendency to increase the pH level, had a toxic effect on *O. niloticus*. It also negatively affected the hematological system of the fish and created differences in blood parameters as a stress response. This study revealed the lethal and physiological effects of PC exposure during the acute period. Case studies are important to raise awareness by showing how the release of toxic PC material into natural waters affects the aquatic environment around the world. Future studies should focus on improving PC with supporting materials that do not increase the pH of water. In addition, using the current study as a reference, awareness should be raised for the construction industry and the use of environmentally friendly materials should be encouraged.

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Ethical Declaration: The current study was checked and approved by the Ethical Local Committee of the Recep Tayyip Erdogan University (Decision no:2021/14).

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Investigation of Benthic Macroinvertebrate Fauna and Water Quality of Animal Drinking Water Troughs by Multivariate Statistical Methods: The Case of Şarkışla

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ABSTRACT

This study was conducted in 2022 to investigate the water quality and benthic macroinvertebrate fauna of water troughs in Şarkışla, a district in the city of Sivas. Fifteen stations were selected from the most used water troughs in the district and sampling was carried out in the spring, summer, and autumn seasons during peak agricultural activities. As a result, a total of 17 taxa were identified: 4 taxa belonging to Oligochaeta, 6 species belonging to Chironomidae, 2 taxa belonging to Amphipoda, and one each belonging to Gastropoda, Hemiptera, Coleoptera, Ostracoda and Trichoptera larvae. All taxa identified are new records for the study area. In addition to benthic sampling, some physicochemical parameters of water (water temperature, pH, electrical conductivity, dissolved oxygen, total hardness, calcium, magnesium, chloride, salinity, total dissolved solids, phosphate, sulphate, nitrate nitrogen, nitrite nitrogen) were also analyzed. The similarity of the stations in terms of physicochemical parameters was analyzed by utilizing the Bray-Curtis similarity analysis. Accordingly, stations 2 and 4 were found to be the most similar, by 96.91%, and stations 3 and 13 were found to be the most different, by 66.46%. In addition, the relationship between the investigated physicochemical parameters was analyzed using the Pearson and Spearman Correlation Analysis. The relationship between the species and physicochemical parameters was revealed using the CCA Analysis.

Keywords: Water troughs, benthic macroinvertebrates, physicochemical parameters, statistical methods

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INTRODUCTION

Water troughs are artificially transformed examples of natural water sources. Water troughs are found all over Turkey. Sometimes they are located in villages and sometimes they are found on the roadsides. They were first built to meet the water needs of animals. They can also be used to control water flow, agricultural irrigation, drinking water, and groundwater (Külköylüoğlu et al., 2013; TODAİE, 2011). Water is vital for all living creatures on this planet. In animals, body temperature regulation, nutrient uptake, and waste removal occur with wa-

ter. Water troughs are artificial water structures created to meet the water needs of animals. They are open-water storage structures of various sizes to accumulate and hold water for animal consumption (Denktaş, 2000; Tay, 2011). These structures are an important source of water for domestic cattle and sheep in cultivated areas and also for wild animals such as wolves, deer, rabbits, birds, etc. in rural and wild areas (Özer, 2010). These water structures, especially in rural areas, are very important to meet the water needs of animals while grazing. Depending on the number of animals relying on the water trough, they can be of different sizes. Water



troughs are sometimes located in forested areas, sometimes on higher parts of the landscape, and sometimes on roadsides. The water in these structures is open to contamination. Especially in the summer months algae layers can form on the inside of the water troughs, due to increased weather temperature and stagnation of the water body.

Benthic macroinvertebrates are often used as indicators of the biological status of water bodies. They are reliable indicators as they spend all or most of their life in water. They are easy to collect due to their high tolerance of pollution (Kazancı et al., 1997). Benthic macroinvertebrates are usually found attached to rocks and vegetation, or they are buried in the sand and sediments at the bottom of the water trough.

To date, studies conducted on water troughs belong to Özkan (2006a; 2006b), Külköylüoğlu ve ark. (2012), Külköylüoğlu et al. (2013), Başak et al., (2013); Külköylüoğlu et al., (2017); Özer & Dikmen (2021); Ataman et al., 2023; Sanbur (2023).

This study aims to determine the water quality and benthic macroinvertebrate fauna of water troughs in the district of Şarkışla using multivariate statistical analyses.

MATERIALS AND METHODS

Study area

Şarkışla is located in the southwest of Sivas province, within the borders of the Upper Kızılırmak region of central Anatolia. Its surface area is 2250 km wide and is at an altitude of 1180 m above sea level.

Şarkışla is the 7th largest district in terms of surface area among the 17 districts of Sivas (<https://sarkisla.gov.tr>). The economy of Şarkışla is based on agriculture and animal husbandry. The district offers a rich variety of agricultural products due to the presence of large agricultural land and agricultural basins. Field crops include cereals, fodder crops, industrial crops, and edible legumes. Sugar beet and potatoes are among the most cultivated crops. The district creates a suitable environment for animal husbandry with its climate and the presence of pasture areas. Cattle and ovine breeding is common in the district (Oran Kalkınma Ajansı, 2017). Animal husbandry is intensively practiced in the region where this research was conducted. Natural water troughs are made of concrete, metal or wood and are generally rectangular in shape. Water troughs exist for the benefit of both humans and animals.

The coordinates of the selected sampling stations and information about the water troughs are detailed in Table 1.

Sampling

The study was carried out at 15 stations in the spring, summer and autumn of 2022, during peak agricultural activity. The stations could not be reached in winter due to bad weather conditions. For this reason, sampling could not be performed during the winter season. pH levels, total dissolved solids, and electrical conductivity of the water samples were measured and recorded during the field study. To analyze other physicochemical parameters, water samples were taken in dark-coloured glass bottles of

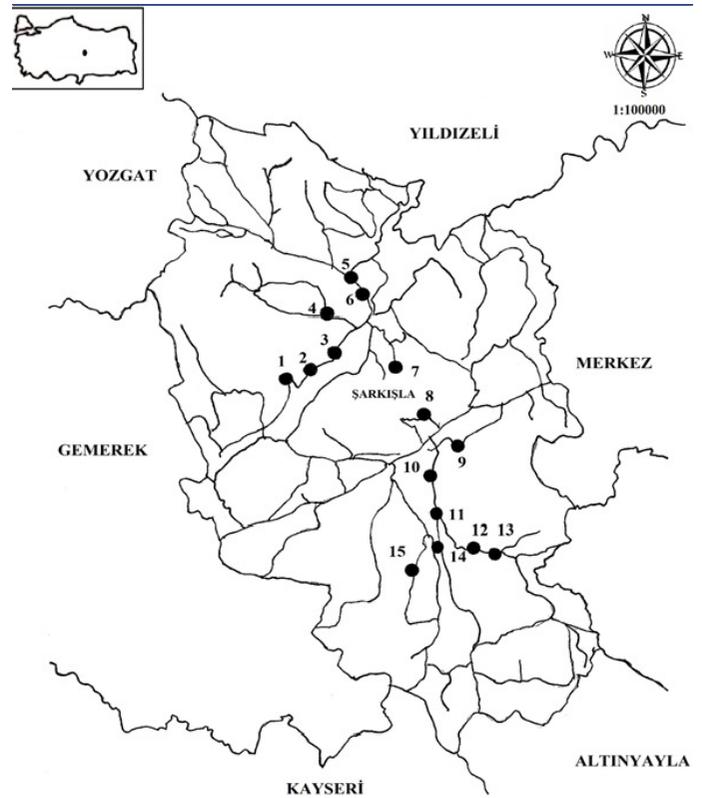


Figure 1. Study area and sampling locations.



Figure 2. Two-partition concrete water trough.

1 liter. At the same time as taking water samples, benthos samples were also collected from the bottom part of the water troughs using hand mud scoops of different sizes. The samples were stored and preserved with 70% ethyl alcohol.

The physicochemical parameters of the water were measured using various titrimetric and spectrophotometric methods (Egemen & Sunlu, 1999).

The benthos samples brought to the laboratory were separated from the debris and placed in 70% ethyl alcohol for species iden-

Table 1. Geographical coordinates and information about the water troughs sampled.

Though Number	Stations	Coordinates	Information about the water troughs
1	Çiçekliyurt	39°26'00"N 36°17'58"E	2- partitions concrete water trough
2	Alaçayır	39°27'08"N 36°19'48"E	3- partitions metal water trough
3	İlyashacı Village	39°27'46"N 36°19'58"E	2- partitions concrete water trough
4	Emlakkaracaören	39°29'24"N 36°21'28"E	Karacaören Meryemin Ahmet Fatma Cemal Koçtürk Hayratı, 4- partitions concrete water trough
5	Akdağmadeni Şarkışla roadside water trough	39°25'50"N 36°24'19"E	2- partitions concrete water trough, surrounded by agricultural land
6	Belen Tepesi roadside water trough	39°25'04"N 36°24'01"E	Muhlis Doğan Hayratı, 3- partitions concrete water trough
7	Cemel Altınyayla road	39°19'12"N 36°30'11"E	5- partitions concrete water trough, surrounded by agricultural land
8	Elmalı Village entrance	39°26'00"N 36°17'58"E	3- partitions concrete water trough, surrounded by agricultural land
9	Maksutlu Village	39°21'38"N 36°27'40" E	Kalembey Oğullarından Kaya Kızı Nalan Polat Hayratı, 2- partitions concrete water trough
10	Üyük (Hüyük) Village	39°22'21"N 36°17'58"E	5- partitions concrete water trough
11	Döllük Village	39°20'80"N 36°27'89"E	2- partitions concrete water trough
12	Cemel roadside water trough	39°18'16"N 36°27'36"E	Asım oğlu Ahmet Demirci ve Cemel Belediyesi Hayratı, 4- partitions concrete water trough, surrounded by agricultural land
13	Cemel Village road	39°20'09"N 36°26'20"E	4- partitions concrete water trough. In addition, daily agricultural labourers live in the surrounding area.
14	Hamlar	39°17'55"N 36°27'24"E	4- partitions concrete water trough, surrounded by agricultural land. Wheat, barley, chickpea, and sunflower are among the most cultivated crops.
15	Aşılık (Osmanpınarı)	39°16'08"N 36°27'48"E	4- partitions concrete water trough, surrounded by agricultural land. Wheat and barley are among the most cultivated crops.

tification. The collected benthos samples were identified to the smallest taxonomic group possible. Kathman & Brinkhurst (1998), Milligan & Michael (1997) and Timm (1999) were consulted for identification of Oligochaeta. Oliver et al., (1978), Saether (1980), Cranston (1982), Pinder & Reiss (1983), and Fittakau & Roback (1983) were consulted for the identification of Chironomidae. For the identification of Amphipoda, the help of subject experts was obtained.

Statistical analyses

All obtained results were transformed by LogBase10 in Microsoft Office Excel 2010 and SPSS 9.0 for Windows to use statistical techniques (Krebs, 1999). The relationship between the physicochemical parameters was evaluated by applying the Spearman and Pearson Correlation Analysis (Krebs, 1999). The similarity of physicochemical parameters in terms of stations was analyzed by Bray-Curtis Cluster analysis using BioDiversity Pro 2.0 (McAleece et al., 1997). The statistical relationship between the benthic macroinvertebrates and environmental parameters Canonical Correspondence Analysis (CCA) was determined using the Past Version 3.14. (Hammer et al., 2001).

RESULTS AND DISCUSSION

A total 17 taxa were identified in this study. These included four taxa belonging to Oligochaeta, six to Chironomidae, two to Amphipoda, and one each to Gastropoda, Hemiptera, Coleoptera, Ostracoda and Trichoptera larvae. All taxa identified are new records for the study area.

In the study, the habitat and station information on taxa as well as information about seasons is as follows. The highest number of taxa was found in the spring season (14 taxa), followed by the summer (10 taxa) and finally the autumn (7 taxa). *Nais elinguis* are abundant in organically polluted waters. They can be found in fast-flowing rivers as well as in muddy, foul-smelling, poorly oxygenated, and cold waters under wide environmental conditions (Timm, 2003). In the study, *Nais elinguis* were found in the spring season at stations 4 and 6. *Nais simplex* is an almost cosmopolitan species, found in fresh waters (Timm et al., 2002). *N. simplex* was found in water samples collected from soft muddy ground. *N. simplex* was found only at station 11 in the spring and summer seasons. *Chaetogaster diaphanus* is found among plants in brackish and fresh waters and in rivers with generally clear water. It is a predatory species that eats small invertebrates such as Oli-

gochaetae, Chironomidae larvae, Cladocera and Copepoda (Timm, 2003). *C. diaphanus* was found in the spring season at station 11. Timm (2013) reported that species belonging to the genus *Potamothenix* sp. are present in freshwater, brackish water and euryhaline forms. *Potamothenix* sp. was found in the spring season at station 13.

Chironomus riparius is a common species and usually lives in eutrophic conditions (Thienemann, 1954). *C. riparius* was found at stations 2, 6, 9, 11, 12 in the spring season (Table 2). It was found at stations 1, 8, 13, 14 in the summer season, and at station 2 in the autumn season (Table 2). The most common species encountered during sampling was *Chironomus anthracinus* (Table 2). This species is the most tolerant species of the Chironomidae family and is very common (Taşdemir et al., 2009). In this study, *C. anthracinus* was found in all three seasons and in all stations. *C. anthracinus* was found at stations 3, 5, 6, 8, 9, 10, 11, 12, 14 and 15 in the spring season (Table 2); at stations 1, 2, 4, 7, 8, 9, 12, 13, 14 and 15 in the summer season (Table 2); and at stations 2, 3, 4, 5, 10, 12 and 13 in the autumn season (Table 2). *Chironomus viridicollis* lives in the muddy parts of stagnant waters (Özkan 2006a). In the study, it was detected at stations 8 and 9 in the spring season and at station 8 in the summer season (Table 2). Taşdemir et al., (2010) reported that *Cricotopus intersectus* was found in stagnant waters. In this study, *C. intersectus* were found at stations 1, 3, 5, 9, 10, 11, 12 in the spring season; at stations 1, 2, 3, 5, 9, 10, 11, 14 in the summer season, and at station 8 in the autumn season (Table 2). Özkan (2007) reported that *Paracladius conversus* is seen on mud-aquatic stream plants. *P. conversus* was found at station 3 in the spring and at station 7 in the sum-

mer seasons. Özkan (2007), reported that *Tanytarsus gregarilus* is found in mud in stagnant waters, and found in sandy, muddy, and stony ground in streams. *T. gregarilus* was found at stations 3, 6, 9, 13, 14, 15 in the spring season (Table 2). In the summer season, it was found at stations 1, 7, 8, 14, 15 (Table 2).

Gastropods are more abundant in spring and summer. Their tolerance to very hard and salty waters is quite high (Robert & Diillon, 1999). Gastropoda individuals were found at station 8 in the spring; at station 15 in the summer; and at stations 12 and 13 in the autumn (Table 2). *Niphargus* sp. is reported to be the largest of all freshwater amphipod genera (Väinölä et al., 2008). Usually, these individuals inhabit caves or groundwater (Karaman & Ruffo, 1986; Sket, 1999). *Niphargus* sp. was found at station 1 in the autumn season. *Gammarus pseudosyrithia* has a wide tolerance range to temperature (usually 5–21 °C), so it was reported that this adaptation to different water temperatures is the main reason for its wide presence (Zamanpoore et al., (2011). *G. pseudosyrithia* was found at station 5 in the the spring season (Table 2). Hemipterans are found in lentic habitats or in backwater or pool areas of streams (<http://lakes.chebucto.org/>). Hemiptera was found at station 12 in the autumn season. Coleopterans larvae are organisms sensitive to pollution. They can be found in moderately polluted waters (Haldar et al., 2016). Coleoptera was found at station 5 in the spring season (Table 2).

Ostracods live in all kinds of aquatic environments including the base of oceans, in rivers, in lakes and even in swamps. Some ostracods live in temporary water bodies and can survive when the pond dries up in summer (www.bgs.ac.uk). Ostracoda was found

Table 2. Benthic macroinvertebrate taxa identified from the water troughs and their seasonal distribution according to stations.

Taxonomic Groups /Seasons	Spring Season	Summer Season	Autumn Season
Oligochaeta			
<i>Nais elinguis</i> Müller, 1773	4,6	-	-
<i>Nais simplex</i> Piquet, 1906	11	11	-
<i>Chaetogaster diaphanus</i> (Gruithuisen, 1828)	11	-	-
<i>Potamothenix</i> sp.	13	-	-
Chironomidae			
<i>Chironomus riparius</i> Meigen, 1804	2,6,9,11,12	1,8,13,14	2
<i>Chironomus anthracinus</i> Zetterstedt, 1860	3,5,6,8,9,10,11,12,14,15	1,2,4,7,8,9,12,13,14,15	2,3,4,5,10,12,13
<i>Chironomus viridicollis</i> Wulp, 1877	2,9	8	-
<i>Cricotopus intersectus</i> (Staeger,1839)	1,3,5,9,10,11,12	1,2,3,5,9,10,11,14	8
<i>Paracladius conversus</i> (Walker, 1856)	3	7	-
<i>Tanytarsus gregarilus</i> Kieffer, 1909	3,6,9,13,14,15	1,7,8,14,15	-
Gastropoda	8	15	12,13
Amphipoda			
<i>Niphargus</i> sp.	-	-	1
<i>Gammarus psedosyrithia</i> Karaman&Pinkster, 1977	5	-	-
Hemiptera	-	-	12
Coleoptera	5	-	-
Ostracoda	-	7,10	3,7,12,13
Trichoptera larvae	3,8	7	-
Total number of taxa	14	10	7

at stations 7 and 10 in the summer season (Table 2). It was found at stations 3, 7, 12 and 13 in the autumn season (Table 2). Trichopterans are indicator species of clean waters. They live in all kinds of aquatic environments, including rivers, lakes, ponds, and swamps (Haldar et al., 2016). Trichoptera larvae were found at stations 3 and 8 in the spring and at station 7 in the summer (Table 2). The benthos samples from the water troughs and the detected species are given in Table 2.

During the study period, water temperature values were measured between 6 °C (spring season, stations 3, 7, 10, 12, 13) and 16.5 °C (summer season, station 13) (Table 3). pH values were found to vary between 7.23 (spring season, station 13) and 8.67 (autumn season, station 11) (Table 3). Electrical conductivity values were found to vary between 359 $\mu\text{S}/\text{cm}$ (autumn season, station 7) and 994 $\mu\text{S}/\text{cm}$ (spring season, stations 8 and 13) (Table 3). During the study period, dissolved oxygen value was measured between 1.52 mg/L (autumn season, station 12) and 6.66 mg/L (spring season, stations 5 and 15) (Table 3). TH values were recorded between 18 FS° (spring season, station 15) and 37 FS° (spring season, station 2). Ca values were found to vary between 5.20 mg/L (autumn season, station 11) and 123.9 mg/L (summer season, station 5) (Table 3). Mg values were found to vary between 1.22 mg/L (spring season, station 12) and 34 mg/L (summer season, station 13) (Table 3). Cl values were found to vary between 0.99 mg/L (summer season, stations 2 and 3) and 53.9 mg/L (summer season, station 10) during the study period (Table 3). Salinity values were measured between 0.008 ‰ (autumn season, 2nd, 3rd, 6th stations) and 0.09 ‰ (spring, 10th station) (Table 3). TDS values were found to vary between 140 ppm (spring, 7th and 10th stations) and 486 ppm (spring season, 8th station) (Table 3). PO₄ values varied between 0.0005 mg/L (autumn season, station 1) and 0.87 mg/L (autumn season, station 11) (Table 3). It was found that the measured SO₄ values varied between 0.12 mg/L (autumn, station 1) and 5.34 mg/L (autumn season, station 8) (Table 3). NO₃-N values varied between 5.83 (summer season, 11th station) and 54.5 mg/L (summer season, 13th station) (Table 3). The NO₂-N values measured in the study varied between 0.002 mg/L (spring season, 11th station) and 0.65 mg/L (summer season, 4th station). (Table 3). Some physicochemical data measured from the water troughs according to seasonal average values of the stations are presented in Table 3.

According to TSWQR (2016), 1st class refers to high-quality or "very good" water. This quality of water can be used for animal production and farm needs. When the results obtained are evaluated, it is determined that all stations are of 1st class quality in terms of pH, sulfate and water temperature (SKKY, 2004). In terms of electrical conductivity, the stations are between 1st and 2nd class water quality (TSWQR, 2016). In terms of NO₃-N, it was determined that stations 2, 5, 8, 9, 10, 11 and 12, were of 3rd class water quality and the other stations were of 4th class water quality. The high NO₃-N values of the waters may be attributed to the presence of agricultural land around some stations. Leaving animal faeces in grazing areas can also increase the amount of nitrate in groundwater. In terms of dissolved oxygen, stations 3, 4 and 8 are in the 4th class water quality grade and the other stations are of between 2nd and 3rd class water quality. When the

chloride values in the water samples were examined, it was found that stations 9 and 10 were between 1st and 2nd class water quality grades and the other stations were of 1st class water quality (TSWQR, 2016).

D.O., Ca, Salinity and NO₃-N values were found to be high in the spring season. W.T., pH, E.C., Mg, Cl, PO₄, SO₄ were found in high amounts in the summer season (Table 4). As the air temperature increases in the summer, the water temperature also increases. The heated water dissolves the rock layers through which it passes and causes both pH and E.C. values to be higher. It was determined that T.H., TDS, NO₂-N values were high in the autumn season. Average values according to seasons and water quality parameters are given in Table 4.

When the average physicochemical values of the sampling stations are compared using the Bray-Curtis similarity index it is found that stations 2 and 4 are the most similar stations (96.91%), followed by station 9 (96.72%). Stations 3 and 13 are the least similar stations (66.46%) (Figure 3). Stations 2 and 4 are the most similar stations. It is very likely that the water coming to these stations comes from the same source. The elevation of the 3rd station is higher than that of the 13th station. The difference in elevation may impact water properties depending on the rock layers the water is passing through.

The average physicochemical values across the three seasons are also analyzed using the Bray-Curtis similarity index. Accordingly, spring and summer seasons were 95.90% similar, followed by the autumn season with 95.89% similarity to both spring and summer. Summer and autumn seasons were found to be the least similar seasons with a rate of 95.82% (Figure 4). It was determined that the similarity is very high across all three seasons.

The taxa and physicochemical values were also analyzed utilizing the CCA Analysis. The eigenvalues of the first two axes were calculated as 0.486 and 0.388, respectively. In the analysis, the two axes explain 40.59% of the variance of the species, 22.56% (Axis 1) and 18.03% (Axis 2). The distributions of *N. elinguis*, *Potamothrix* sp., and Gastropoda were affected by NO₂-N, NO₃-N, TDS, E.C., and salinity. Environmental variables do not affect the distributions of *C. intersectus*, *P. conversus*, *G. psedosyriacus*, Coleoptera, and Trichoptera. While W.T., Cl, PO₄, D.O., SO₄, and pH are effective in the distribution of *C. riparius*, *C. viridicollis*, *N. simplex*, *C. diaphanus*, *Niphargus* sp., Ostracoda, and Hemiptera; T.H., Mg, and Ca are not effective. The species *C. anthracinus* is affected by T.H., Mg, and Ca (Figure 5).

In addition, Pearson correlation analysis was applied to the normally distributed physicochemical values. Accordingly, a significant positive correlation was found between Ca and TH levels. Similarly, a significant negative correlation was found between pH and Ca levels. The relationship and correlation coefficients between the physicochemical parameters analyzed are given in Table 5.

Finally, the Spearman correlation analysis was applied to the values that did not show a normal distribution. Accordingly, a significant negative correlation was found between D.O. and NO₂-N; NO₂-N and PO₄; W.T. and D.O. There is a significant positive correlation be-

Table 3. Physicochemical data measured at sampling stations in terms of seasons.

Parameters	Unit	1 st	2 nd	3 rd	4 th	5 th	6 th	7 th	8 th	9 th	10 th	11 th	12 th	13 th	14 th	15 th
Spring Season																
W.T.	°C	7	7.5	6	6.5	7	7	6	7	6.5	6	7	6	6	6.5	6.5
pH		7.85	7.70	7.89	7.73	7.64	7.69	7.72	7.68	7.63	7.78	7.83	7.55	7.23	7.40	7.79
E.C.	µs/cm	398	652	366	645	521	394	362	994	455	701	381	398	994	655	408
D.O	mg/L	4.76	4.18	3.80	3.42	6.66	4.32	5.71	2.85	4.76	5.90	5.52	5.71	5.33	6.09	6.66
T.H	°F	20	37	19	32.8	27.6	21.6	20	35.6	27.8	27	18.6	31.2	35.2	28.4	18
Ca	mg/L	48	59.3	38.4	83.36	96.19	73.74	68.13	102.6	84.16	83.36	50.55	76.15	103	74.54	48.08
Mg	mg/L	6.83	5.42	4.73	12.29	16.68	12.68	11.7	16.29	13.7	11.36	7.75	1.22	16.48	11.22	6.3
Cl	mg/L	6.99	6.99	2.99	25.99	2.99	1.99	3.99	8.99	27.99	3.99	4.99	23.99	19.99	21.99	7.99
Salinity	%	0.02	0.01	0.03	0.03	0.02	0.01	0.01	0.04	0.03	0.09	0.03	0.05	0.07	0.06	0.03
TDS	ppm	183	321	181	316	248	173	140	486	250	245	140	249	247	249	152
PO ₄	mg/L	0.62	0.20	0.62	0.32	0.39	0.67	0.71	0.74	0.43	0.38	0.61	0.49	0.67	0.48	0.78
SO ₄	mg/L	0.08	0.59	0.20	1.38	1.10	0.26	0.48	4.80	1.16	0.99	0.12	1.72	3.20	1.63	0.34
NO ₃ -N	mg/L	52.7	24.9	37.6	34.28	10.94	29.06	33.04	40.04	26.66	23.65	30.16	18.56	17.04	28.17	29.9
NO ₂ -N	mg/L	0.01	0.01	0.01	0.01	0.02	0.01	0.01	0.02	0.003	0.01	0.002	0.02	0.01	0.02	0.01
Summer Season																
W.T.	°C	16	15.5	16	15	15	14.5	16	15	15.5	15	14.5	16	16.5	15	15.5
pH		7.91	7.71	7.98	7.86	7.81	7.72	7.91	7.86	7.80	7.91	7.70	7.88	7.86	7.94	7.90
E.C.	µs/cm	395	644	370	641	552	458	788	641	728	705	378	640	737	643	412
D.O	mg/L	3.04	2.66	2.47	1.90	2.09	3.72	4.37	1.52	5.71	4.76	3.61	3.80	4.56	4.76	5.14
T.H	°F	20	35.4	18.8	32	29.8	21.7	40	29	26.8	29	17.9	30.4	34	30	19
Ca	mg/L	48.09	68.13	40.08	80.16	123.9	80.15	60.92	91.3	20.04	113.8	48.7	74.54	97.7	61.7	48.8
Mg	mg/L	6.83	7.95	5.17	11.71	16.4	13.60	5.08	15.1	1.64	29	9.53	30.4	34	30	19
Cl	mg/L	2.99	0.99	0.99	1.99	8.99	4.99	4.99	20.99	36.9	53.9	2.99	26.9	31.9	26.9	9.99
Salinity	%	0.008	0.04	0.02	0.05	0.03	0.009	0.01	0.05	0.02	0.04	0.03	0.02	0.02	0.04	0.02
TDS	ppm	158	250	161	283	222	334	293	429	330	319	158	239	317	288	179
PO ₄	mg/L	0.65	0.18	0.64	0.37	0.41	0.69	0.70	0.73	0.55	0.50	0.87	0.50	0.68	0.53	0.79
SO ₄	mg/L	5.13	1.46	2.40	1.73	1.01	0.13	0.35	3.71	0.97	0.73	0.36	1.51	2.78	1.59	0.74
NO ₃ -N	mg/L	32.79	21.79	33.04	35.9	13.24	26.52	30.8	13.7	14.1	6.08	5.82	12.4	54.5	26.4	39.8
NO ₂ -N	mg/L	0.04	0.05	0.009	0.65	0.04	0.03	0.04	0.32	0.04	0.03	0.02	0.04	0.04	0.05	0.04

W.T: Water Temperature. E.C: Electrical Conductivity. D.O: Dissolved Oxygen. T.H: Total Hardness. TDS: Total Dissolved Solids

Table 3 (Continue). Physicochemical data measured at sampling stations according to seasons.

Parameters	Unit	1 st	2 nd	3 rd	4 th	5 th	6 th	7 th	8 th	9 th	10 th	11 th	12 th	13 th	14 th	15 th
Autumn Season																
W.T.	°C	8	8.5	8	9	9.5	9.5	8.5	10	10.5	10	10	11	11	10.5	11
pH		7.85	7.70	7.82	7.75	7.74	7.52	7.61	7.79	7.76	7.80	8.67	7.62	7.66	7.80	7.70
E.C.	µs/cm	404	612	365	567	509	416	359	886	737	581	395	639	656	630	442
D.O	mg/L	2.66	2.47	2.28	1.71	2.09	3.23	3.04	3.80	2.47	2.85	3.42	1.52	1.90	2.28	2.66
T.H	°F	26.2	38	22.4	37	30	26.8	24.4	35	31	24.4	20.6	30.6	32.6	32	24.4
Ca	mg/L	57.7	41.68	44.08	72.14	89.77	86.57	72.14	97.79	53.70	62.52	5.20	80.16	84.96	65.73	57.71
Mg	mg/L	7.66	0.89	5.27	8.54	14.53	14.53	12.68	15.27	5.52	9.90	8.70	12.05	12.84	8.54	6.25
Cl	mg/L	7.99	8.99	7.99	2.99	8.99	7.99	10.99	9.99	28.99	44.98	3.99	12.99	13.99	14.99	2.99
Salinity	%	0.01	0.008	0.008	0.01	0.01	0.008	0.02	0.01	0.05	0.02	0.01	0.01	0.02	0.05	0.03
TDS	ppm	202	306	182	283	254	208	179	443	368	290	305	319	348	178	300
PO ₄	mg/L	0.0005	0.01	0.006	0.004	0.01	0.001	0.003	0.012	0.005	0.003	0.02	0.012	0.008	0.009	0.01
SO ₄	mg/L	0.12	0.06	0.09	0.93	1.07	0.01	0.17	5.34	1.44	0.65	0.19	1.66	3.33	1.83	1.27
NO ₃ -N	mg/L	17.04	10.96	17.93	13.94	15.82	27.79	0.90	7.75	12.39	11.95	15.72	11.51	30.20	10.84	24.34
NO ₂ -N	mg/L	0.04	0.04	0.04	0.04	0.04	0.05	0.04	0.04	0.04	0.04	0.08	0.05	0.05	0.05	0.05
Average values of the stations																
W.T.	°C	10.3	10.5	10	10.1	10.5	10.3	10.1	10.6	10.8	10.3	10.5	11	11.1	10.6	11
pH		7.87	7.89	7.89	7.78	7.73	7.64	7.74	7.77	7.73	7.83	8.06	7.68	7.58	7.71	7.79
E.C.	µs/cm	399	636	367	618	527	423	503	840	640	662	385	559	796	643	421
D.O	mg/L	3.48	3.10	2.85	2.34	3.61	3.75	4.37	2.72	4.31	5.50	4.18	3.67	3.93	4.37	4.82
T.H	°F	22.06	36.8	20.06	33.93	29.13	23.36	28.13	33.2	28.53	26.8	19.03	30.73	33.93	30.13	20.46
Ca	mg/L	51.26	56.37	40.85	78.55	103.2	80.15	67.06	97.23	52.63	86.56	34.81	76.95	95.22	67.32	51.53
Mg	mg/L	7.10	4.75	5.05	10.84	15.87	13.60	9.82	15.53	6.95	16.7	8.66	14.55	21.10	16.58	10.51
Cl	mg/L	5.99	5.65	3.99	10.32	6.99	4.99	6.65	13.32	31.29	34.29	3.99	21.29	21.96	21.96	6.99
Salinity	%	0.01	0.01	0.01	0.03	0.02	0.009	0.01	0.03	0.03	0.05	0.02	0.02	0.03	0.05	0.02
TDS	ppm	181	292	174	294	241	238	204	452	316	284	201	269	304	238	210
PO ₄	mg/L	0.42	0.13	0.42	0.23	0.27	0.45	0.35	0.49	0.32	0.29	0.50	0.33	0.45	0.34	0.52
SO ₄	mg/L	1.77	0.70	0.89	1.34	1.06	0.13	0.33	4.61	1.19	0.79	0.22	1.63	3.10	1.68	0.78
NO ₃ -N	mg/L	34.17	19.21	29.52	28.04	13.33	27.79	21.58	20.49	17.71	13.89	17.23	14.15	33.91	21.80	31.34
NO ₂ -N	mg/L	0.03	0.03	0.01	0.23	0.03	0.03	0.03	0.12	0.02	0.02	0.03	0.03	0.03	0.04	0.03

W.T: Water Temperature. E.C: Electrical Conductivity. D.O: Dissolved Oxygen. T.H: Total Hardness. TDS: Total Dissolved Solids

Table 4. Water quality parameters of water troughs according to seasonal average values

	Spring	Summer	Autumn	Average
W.T. (°C)	6.57	15	9.7	10.42
SKKY, 2004	I	I	I	I
pH	7.67	8	7.8	7.82
TSWQR,2016	I	I	I	I
E.C.(µs/cm)	554.93	582	546.5	561.14
TSWQR,2016	I-II	I-II	I-II	I-II
D.O (mg/L)	5.04	4	2.6	3.88
TSWQR 2016	II-III	II-III	IV	II-III
T.H (°F)	26.65	28	29	27.88
Ca (mg/L)	72.64	71	64.8	69.48
Mg (mg/L)	10.31	16	9.5	11.94
Cl (mg/L)	11.46	16	12.6	13.35
SKKY, 2004	I	I	I	I
Salinity (‰)	0.04	0.01	0.01	0.02
TDS (ppm)	238.67	264	277.7	260.12
PO ₄ (mg/L)	0.54	1	0.01	0.52
SO ₄ (mg/L)	1.2	2	1.2	1.47
SKKY, 2004	I	I	I	I
NO ₃ -N (mg/L)	29.11	24	15.3	22.80
TSWQR,2016	IV	IV	II-III	IV
NO ₂ -N (mg/L)	0.01	0.01	0.1	0.04
SKKY, 2004	II	II	IV	IV

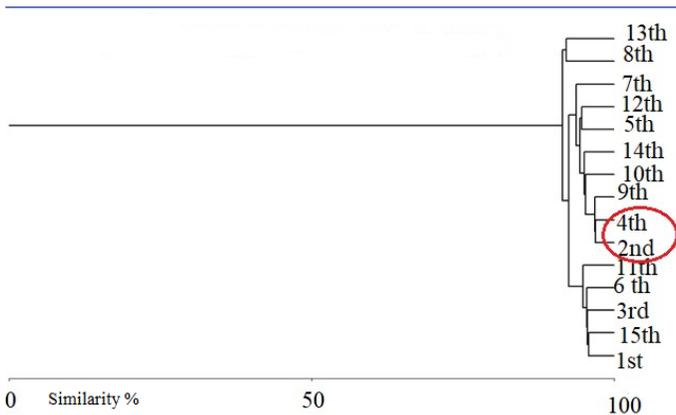


Figure 3. Bray Curtis Similarity dendrogram in terms of physicochemical values.

tween W.T. and NO₂-N; D.O. and PO₄; E.C. and Cl; E.C. and SO₄; E.C. and TDS; Cl and salinity; Cl and TDS. As temperature increases in aquatic systems, oxygen concentration decreases. This causes the release of phosphorus from the sediment (IPCC, 2008). The results support the literature. Significant positive correlation was found between E.C. and Cl, and between SO₄ and TDS. The conductivity of water increases in the presence of inorganic dissolved solids such as Cl, NO₃, SO₄ and PO₄ anions or Na, Mg, Ca, Fe and Al cations (Spellman, 2003). Salinity consists of four cation groups such as Ca, Mg, Na, K and four anion groups such as HCO₃, CO₃, SO₄ and Cl (Egemen & Sunlu 1999). Cl, SO₄ and TDS were all found to be significantly positively correlated with E.C. The results are consistent with the literature. Correlation coefficients are given in Table 6.

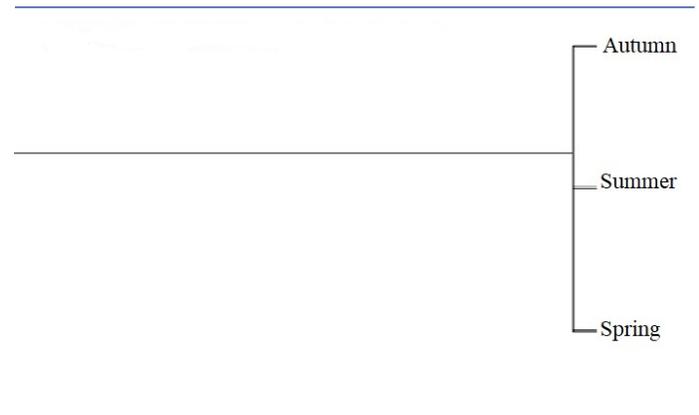


Figure 4. Bray-Curtis similarity dendrogram in terms of seasons.

When the study is compared with previous studies, it can be said that some similar features are observed. Özkan (2006a) taxonomically examined samples from six different localities in Bozcaada between 1999-2002. The same study also identified a total of 14 species and made diagnostic keys of the species. All species are reported as new records for Bozcaada. *Chironomus salinarius* K was reported as a new species for the Chironomidae fauna of Turkey. *C. thummi*, *C. anthracinus*, *C. viridicollis*, *Einfeldia pagana* and *Polypedilum aberrans* were recorded as species emerging from the water troughs. Özkan (2006a) reported that *C. ripar-*

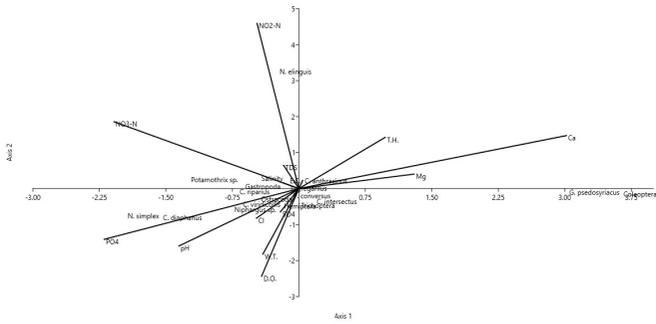


Figure 5. CCA Analysis of physicochemical values with taxa.

Table 5. Pearson Correlation Analysis.

	Ca	TH	pH	NO ₃ -N
Ca	1			
TH	.468**	1		
pH	-.500**	-.294	1	
NO ₃ -N	-.084	-.173	.092	1

** : correlation is significant at 0.01 level (p < 0.01);
 -: no statistically significant correlation was detected
 ** : Correlation is significant at the 0.01 level (2-tailed).

Külköylüoğlu et al. (2013) collected samples from water troughs in Bolu, Erzincan, Gaziantep, Kahramanmaraş, Ordu and Van provinces at different times between 2006 and 2010. They identified 32 Ostracoda species in total from 105 water troughs and investigated the composition and diversity of these species. The species belonging to Ostracoda were reported as *Heterocypris incongruens*, *Ilyocypris bradyi*, *Psychrodromus olivaceus*, *Candona neglecta*. The authors of the above-mentioned study identified water troughs as important structures to be studied due to their high-altitude locations and the unique species diversity to be found in them. They stated that zooplankton, phytoplankton and molluscs of different taxonomic groups found in water troughs should also be studied. In this study, ostracoda were found at stations 7 and 10 in the summer season and at stations 3, 7, 12 and 13 in the autumn season. However, the exact species could not be identified.

Başak et al. (2013) collected samples belonging to Copepoda and Cladocera groups from zooplankton in Ankara province. They reported 12 new zooplankton species for Ankara. The most common species found was *Eucyclops serrulatus*.

Külköylüoğlu et al. (2017) investigated the diversity and distribution of Ostracods between Karabük and Düzce provinces. They reported that the most widespread species was *Heterocypris incongruens*.

Table 6. Spearman Correlation Analysis.

	NO ₂ -N	Mg	D.O.	E.C.	W.T.	Cl	Salinity	SO ₄	TDS	PO ₄
NO ₂ -N	1									
Mg	.137	1								
D.O.	-.600**	.069	1							
E.C.	.206	.299*	.010	1						
W.T.	.594**	.175	-.424**	.209	1					
Cl	.110	.330*	.111	.471**	.054	1				
Salinity	.272	.313*	-.055	.292	.356*	.443**	1			
SO ₄	.209	.323*	-.170	.527**	.287	.253	.380*	1		
TDS	.371*	.170	-.259	.731**	.231	.421**	.202	.342*	1	
PO ₄	-.425**	.155	.488**	-.027	.052	-.116	.031	.129	-.197	1

*: correlation significant at 0.05 level (p < 0.05);
 **: correlation is significant at 0.01 level (p < 0.01);
 -: no statistically significant correlation was detected

ius and *C. anthracinus* lived mostly in the muddy bottom of stagnant waters, while *C. viridicollis* lived in the mud of stagnant waters. In this study, *C. riparius*, *C. anthracinus* and *C. viridicollis* species were found. Our study supports the literature in terms of both habitats and species.

Özkan (2006b) reported that *Paracladius conversus* and *Tanytarsus gregarilus* species were observed in stagnant water bodies in Gökçeada. *Paracladius conversus* and *T. gregarilus* were also identified in this study, in line with the larger literature.

In their study, Külköylüoğlu et al. (2012) investigated whether water troughs constructed as artificial sources are beneficial for biodiversity and for the ecological characteristics of Ostracods in water troughs in some regions of Turkey.

Özer & Dikmen (2021) analyzed water troughs across the world, but especially in Erzurum, Turkey. They classified the water troughs according to type of material, number of sections, aesthetic features, and shapes, and made some suggestions on the protection and development of these structures.

Ataman et al. (2023) investigated the diversity and seasonal cycles of freshwater Ostracoda in man-made water troughs. They reported that *Heterocypris incongruens*, *Heterocypris salina* and *Ilyocypris bradyi* species were found in the water troughs.

Sanbur (2023) investigated the diatom species diversity of rural fountains and water troughs in various regions of Eskişehir. The

conclusion of the study showed that 200 diatom taxa from the water troughs had been identified. The researcher reported that diatom diversity was high in the water troughs.

As a result of this multivariate statistical study carried out on water samples collected from 15 water trough stations across the Şarkışla region of Sivas, statistically significant relationships were found showing that altitude and agricultural activity impact water quality. These factors contribute to variations in water quality from the 1st to the 4th water quality grades. Within agricultural activities, the water quality of troughs may deteriorate due to pesticide applications. Pesticides can cause deterioration of groundwater quality and deterioration of the physical, chemical, and biological structure of water through irrigation and precipitation due to rain and snow. In areas where livestock breeding is intensive, other factors polluting groundwater resources should be identified and water troughs should be evaluated based on integrated basin management. In order to identify this, it is very important to carry out studies of this kind at regular intervals to monitor the water quality and benthic macroinvertebrate fauna of surface and groundwater resources and to catalogue and maintain the hydrobiological diversity of Turkey.

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Bio-indicator Bacteria Levels in Riva Stream, an Important Stream in İstanbul, Türkiye

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ABSTRACT

Bacterial pollution is the presence of harmful bacteria in aquatic ecosystems, which can cause health problems if ingested. These bacteria originate from a variety of sources, including human and animal waste, agricultural runoff, and untreated sewage. This study aimed to assess the levels of bioindicator bacteria and their associations with various environmental parameters in surface water samples. The samples were collected seasonally from five stations along Riva Stream between March 2018 and January 2019. Indicator bacterial levels were determined using membrane filtration. Temperature, salinity, pH, dissolved oxygen, and conductivity values of the stations were measured in situ during sampling using a multiparameter (YSI). As a result of the sampling, maximum total coliform, fecal coliform, and intestinal enterococci levels were recorded as 24×10^3 CFU/100ml, 18×10^3 CFU/100ml and 24×10^2 CFU/100 ml, respectively. During sampling, the indicator bacterial levels were above the limit at all stations. When the FC/IE ratios were analyzed, the maximum FC/IE ratio was recorded as 52.10 during the winter sampling period at the station. The detection of bioindicator bacteria levels above the standard limit throughout the sampling period (especially during the summer period) was associated with a high level of domestic waste in the region. The results showed that more frequent and detailed monitoring studies should be conducted in this region.

Keywords: Total coliform, Fecal coliform, Intestinal enterococci, Environmental parameters, Riva stream

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INTRODUCTION

Microbial communities have a significant impact on the regulation of water quality and biogeochemical cycling in freshwater ecosystems. It is well-known that bacteria play a pivotal role in the decomposition and stabilization of organic matter in both natural environments and biological sewage treatment processes. Comprehending the ecology and evolution of microbial communities and the implications of these changes for public health, it is imperative to evaluate physical and chemical parameters, including salinity, temperature, and pH (Ibekwe et al., 2012).

Accelerating urbanization, increased population density, and ineffective sewerage infrastructure systems have important implications for public health (Mallin, 2000; 2001; Garbossa, 2017). Water sources contaminated with sewage and human fecal contamination are related to the occurrence of infectious waterborne diseases (McKee and Cruz, 2021; Manetu and Karanja, 2021; Kahn et al., 2018).

Coliform bacteria, especially enteric bacteria, serve as indicators of the presence of bacterial pathogens. Although fecal coliforms themselves are generally not directly harmful to human health, their detection indicates the pres-



ence of fecal waste that may harbor harmful pathogens (Farrell, 2013). The microbial water quality standard includes the evaluation of bacterial indicator organisms, which often includes the coliform group, fecal streptococci, enterococci, and *Escherichia coli* (*E. coli*) (Wen et al., 2020; Davene et al., 2020; Augustyn et al., 2016; APHA, 2013). When pollution originates from human waste, the fecal coliform/fecal streptococcal ratio (FC/IS ratio) is greater than 4, between 0.1 and 4.0 when it originates from domestic animals such as cattle, and below 0.1 in wild animals (Clinton et al., 2006, Benhassane et al., 2019, Gerba, 2009).

The Marmara Sea, an inland sea between Asia and Europe and the gateway to the Aegean and Black Seas through the Bosphorus (Bosphorus) and the Dardanelles (Dardanelles), is important for the fisheries sector, maritime transport, and ecosystem health (Altuğ et al., 2009). The Riva Stream flows through Riva village, İstanbul and it is approximately 33 km long. It is located between the Black Sea and the exit of the Marmara Sea and is one of the largest rivers in the region. Ömerli Dam, the largest drinking water reservoir on Riva Stream, was established by the General Directorate of State Hydraulic Works (DSİ) in 1972 (Tarkan, 2010; Özuluğ et al., 2005). The area around Riva Stream and Riva Beach offers opportunities for a variety of outdoor activities such as hiking, picnics, camping and water sports. Due to these features, it has become a popular touristic and recreational destination.

Studies indicating the presence of pollutants and including topics such as fishing, mollusks, indicator bacteria and eutrophication have emerged in previous studies on the Riva Stream (Dökümcü and Koşal Şahin, 2022; Altuğ et al., 2016; Garipağaoğlu et al., 2015; Tarkan, 2010; Özuluğ et al., 2005; Dinseven and Çurğunlu, 1988).

This study monitored the level of indicator organisms associated with bacterial pollution at determined stations in Riva stream and determined the relationship of bio-indicators with variable environmental parameters.

MATERIALS AND METHODS

Surface water (0-30 cm) was collected seasonally from five stations between March 2018 and January 2019. The samples were delivered to a laboratory under cold chain conditions and were analyzed on the same day as the sampling. The stations are shown in Figure 1 and listed in Table 1.

The membrane filtration method was used to determine indicator bacteria levels. The 100ml samples filtered through sterile filters were placed in Endo-NKS (sartorius) medium for total coliform bacteria, m-FC - NKS (Sartorius) for fecal coliform and azide - NKS (Sartorius) medium for intestinal enterococci. The media were incubated for 24 hours at 37±0.1°C for total coliforms and intestinal enterococci, and at 44±0.1°C for fecal coliforms. The colonies grown at the end of the incubation were counted and recorded as CFU/100ml. The pink-red colonies exhibiting a yellow-green metallic sheen in the m-Endo medium were considered as potential indicators of coliforms. The Cytochrome oxidase test (API Strep, BioMereux) was performed on the pink-red colonies with a yellow-green metallic sheen, which were evaluated as suspicious for coliform, and oxidase-negative colonies were taken into numerical evaluation. The

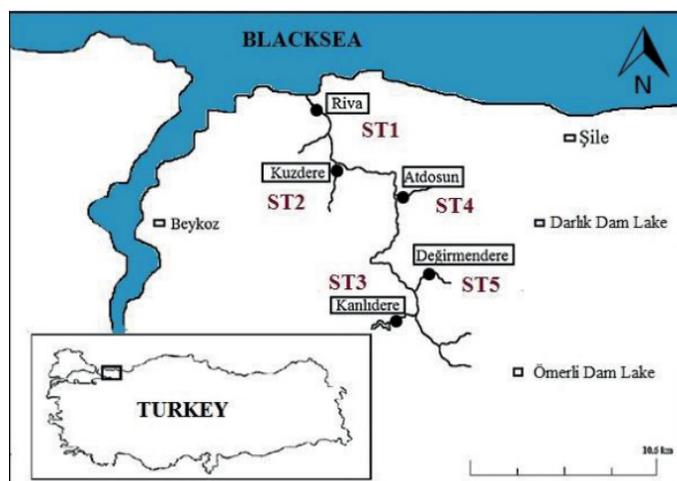


Figure 1. The positioning of Riva Stream and the designated sampling sites.

Table 1. The geographical coordinates of the sampling stations along Riva Stream.

Station No	Station Name	Coordinate
S 1	Riva Stream	N 41.12763° E 29.13521°
S 2	Kuzdere	N 41.12763° E 29.13521°
S 3	Kanlıdere	N 41.07127° E 29.16175°
S 4	Atdosun Stream	N 41.10403° E 29.16333°
S 5	Değirmendere	N 41.08222° D 29.17393°

blue colonies observed in the m-FC medium were considered as potential indicators of fecal coliforms. Cytochrome oxidase (API Strep, bioMérieux) and indole tests were performed on these colonies with suspected fecal coliform. Oxidase negative and indole positive colonies were evaluated numerically (APHA, 2013). To determine if the bacteriological contamination was of human or animal origin (such as cattle), the ratio of fecal coliform bacteria levels (FC) to intestinal enterococci (IE) bacteria levels was evaluated. Saturated oxygen (mg/L), water temperature (°C), salinity (‰), conductivity (µS/cm) and pH values were measured in situ during sampling using multiparameters (YSI). Spearman's rank correlation statistical analysis was performed to determine the relationship between environmental parameters and indicator bacterial levels. The significance level for differences between the parameters was set at 0.05. The differences between the measured environmental parameters and bacterial levels were examined by two-way ANOVA test. Statistical analyses were performed using SPSS 28.0.

RESULTS AND DISCUSSION

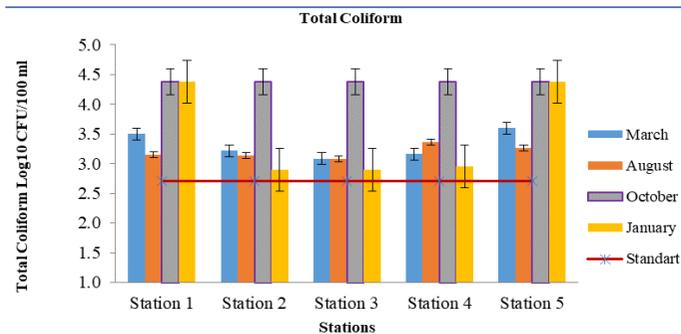
The variable environmental parameters of the sampling areas, including dissolved oxygen, temperature, salinity, pH, and conductivity, are shown in Table 2.

Table 2. Physicochemical parameters on Riva Stream.

Stations	Months	Temp (°C)	Salinity (‰)	pH	O ₂ (mg/L)	Conductivity (µS/cm)
Station 1	Mar-18	11.47	1.25	8.67	4.82	2.44
	Aug-18	24.50	2.40	7.70	3.46	1.12
	Oct-18	18.10	1.40	7.14	3.47	3.07
	Jan-19	5.40	0.20	7.91	11.50	0.97
Station 2	Mar-18	10.47	0.34	8.82	9.70	0.41
	Aug-18	21.50	0.00	7.69	5.82	0.60
	Oct-18	17.70	0.00	7.07	4.70	0.44
	Jan-19	5.80	0.00	8.04	13.76	0.32
Station 3	Mar-18	11.46	0.17	8.88	10.08	0.31
	Aug-18	16.6	0.00	6.32	6.78	0.04
	Oct-18	15.7	0.00	7.16	6.80	0.02
	Jan-19	6.5	0.00	9.63	12.39	0.15
Station 4	Mar-18	11.0	0.16	8.59	10.34	0.58
	Aug-18	21.6	1.20	7.57	8.83	0.01
	Oct-18	15.9	0.00	7.20	6.02	0.04
	Jan-19	4.4	0.00	8.96	12.65	0.25
Station 5	Mar-18	11.97	0.27	7.92	7.58	0.55
	Aug-18	24.9	0.40	7.51	5.52	0.39
	Oct-18	16.2	0.30	7.40	6.76	1.10
	Jan-19	5.4	0.00	8.10	12.57	0.48

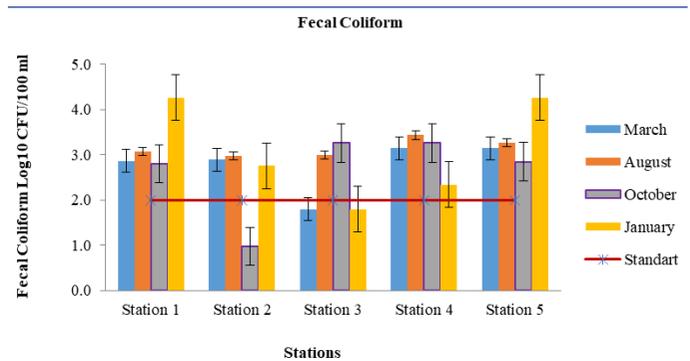
pH values 6.32 to 9.63, dissolved oxygen values 3.46 to 13.76mg/L, temperature values 4.4-24.9°C, salinity values and ppt values 0 to 2.4 recorded as the maximum and minimum, respectively.

The detected total coliform levels in the samples are shown in Figure 2.

**Figure 2.** Total coliform bacteria levels are determined in water samples by seasons and stations.

The levels of coliform bacteria in samples from all stations were found to exceed national limit values. The highest total number of coliform bacteria was detected as 24×10^3 CFU/100ml in October and January 2018.

The detected fecal coliform levels in the samples are shown in Figure 3. The fecal coliform levels were higher than the recommended national limit values. Fecal coliform counts were recorded at a minimum of 10 CFU/100ml and a maximum of 18×10^3 CFU/100ml during the study period.

**Figure 3.** Fecal coliform levels are determined in water samples by seasons and stations.

The detected intestinal enterococci levels in the samples are shown in Figure 4. The levels of intestinal enterococci were higher than the recommended national limit. Intestinal enterococci counts were recorded at a minimum of 38 CFU/100ml and a maximum of 24×10^2 CFU/100ml during the study period.

The fecal coliform/ intestinal enterococci ratios (FC/IE ratio) analyzed in the surface water samples during the study period are shown in Table 3. Data presented in Table 3 shows that the FC/IE counts recorded a minimum 0.04 and maximum 52.10 during the study period.

Values are shown in bold. If the FC/IE ratio is greater than 4, the pollution is considered as human origin, and if the FC/IE ratio is between 0.1 and 4, the pollution is considered as animal origin.

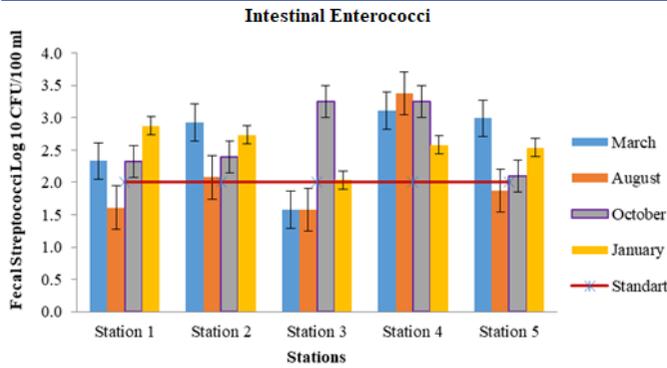


Figure 4. Intestinal enterococci bacteria levels are determined in water samples by seasons and stations.

Table 3. Fecal coliform/Intestinal Enterococci ratio at all stations.

Stations	FC/IE			
	Mar-18	Aug-18	Oct-18	Jan-19
S 1	3.38	28.34	2.95	23.89
S 2	0.91	7.78	0.04	1.05
S 3	1.64	25.84	1.00	0.57
S 4	1.08	1.12	1.00	0.58
S 5	1.42	24.21	5.60	52.10

Many river and stream studies conducted worldwide pointed out that the pollution rate is above the WHO and EPA standards. Shittu et al. (2008) reported that the total coliform count of Abeokuta, Southwest Nigeria, exceeded the EPA's maximum contamination standards for coliform bacteria in drinking water. Viau et al. (2011) reported that Enterococci were associated with bacterial pathogens from the five indicators tested in Hawaiian Oahu coastal streams. Musyoki et al. (2013) reported that the microbiological contamination of the Nairobi and Athi rivers was unacceptably high, according to Kenya standards and WHO guidelines for drinking water and agricultural use. Akkan et al. (2019) reported that the Total Coliform (TC), Fecal Coliform (FC) and Fecal Streptococcus (FS) rates of surface water samples in Yağlıdere were 45%, 71.66% and 56.66%, respectively. Altug et al. (2016) reported that the total coliform and fecal coliform counts were higher than advised national limit values in Riva stream.

The data obtained within the scope of the study was analyzed in accordance with national and international water quality standards. The standard values for the European Parliament are a total coliform count 500 CFU/100ml (2.69/100 ml based on log₁₀), fecal coliform count 100 CFU/100ml (2.0/100ml based on log₁₀), and intestinal enterococci counts 100 CFU/100ml (2.0/100ml based on log₁₀ value), which are indicated in dark red in the graphs below (European Parliament, 2006). Standard values for sea water used for recreational purposes were determined by the National Regulation published on 31 December 2004. According to this, limit values were established as 1000 cfu/100ml

for total coliform, 200 cfu/100ml for fecal coliforms and 100 cfu/100ml for fecal enterococcus (Official Gazette 2004).

When total coliform, fecal coliform and intestinal enterococci data were examined, it was determined that the bacteriological pollution level in the Riva River was above the standard limits in all seasons. During the sampling period, the highest levels of total coliforms were recorded in summer, which was associated with increased human activities. Fecal coliform counts peaked at a maximum of 18×10^3 CFU/100ml throughout the study, while intestinal enterococci counts ranged from a minimum of 38 CFU/100ml to a maximum of 24×10^2 CFU/100ml. The fact that the levels of total coliforms, fecal coliforms and fecal enterococci consistently exceeded the recommended national limits draws attention to the persistent presence of bacteriological pollution sources in the region.

Environmental variables significantly impact the levels of indicator bacteria in rivers (Herrig et al., 2019; Islam et al., 2017). Indicator bacteria such as *E. coli* and fecal coliforms which are used to measure the presence of fecal contamination in water can indicate the potential presence of other harmful bacteria and viruses (Saxena et al., 2015; Lin and Ganesh, 2013). Temperature is an important factor in bacterial growth in aquatic ecosystems. Warm temperatures can accelerate bacterial growth, leading to higher levels of indicator bacteria in the water. Conversely, cold temperatures can slow bacterial growth and reduce the levels of the indicator bacteria (Kunili and Ateş, 2021). During the summer months, when temperatures are warmer and recreational use of rivers is more common, there is typically an increase in the levels of indicator bacteria owing to higher levels of human and animal waste entering the water. In our study, no significant relationship was found between the total, fecal coliform levels, and water temperature. This situation is due to the fact that there is a continuous input of pollution to the study area. Heavy rainfall washes pollutants, including bacteria, from the surrounding land into the river, leading to increased levels of the indicator bacteria. This is particularly true for urban areas, where rainwater runoff can carry pollutants from streets and parking lots into nearby rivers (Shehane et al., 2005). Higher flow rates dilute the bacteria in water, thereby reducing the levels of indicator bacteria. However, during periods of high flow, bacteria is more easily resuspended from the riverbed and carried downstream, potentially increasing the levels of the indicator bacteria (Crump et al., 2007). Agricultural and urban land use contributes to higher levels of indicator bacteria in rivers. Agricultural runoff contains animal waste and fertilizers, whereas urban runoff contains a variety of pollutants, including pet waste, oil and grease from roads, and chemicals from industrial areas (Gotkowska-Płachta, et al., 2016; Paule-Mercado et al., 2016).

When examining the relationship between measurements of various environmental parameters, a significant correlation was found between the pH values and temperature values ($r = 0.005$, $p < 0.05$). Furthermore, pH values exhibited variations at the different stations (< 0.001), but no seasonal variation was observed. A significant correlation was identified between the dissolved oxygen values, temperature, pH, and conductivity

Table 4. Spearman's rank correlations between environmental parameters and indicator bacteria levels in Riva stream (N 20; marked correlations are significant at $p < 0.05$).

		Correlations										
		Depth	Width	FV	Temp	pH	Cond	DO	Salinity	TC	FC	IE
D	PC	1										
	Sig. (2-tailed)											
W	N	20										
	PC	.110	1									
FV	Sig. (2-tailed)	.645										
	N	20	20									
T	PC	-.133	-.319	1								
	Sig. (2-tailed)	.576	.171									
pH	N	20	20	20								
	PC	-.354	.066	-.283	1							
Cond	Sig. (2-tailed)	.126	.782	.227								
	N	20	20	20	20							
DO	PC	.313	-.023	.500*	-.607**	1						
	Sig. (2-tailed)	.179	.924	.025	.005							
Salinity	N	20	20	20	20	20						
	PC	.188	.802**	-.290	.101	-.034	1					
TC	Sig. (2-tailed)	.427	<.001	.215	.672	.888						
	N	20	20	20	20	20	20					
FC	PC	.149	-.358	.372	-.826**	.600**	-.461*	1				
	Sig. (2-tailed)	.530	.122	.106	<.001	.005	.041					
IE	N	20	20	20	20	20	20	20				
	PC	.040	.715**	-.249	.489*	-.080	.591**	-.527*	1			
FC	Sig. (2-tailed)	.866	<.001	.289	.029	.737	.006	.017				
	N	20	20	20	20	20	20	20	20			
IE	PC	-.061	.169	-.349	-.029	-.462*	.219	-.204	-.137	1		
	Sig. (2-tailed)	.799	.477	.131	.904	.041	.354	.388	.566			
FC	N	20	20	20	20	20	20	20	20	20		
	PC	.078	.253	-.226	-.381	-.015	-.003	.373	-.131	.453*	1	
IE	Sig. (2-tailed)	.743	.281	.337	.098	.950	.991	.105	.582	.045		
	N	20	20	20	20	20	20	20	20	20	20	
FC	PC	-.238	-.250	-.102	.038	-.137	-.350	.104	-.055	.165	.063	1
	Sig. (2-tailed)	.312	.289	.668	.873	.566	.130	.662	.818	.487	.791	
IE	N	20	20	20	20	20	20	20	20	20	20	20

*. Correlation is significant at the 0.05 level (2-tailed); **. Correlation is significant at the 0.01 level (2-tailed).

F.V. Flow Velocity °C; Temperature Cond; Conductivity ($\mu\text{S}/\text{cm}$) D.O; Dissolved Oxygen %; Salinity/ TC: Total Coliform levels, FC: Fecal Coliform Levels, IE: Intestinal Enterococci levels

values. Additionally, a significant difference was noted in the dissolved oxygen values among different stations. In the analysis of water temperature values, a disparity was observed between stations ($p < 0.05$), while no significant difference was observed between seasons. The total coliform values displayed significant differences among stations. Overall, environmental variables have a significant impact on the levels of indicator bacteria in rivers. Monitoring these variables and understanding their effects can help water managers and policymakers develop strategies to reduce bacterial pollution in rivers and protect public health.

CONCLUSION

Riva Stream is important due to its ecosystem structure and it is also a popular place for tourism and recreation activities. This study, carried out between March 2018 and January 2019, investigated the seasonal levels of indicator bacteria in Riva stream. It was determined that indicator bacteria levels throughout the study area exceeded national and international standard values throughout the year in every season. Human activities were identified as the primary source of pollution. Seasonal sampling from the Riva stream provided important data for the evaluation of the pollution load carried by the stream and the possible introduc-

Table 5. Two-way ANOVA between bio-indicator bacteria and environmental parameters.

	Stations		Seasons	
	F.	Sig.	F.	Sig.
Depth	1.245	.326	1.645	.215
Width	.003	1.000	4628.520	<.001
Flow Velocity	2.192	.129	2.015	.144
Temperature	75.211	<.001	.071	.990
pH	10.356	<.001	.088	.985
Conductivity	.529	.669	8.256	<.001
Dissolved Oxygen	17.106	<.001	.712	.596
Salinity	1.207	.339	4.494	.014
Total Coliforms	13.309	<.001	.345	.844
Fecal Coliforms	2.056	.147	.804	.542
Intestinal Enterotococci	.288	.834	2.476	.089

tion of bacteria to the Black Sea coast. Bacterial pollution, which was found to flow from both sides of Riva Stream to the Black Sea poses a potential risk for recreational activities in the region. To improve water quality, it is imperative to implement a comprehensive management strategy and adopt measures to address water quality issues. Additionally, improving national awareness of the problems associated with fecal pollution in freshwater sources is crucial to maintaining overall ecosystem health.

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Seasonal Feeding Biology of Catfish, *Pachypterus atherinoides* (Bloch, 1794) with Special Reference to Lentic and Lotic Ecosystem

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ABSTRACT

The study of food and feeding biology is extremely crucial in fishery. *Pachypterus atherinoides* (Bloch, 1794) is a good food and ornamental catfish in the world. The present study aims to examine the seasonal variation of feeding habits of *P. atherinoides* in both lentic and lotic ecosystems. The study was conducted over two years in selected lentic and lotic aquatic ecosystems of the Jhargram and Paschim Medinipur districts of West Bengal, India. The study reveals that *P. atherinoides* shows carni-omnivorous nature. Pearson's correlation represents gastro-somatic index, relative gut length, and hepato-somatic index have significant positive correlation among them. Based on posthoc tests, it has been observed that the gastro-somatic index, relative gut length, and hepato-somatic index vary significantly both seasonally and in both lentic and lotic ecosystems. This study would help aquaculturists in the culture, captive breeding, and conservation of *P. atherinoides* and also help researchers in the biometric study of another fish.

Keywords: Carni-omnivorous, Catfish, Feeding biology, Lentic, Lotic, *Pachypterus atherinoides*

INTRODUCTION

The study of food and feeding behaviors is extremely crucial in fishery biology. Fish mostly obtain their energy from food, which has a big impact on their population size, growth rate, and overall health. Because they enable fish to consume all of the potential food in water bodies without competing with one another and living in communities with other fish, diet and feeding habits are crucial in aquaculture (Victor *et al.*, 2014). To determine the fullness of the stomach and the feeding status of the species, the gastro-somatic index (GSI) is used and the relative gut length (RGL) is applied to determine the eating habits and to measure a certain relationship with the length and life stage of the fish species. The RGL value is often used to predict a wide range of food groups and show which foods different fish species like to eat. The study of fish-eating habits and gut content analysis has several objectives, including learning about their

food preferences, determining whether they obtain enough food seasonally, and determining when feeding intensity peaks and troughs. The study of a species' feeding ecology can indicate how the creature has evolved ecologically to manage the challenge. It is also vital for the species' propagation to get a better understanding of its feeding behavior, as this information can be used to exploit natural fish food. The HSI measures the liver weight to body weight ratio. An indicator of eating and metabolism status is the HSI biomarker. Changes in HSI and GSI values are good indicators of fish with compromised reproductive function. HSI and GSI can go up during Vitelogenesis. *Pachypterus atherinoides* (Bloch, 1794), often known as Indian potasi, is a common catfish belonging to the family Schibeidae found in rivers and ponds throughout India (except Kerala), Pakistan, Bangladesh, Nepal, Myanmar, Bunna (Menon, 1999). It has high nutritional and also ornamental value. The fish meal of the species contains energy 343

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Kcal, moisture 16 gm, protein 58gm, fat 09 gm, mineral 17 gm, calcium 1597 mg, phosphorus 595 mg, Iron 41 mg each 100 gm (Gopalan et al., 2004). Aquarists are attracted to it because of its bright color and small size. This species has been gradually decreasing due to pollution, habitat destruction, selective captive breeding, removal from aquatic bodies for their carnivorous nature, and also the high preferences of exotic catfish [African catfish *Clarius gariepinus* (Burchell, 1822) and Thai catfish *Pangasius sutchi* due Fowler, 1937]. Globally, there has been very little research on the feeding biology of *Pachypterus atherinoides* (Bloch, 1794). In India, except for Gogai et al., 2020, there is no study on the feeding biology of this species. However, they did not observe seasonal feeding behavior and habitat/ecosystem differentiation, and they did not study both lentic and lotic ecosystems. In West Bengal, some aspects of various indigenous fish species have been studied by different researchers (Dasgupta, 2004; Chattopadhyay et al., 2014; Gupta and Banerjee, 2014; Jana et al., 2021; Jana et al., 2022A; Jana et al. 2022B; Sit et al. 2020; Chanda and Jana., 2021; Sahil et al., 2022; Sit et al. 2022A; Sit et al. 2022B; Sit et al., 2023A; Sit et al., 2023B) but not to observe the feeding habit of the said species. The interspecific interaction and productivity of water bodies can be determined by studying fish feeding habits. Fish nutrition and feeding have an impact on their development, reproduction, and health (Srivastava et al., 2014). Most research on the food and feeding habits of fish from diverse habitats has shown that those species varied in time and place, as well as at different stages of growth, emphasizing the need for more research into a species' food and feeding habits.

MATERIALS AND METHODS

Collection of fish specimens: Specimens were collected every fifteen days from lentic (selected Ponds) and lotic (selected River sites) ecosystems/habitat of Paschim Medinipur and Jhargram districts, West Bengal, India, during the Pre-monsoon/Summer (March-June), monsoon (July-Oct), and Post-monsoon/Winter (Nov-Feb) seasons since March 2020 to February 2022 (Figure 1).

Measured length and weight: Seasonally and ecosystem/habitat-wise, the total weight and total length of each specimen were measured by a digital weighing machine with 0.01 gm accuracy and a digital slide caliper instrument with 0.01 mm accuracy, respectively.

Dissection and Internal organ measured: All internal organs were collected by dissection with the help of scissors, forceps, needles, and a brush. The liver and digestive tract were dissected out and kept in a vial with a labeled 10% formalin solution. Digital slide calipers were used to measure the length of the digestive tract. After that, the stomach was separated from the digestive system. An electronic balance was used to weigh the liver and stomach.

Observed stomach fullness: Stomach distention per species, as determined by season, has been observed. According to Nagar and Sharma (2016), the stomach's distension was graded as 'full', '3/4 full', '1/2 full', '1/4 full', and 'empty' by eye assessment.

Gastrosomatic index (GaSI): The following formula (Desai, 1970) was used to figure out the Gastrosomatic Index (GaSI), or the amount of food eaten each month:

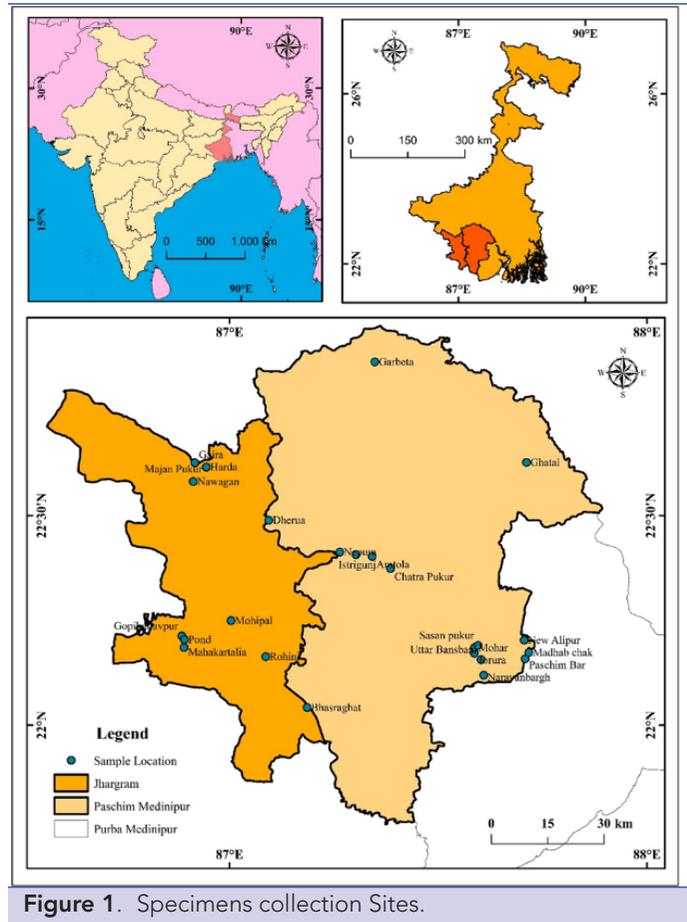


Figure 1. Specimens collection Sites.

$$\text{GaSI} = \frac{\text{Weight of the stomach}}{\text{Weight of fish}} \times 100$$

Relative gut length (RGL): The following formula was used to compute the Relative Gut Length (RGL) (Al-Hussain, 1949).

$$\text{RLG} = \frac{\text{Length of gut}}{\text{Total body length}}$$

Hepatosomatic index (HSI): HSI was determined by Rajaguru's 1992 formula.

$$\text{HSI} = \frac{\text{Weight of liver}}{\text{Weight of fish}} \times 100$$

Food content analysis: To identify the various foods that the fish swallowed, the stomachs were dissected, and 1 mL of the food contents were taken in a glass vial and examined under a light microscope (XSP L101). Hynes's (1950) frequency of occurrence method was used to look at the foods, and the following equation was used:

$$\text{Percentage of occurrence of a food type} = \frac{\text{Number of guts where the food occurred}}{\text{Total number of guts analyzed}} \times 100$$

Data analysis

Finally, data were analyzed (Descriptive statistics, MANOVA, Pearson's Correlation, Post Hoc test,) with the help of Microsoft Excel (2019), SPSS (2021), and Origin Pro (2023) software systems.

RESULTS AND DISCUSSION

Seasonally, habitat-wise maximum, minimum, and average values of gastrointestinal length and weight of *P. atherinoides* in both districts were represented in Table 1. GaSI values were highest during the Summer season (6.65 ± 1.11 , 7.06 ± 1.81 , 7.95 ± 1.90 , and 7.08 ± 1.56), followed by the Winter (6.09 ± 1.05 , 4.85 ± 1.37 , 6.40 ± 1.01 , and 6.09 ± 1.41) and the monsoon (2.93 ± 1.00 , 4.70 ± 1.53 , 6.16 ± 0.96 , and 5.41 ± 1.42) in both habitats (pond and river) of two districts (Table 2 and Figure 2-3). Gogoi et al. (2020) recorded the average GaSI value of *P. atherinoides* was 2.683 ± 0.495 gm, with the highest GaSI value (2.832 ± 0.754) during the winter-spring season (Feb-April) and minimum (2.525 ± 0.363 gm) during the summer-rainy (May-July) in the Subansiri river, Assam. In the present study, the value of GaSI is great-

er than that of Gogoi et al. The GaSI values change seasonally, and the highest during pre-breeding and lowest during breeding have been similar to the studies of Gupta and Banerjee (2014), Kurbah and Bhuyan (2018), Sharma et al. (2018) and Gogoi et al. (2020) but not similar to the study of Chaturvedi and Sakseena (2013). GaSI values indicated maximum food intake during the pre-spawning season and minimum during the breeding season. GaSI ranges from 2.91 to 6.08, representing the voracious nature of fish (Lanthaimeilu and Bhattacharjee, 2018), so the present result indicates the species has been the same (Table 2). In the present study, the RGL values ranged from 0.52 ± 0.057 to 0.572 ± 0.059 , and seasonal fluctuation was observed in the ecosystem of both districts for *P. atherinoides* (Table 2 and Figure 3). Gogoi et al. (2020) observed that the average RGL value was 0.486 ± 0.046 with a minimum of 0.440 ± 0.058 mm and a maximum of

Table 1. Gastro- intestinal length and gastro-intestinal weight of *Pachypterus atherinoides* in both ecosystem of Jhargram and Paschim Medinipur districts.

District	Habitat	Season	GiL (cm)				GiWt (gm)			
			Min.	Max.	Mean	SD	Min.	Max.	Mean	SD
JHARGRAM	POND	Pre-monsoon	2.20	5.80	4.1094	0.90531	0.16	0.47	0.2988	0.09967
		Monsoon	3.60	6.80	5.0188	0.95189	0.04	0.29	0.1763	0.09879
		Post-monsoon	2.20	5.10	3.8594	0.77078	0.04	0.39	0.2338	0.06757
	RIVER	Pre-monsoon	2.6	5.8	4.4281	0.90313	0.19	0.47	0.3175	0.08281
		Monsoon	3.80	6.90	5.2219	0.89973	0.08	0.30	0.1806	0.06345
		Post-monsoon	2.40	5.60	4.1938	0.81119	0.06	0.45	0.2603	0.09654
PASCHIM MEDINIPUR	POND	Pre-monsoon	2.20	5.90	4.1594	0.90371	0.09	0.45	0.2500	0.11843
		Monsoon	3.60	6.80	5.0344	0.95161	0.06	0.31	0.1522	0.07417
		Post-monsoon	2.30	5.10	3.9219	0.76189	0.04	0.33	0.1931	0.07464
	RIVER	Pre-monsoon	2.60	5.90	4.5000	0.92632	0.10	0.44	0.3097	0.09163
		Monsoon	3.90	6.90	5.2375	0.89542	0.04	0.38	0.1991	0.09730
		Post-monsoon	2.40	5.60	4.2063	0.81317	0.04	0.31	0.2013	0.06748

Table 2. Gastro Somatic index (GaSI) and Relative gut length (RGL) of *P. atherinoides* in both ecosystem of Jhargram and Paschim Medinipur districts.

District	Habitat	Season	GaSI				RGL			
			Min	Max.	Mean	SD	Min.	Max.	Mean	SD
JHARGRAM	POND	Pre-monsoon	4.167	8.861	6.65298	1.116203	0.400	0.659	0.52981	0.057801
		Monsoon	0.727	4.578	2.93009	1.001402	0.474	0.63	0.5432	0.049304
		Post-monsoon	3.417	7.947	6.09778	1.053266	0.395	0.622	0.53590	0.058751
	RIVER	Pre-monsoon	5.324	14.331	7.95290	1.906884	0.464	0.949	0.57953	0.091541
		Monsoon	3.095	7.500	6.16186	.967922	0.451	0.633	0.55654	0.048127
		Post-monsoon	4.310	9.845	6.40334	1.012078	0.421	0.691	0.57232	0.059477
PASCHIM MEDINIPUR	POND	Pre-monsoon	3.217	12.925	7.06904	1.812687	0.393	0.678	0.53297	0.057550
		Monsoon	2.727	8.989	4.70767	1.539878	0.468	0.630	0.54480	0.047737
		Post-monsoon	2.639	6.860	4.85507	1.370543	0.411	0.614	0.54326	0.053157
	RIVER	Pre-monsoon	3.689	10.553	7.08806	1.563371	0.441	0.702	0.57125	0.058865
		Monsoon	2.597	8.491	5.41102	1.425112	0.446	0.641	0.55706	0.049073
		Post-monsoon	3.003	8.616	6.09162	1.412353	0.421	0.691	0.56461	0.060830

N=128 Min=Minimum; Max=Maximum; SD=Standard Deviation

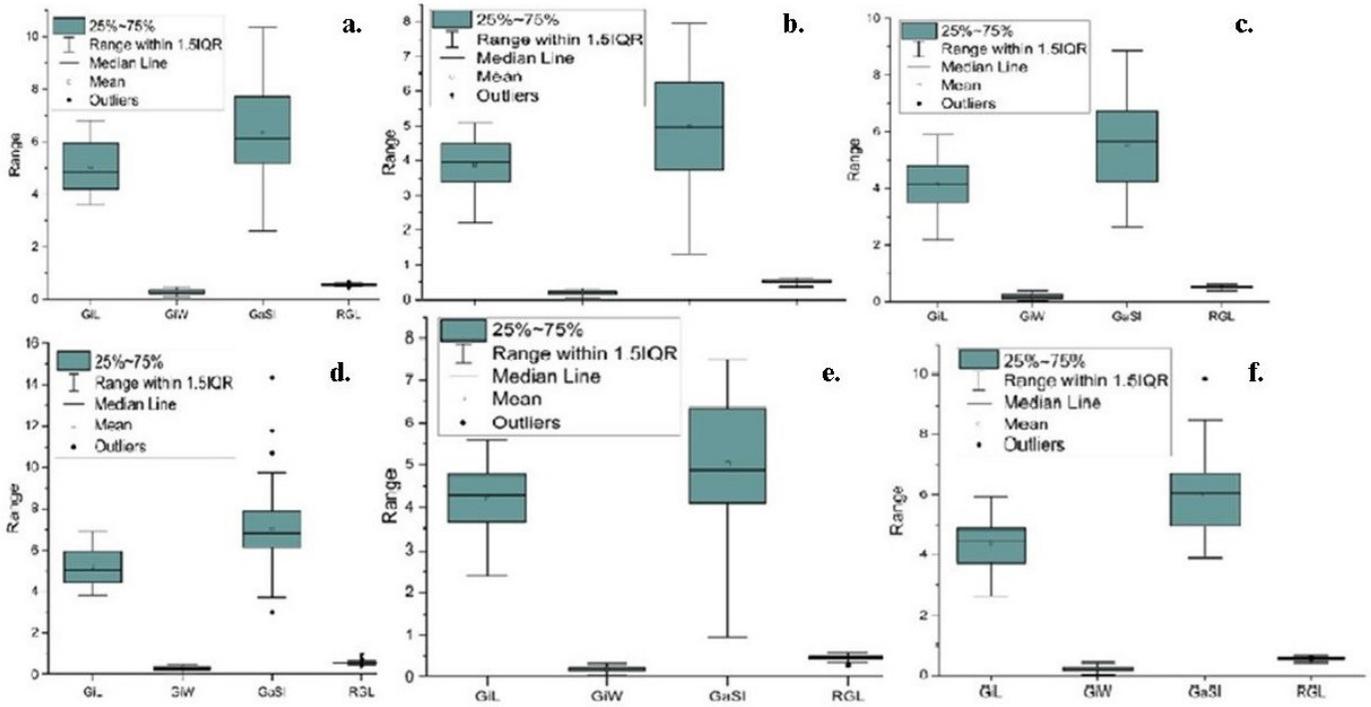


Figure 2. Seasonally GiL, GiWt, GaSI and RGL in both habitat in Jhargram (a,b &c) and Paschim Medinipur (d,e &f) district; a & d-Summer; b & e-Monsoon; c & f-Winter.

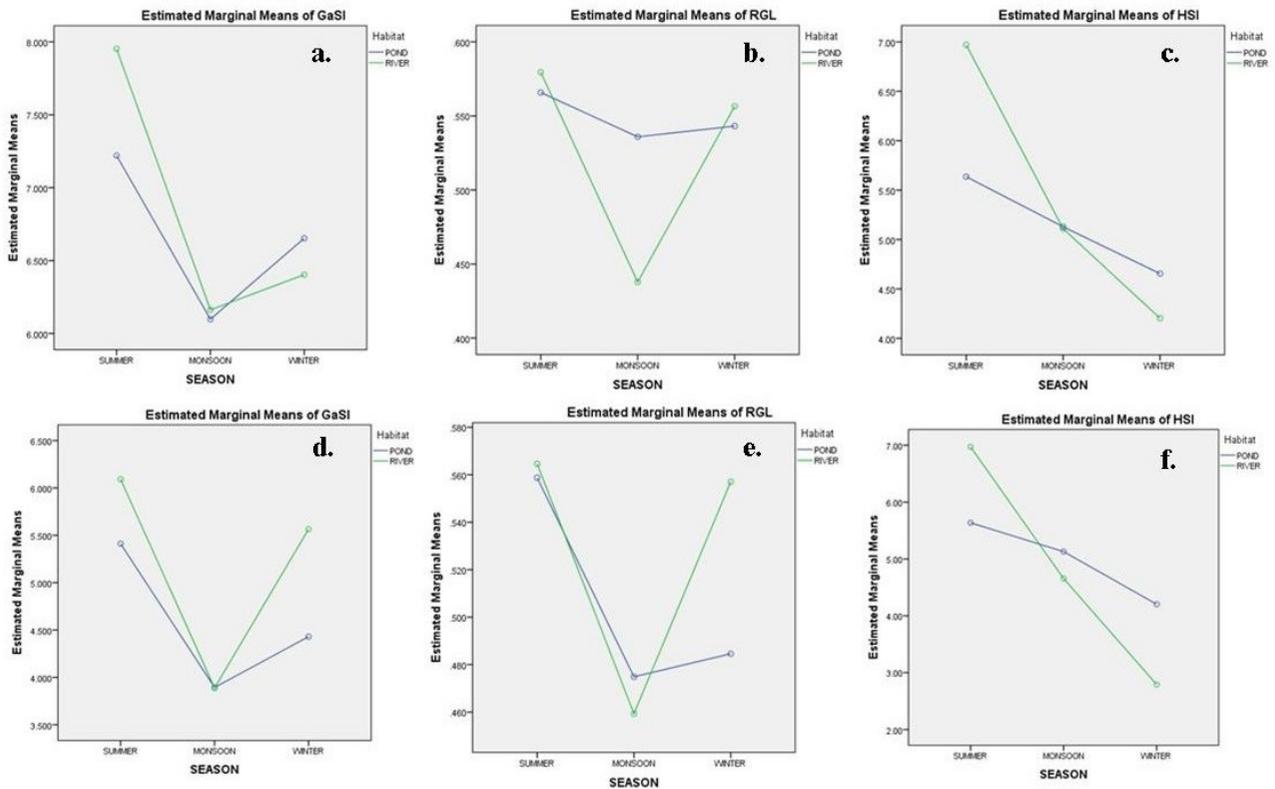


Figure 3. Seasonal variation of GaSI, RGL, HSI in both habitat in Jhargram (a,b &c) and Paschim Medinipur (d,e &f) district; a & d-Summer; b & e-Monsoon; c & f-Winter.

0.511±0.029 mm for *P. atherinoides*. RGL values above 0.8 indicated herbi-omnivorous, 0.7 to 0.8 represented carni-omnivorous, below 0.7 was carnivorous fish, and above 1.6 always herbivorous fish (Koundal *et al.*, 2013), RGL ranged from 0.60 to 0.87 representing omnivorous nature (Dinh *et al.*, 2018), the average RGL value 1.08±0.16 indicated carni-omnivorous nature (Mojumder *et al.*, 2020), RGL ranged between 0.6 to 0.67 with mean SD 0.64 ± 0.003 indicated carnivorous nature (Alam *et al.*, 2020) and very low (<0.32) indicated species as highly carnivorous nature (Renjit Kumar and Roshni, 2021). The present finding agrees with those of Koundal *et al.* (2013), indicating that fish are carnivorous. However, in the current study, the gut contents of *P. atherinoides* show that it eats both plant- and animal-based feeds, including phytoplanktons, zooplanktons, crustaceans, larvae, molluscs, eggs, scales, protozoa, plant parts, and so on. This research revealed the largest proportion of zooplankton accepted as animal food (Figures 4-5). The enormous variety of feeding behaviors that fish have developed is due to evolution, which produced structural adaptations for getting food from an equally large variety of environmental situations (Gupta, 2015). HSI values of female *P. atherinoides* ranged from 4.203 ± 0.979 to 6.97 ± 3.32 with the maximum during Summer (5.63 ± 1.62 and 6.97 ± 3.325) followed by monsoon (7.38 ± 5.23 and 5.11 ± 0.933), Winter (4.65 ± 1.625 and 4.203 ± 0.979) and in male HSI values ranged from 4.10 ± 0.989 to 6.486 ± 1.618 with maximum during Summer (5.51 ± 1.612 and 6.486 ± 1.618) followed by monsoon (5.12 ± 1.093 and 5.00 ± 0.91), Winter (4.567 ± 1.634 and 4.163 ± 0.966) in both pond and river (Table 3 and Figure 3). Cek *et al.* (2001) and Mitu

(2017) reported maximum HSI during the fish breeding season. Jan and Jan (2017), Mojumder *et al.* (2020), Gosavi *et al.* (2020) and Paul *et al.* (2021) observed the minimum HSI during the breeding season. The HSI value in the present study has been supported by the previous work of Jan and Jan (2017), Mojumder *et al.* (2020), Gosavi *et al.* (2020), and Paul *et al.* (2021). The gut of *P. atherinoides* was a maximum of 100 % full from March to May, ¾ full from February to April, ½ full from November to January, and ¼ full and empty from July to October in both habitats in the study area (Figures 6-7). The feeding intensity may be highest before breeding season and lowest during breeding season (Mojumder *et al.*, 2020). The present feeding intensity result is supported by research done by Gupta (2015). Begum *et al.* (2008) stated that fish started to feed heavily after spawning in July, with more than 60% of *M. gulio* having full stomachs; the current data contradicts this. Pearson correlations represent GiLs high positive significant correlation (> 0.01) with GiWt, GaSi with very low positive significant correlation with Giwt, Gil and RGL, HSI with low positive significant correlation (> 0.01) with GiL, GIWt, GaSi, RGL and Liv Wt in both aquatic ecosystems (Tables 4-5 and Figures 8-9). There is a significant difference between the groups in the feeding parameters related to the seasonal and ecosystem (lentic and lotic), according to the multivariate test of *P. atherinoides* in both districts, according to Wilk's lambda (Tables 6 and 9). When studied individually for the six variables, except RGL in Jhargram district, there are no significant differences between the other five parameters towards the lentic and lotic ecosystems in both districts; RGL varied due to food availability in the aquat-

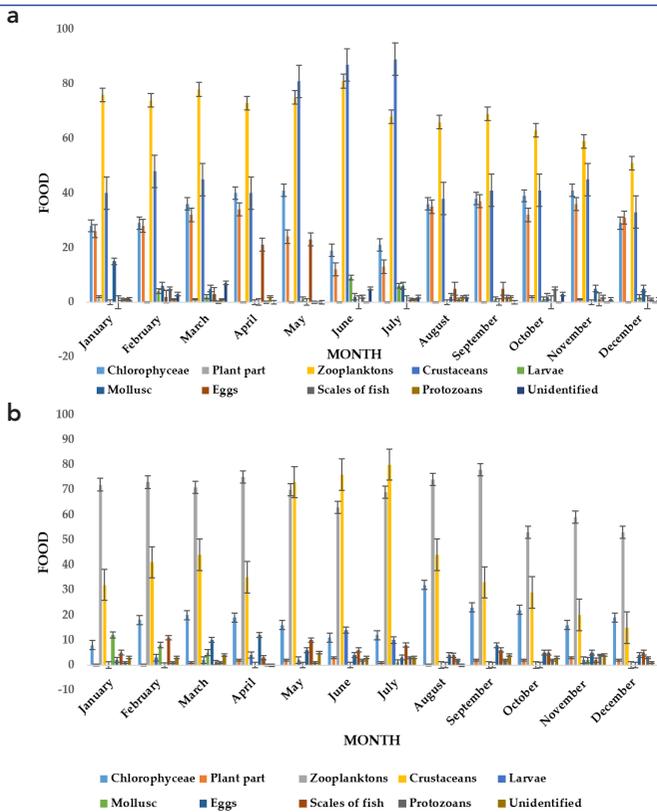


Figure 4. Monthly food item found in gut content of *P. atherinoides* in Jhargram district. a. pond; b. river.

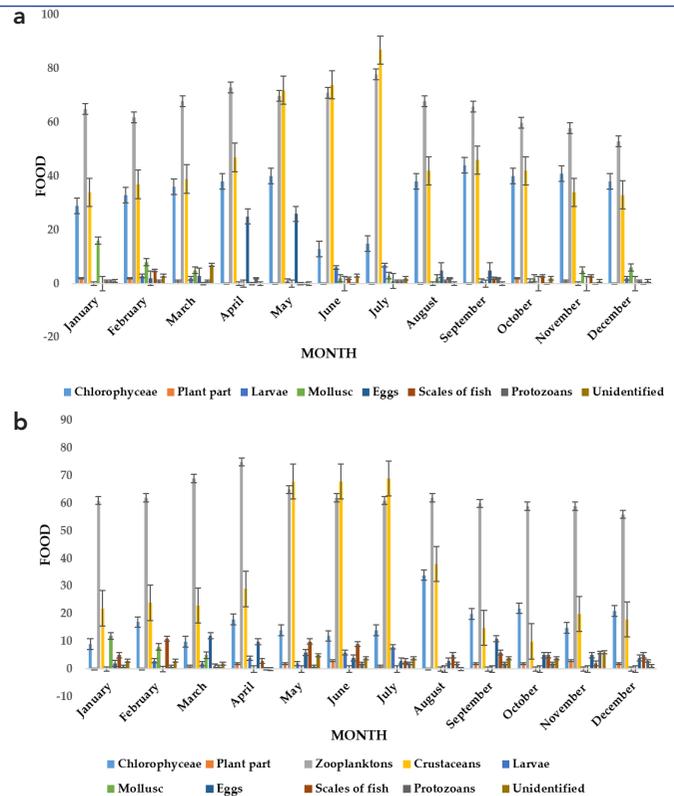


Figure 5. Monthly food item found in gut content of *P. atherinoides* in Paschim Medinipur. a. pond; b. river.

Table 3. Seasonally liver weight and Hepatosomatic index (HSI) of *P. atherinoides* in both ecosystem.

Habitat	Season	Sex	LvWt (g)				HSI			
			Min.	Max.	Mean	SD	Min.	Max.	Mean	SD
POND	PREMONSOON	Female	0.02	0.35	0.1859	0.08904	1.90	8.44	5.6362	1.62325
		Male	0.03	0.26	0.1169	0.06104	1.89	8.34	5.5162	1.61245
	MONSOON	Female	0.12	0.41	0.2531	0.09413	2.34	7.38	5.2297	1.09303
		Male	0.07	0.41	0.2097	0.11338	2.34	7.28	5.1297	1.08304
	POSTMONSOON	Female	0.02	0.24	0.1281	0.06114	1.33	9.61	4.6555	1.62502
		Male	0.02	0.29	0.1513	0.07052	1.33	9.91	4.5675	1.63401
RIVER	PREMONSOON	Female	0.02	0.31	0.1575	0.08673	4.08	12.92	6.9707	3.32530
		Male	0.04	0.39	0.2172	0.09085	4.08	12.42	6.4868	1.61879
	MONSOON	Female	0.14	0.41	0.2653	0.08048	2.62	6.27	5.1103	0.93395
		Male	0.08	0.43	0.2241	0.03422	2.39	6.17	5.0021	0.91013
	POSTMONSOON	Female	0.04	0.24	0.1225	0.05691	2.31	6.22	4.2036	0.97960
		Male	0.02	0.27	0.1591	0.05943	2.21	6.02	4.1063	0.96600

N=128 Min=Minimum; Max=Maximum; SD=Standard Deviation

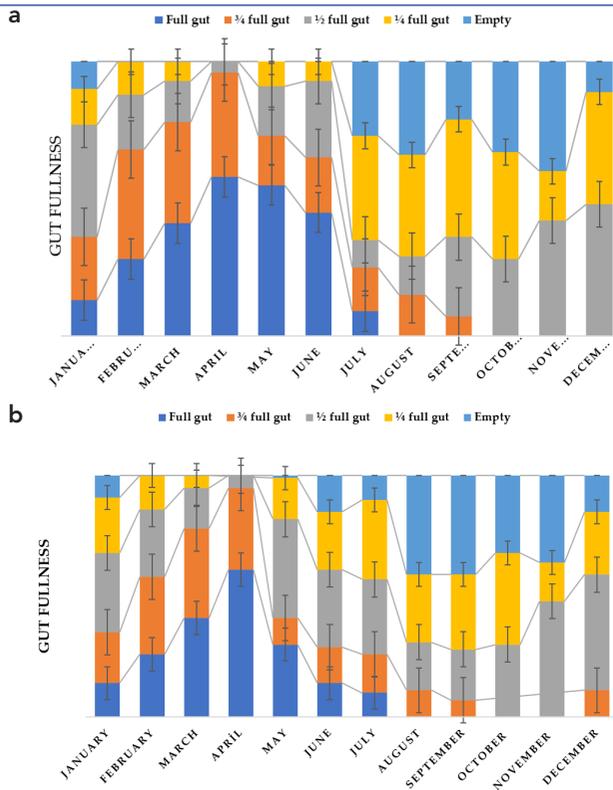


Figure 6. Monthly gut fullness of *P. atherinoides* in Jharghram district. a. pond; b. river.

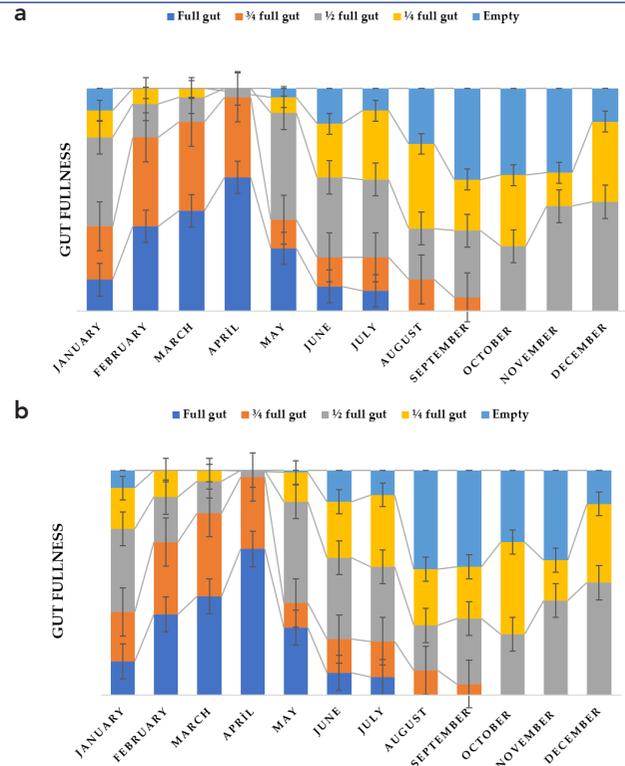


Figure 7. Monthly gut fullness of *P. atherinoides* in Paschim Medinipur district. a. pond; b. river.

ic ecosystem (Tables 7 and 10). Gogoi et al. (2020) stated that Season-wise, GaSI, and RGL were not significant statistically ($p > 0.05$) of *P. atherinoides* in floodplain wetlands of Northeast India; this is not related to the present study. But the seasonal variation of GaSI, and RGL is supported by the study of Kurbah and Bhuyan (2018), Dinh et al. (2018), Gosavi et al., (2020), and Mojumder et al. (2020).

CONCLUSION

The results of the current study make it clear that zooplankton is a vital source of natural food for *P. atherinoides*, although these fish also favored other foods with a plant origin. This finding raises questions about the conventional wisdom that the catfish species only display carnivorous feeding behaviors. On the other hand, it shows a distinct preference for carnivory over herbivory. The availability of such a diverse and partial preference for Phy-

Table 4. Pearson's correlation among feeding biological parameters in Riverine ecosystem for both district.

	GIL	GiWT	LivWt	GaSI	RGL	HSI
GIL	1	.918**	.112	.227**	.436**	.317**
GiWT	.918**	1	.080	.316**	.433**	.351**
LivWt	.112	.080	1	-.050	-.111	.165*
GaSI	.227**	.316**	-.050	1	.330**	.562**
RGL	.436**	.433**	-.111	.330**	1	.437**
HSI	.317**	.351**	.165*	.562**	.437**	1

*: 0.05 level of significance (2-tailed). **: 0.01 level of significance (2-tailed).

Table 5. Pearson's correlation among feeding biological parameters in Pond ecosystem for both district.

	GIL	GiWT	LivWt	GaSI	RGL	HSI
GIL	1	.909**	.037	.244**	.415**	.348**
GiWT	.909**	1	.105	.303**	.457**	.346**
LivWt	.037	.105	1	.115	.154*	.466**
GaSI	.244**	.303**	.115	1	.483**	.412**
RGL	.415**	.457**	.154*	.483**	1	.315**
HSI	.348**	.346**	.466**	.412**	.315**	1

*: 0.05 level of significance (2-tailed). **: 0.01 level of significance (2-tailed).

to-planktivorous resources suggests that the species' eating behaviors are dynamic. This alternate choice of feeding may result from transient adaptation or a phenomenon related to increased resource accessibility. Therefore, this finding represents that the species culture in any aquatic system like a pond, aquarium, or any aquatic body, does not depend on a definite ecosystem for their good growth and development, and it does not require pricey animal protein in its feed, its diverse diet suggests that it may be a suitable species for aquaculture. Additionally, the fact that these little fish species feed could be a crucial evolutionary factor. The study's findings would be valuable tools for developing management and protection strategies for conservation and

Table 6. Multivariate Tests among feeding biological parameters in Jhargram district.

Effect		Value	F	Hypothesis df	Error df	Sig.
Intercept	Pillai's Trace	.999	16493.140 ^b	13.000	174.000	.000
	Wilks' Lambda	.001	16493.140 ^b	13.000	174.000	.000
	Hotelling's Trace	1232.246	16493.140 ^b	13.000	174.000	.000
	Roy's Largest Root	1232.246	16493.140 ^b	13.000	174.000	.000
HABITAT	Pillai's Trace	.241	4.247 ^b	13.000	174.000	.000
	Wilks' Lambda	.759	4.247 ^b	13.000	174.000	.001
	Hotelling's Trace	.317	4.247 ^b	13.000	174.000	.000
	Roy's Largest Root	.317	4.247 ^b	13.000	174.000	.000
SEASON	Pillai's Trace	1.599	53.753	26.000	350.000	.000
	Wilks' Lambda	.033	59.787 ^b	26.000	348.000	.001
	Hotelling's Trace	9.971	66.347	26.000	346.000	.000
	Roy's Largest Root	7.423	99.924 ^c	13.000	175.000	.000

b. Exact statistic; c. The statistic is an upper bound on F that yields a lower bound on the significance level.

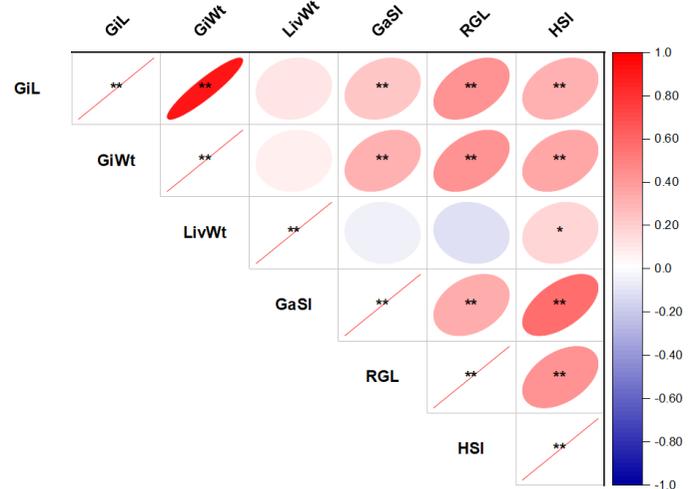


Figure 8. Pearson's correlation of feeding parameters in rivers of both district: *0.05 level of significance, ** 0.01 level of significance.

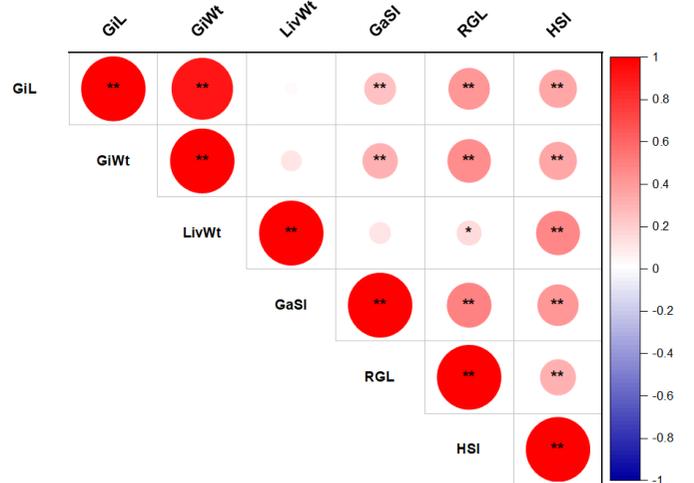


Figure 9. Pearson's correlation of feeding parameters in ponds of both district: *0.05 level of significance, ** 0.01 level of significance.

Table 7. Tests of Between-Subjects Effects of *P. atherinodes* in Jhargram district.

Source	Dependent Variable	Type III Sum of Squares	df	Mean Square	F	Sig.
HABITAT	GaSI	1.594	1	1.594	.903	.343
	RGL	.027	1	.027	6.947	.001
	HSI	3.974	1	3.974	1.232	.269
	GiL	3.910	1	3.910	5.097	.025
	GiWt	.013	1	.013	1.777	.184
	LvWt	.008	1	.008	1.189	.277
SEASON	GaSI	72.574	2	36.287	20.543	.001
	RGL	.253	2	.126	32.690	.001
	HSI	114.963	2	57.482	17.815	.001
	GiL	42.242	2	21.121	27.532	.001
	GiWt	.539	2	.269	36.355	.001
	LvWt	.577	2	.289	45.002	.001

N=256

Table 8. Post Hoc Tests seasonally of *P. atherinodes* in Jhargram district.

Dependent Variable	(I) SEASON	(J) SEASON	Mean Difference (I-J)	Std. Error	Sig.	95% Confidence Interval	
						Lower Bound	Upper Bound
GaSI	SUMMER	MONSOON	1.45692*	.234945	.001	.90183	2.01201
		WINTER	1.05858*	.234945	.001	.50350	1.61367
	MONSOON	SUMMER	-1.45692*	.234945	.001	-2.01201	-.90183
		WINTER	-.39833	.234945	.210	-.95342	.15675
	WINTER	SUMMER	-1.05858*	.234945	.001	-1.61367	-.50350
		MONSOON	.39833	.234945	.210	-.15675	.95342
RGL	SUMMER	MONSOON	.08580*	.010992	.001	.05983	.11177
		WINTER	.02284	.010992	.097	-.00313	.04881
	MONSOON	SUMMER	-.08580*	.010992	.001	-.11177	-.05983
		WINTER	-.06297*	.010992	.001	-.08894	-.03700
	WINTER	SUMMER	-.02284	.010992	.097	-.04881	.00313
		MONSOON	.06297*	.010992	.001	.03700	.08894
HSI	SUMMER	MONSOON	1.1834*	.31754	.001	.4332	1.9337
		WINTER	1.8739*	.31754	.001	1.1237	2.6242
	MONSOON	SUMMER	-1.1834*	.31754	.001	-1.9337	-.4332
		WINTER	.6905	.31754	.078	-.0598	1.4407
	WINTER	SUMMER	-1.8739*	.31754	.001	-2.6242	-1.1237
		MONSOON	-.6905	.31754	.078	-1.4407	.0598
GiL	SUMMER	MONSOON	1.0938*	.15483	.001	.7279	1.4596
		WINTER	.8516*	.15483	.001	.4857	1.2174
	MONSOON	SUMMER	-1.0938*	.15483	.001	-1.4596	-.7279
		WINTER	-.2422	.15483	.264	-.6080	.1236
	WINTER	SUMMER	-.8516*	.15483	.001	-1.2174	-.4857
		MONSOON	.2422	.15483	.264	-.1236	.6080
GiWt	SUMMER	MONSOON	.1297*	.01522	.001	.0937	.1656
		WINTER	.0611*	.01522	.001	.0251	.0970
	MONSOON	SUMMER	-.1297*	.01522	.001	-.1656	-.0937
		WINTER	-.0686*	.01522	.001	-.1045	-.0326
	WINTER	SUMMER	-.0611*	.01522	.001	-.0970	-.0251
		MONSOON	.0686*	.01522	.001	.0326	.1045
LvWt	SUMMER	MONSOON	-.0577*	.01416	.001	-.0911	-.0242
		WINTER	.0763*	.01416	.001	.0428	.1097
	MONSOON	SUMMER	.0577*	.01416	.001	.0242	.0911
		WINTER	.1339*	.01416	.001	.1005	.1674
	WINTER	SUMMER	-.0763*	.01416	.001	-.1097	-.0428
		MONSOON	-.1339*	.01416	.001	-.1674	-.1005

N=256; *. The mean difference is significant at 0.05 level.

Table 9. Multivariate tests of *P. atherinodes* in Paschim Medinipur district.

Effect		Value	F	Hypothesis df	Error df	Sig.
Intercept	Pillai's Trace	.999	20952.388 ^b	13.000	174.000	.000
	Wilks' Lambda	.001	20952.388 ^b	13.000	174.000	.000
	Hotelling's Trace	1565.408	20952.388 ^b	13.000	174.000	.000
	Roy's Largest Root	1565.408	20952.388 ^b	13.000	174.000	.000
HABITAT	Pillai's Trace	.272	4.994 ^b	13.000	174.000	.000
	Wilks' Lambda	.728	4.994^b	13.000	174.000	.001
	Hotelling's Trace	.373	4.994 ^b	13.000	174.000	.000
	Roy's Largest Root	.373	4.994 ^b	13.000	174.000	.000
SEASON	Pillai's Trace	1.585	51.491	26.000	350.000	.000
	Wilks' Lambda	.039	54.783^b	26.000	348.000	.001
	Hotelling's Trace	8.751	58.230	26.000	346.000	.000
	Roy's Largest Root	6.096	82.063 ^c	13.000	175.000	.000
	Roy's Largest Root	.314	4.221 ^c	13.000	175.000	.000

b. Exact statistic; c. The statistic is an upper bound on F that yields a lower bound on the significance level.

Table 10. Tests of Between-Subjects Effects of *P. atherinodes* in Paschim Medinipur district.

Source	Dependent Variable	Type III Sum of Squares	df	Mean Square	F	Sig.
HABITAT	GaSI	17.461	1	17.461	10.818	.347
	RGL	.021	1	.021	9.701	.001
	HSI	11.295	1	11.295	3.598	.059
	GiL	3.658	1	3.658	4.746	.031
	GiWt	.070	1	.070	8.858	.178
	LvWt	6.024	1	6.024	1.223	.270
SEASON	GaSI	112.068	2	56.034	34.716	.001
	RGL	.288	2	.144	66.387	.001
	HSI	196.681	2	98.340	31.325	.001
	GiL	39.883	2	19.941	25.877	.001
	GiWt	.387	2	.194	24.459	.001
	LvWt	13.907	2	6.953	1.411	.246

N=256

Table 11. Post Hoc Test seasonally of *P. atherinodes* in Paschim Medinipur district.

Dependent Variable	(I) SEASON	(J) SEASON	Mean Difference (I-J)	Std. Error	Sig.	95% Confidence Interval	
						Lower Bound	Upper Bound
GaSI	SUMMER	MONSOON	1.86032*	.224588	.001	1.32970	2.39094
		WINTER	.75404*	.224588	.003	.22343	1.28466
	MONSOON	SUMMER	-1.86032*	.224588	.001	-2.39094	-1.32970
		WINTER	-1.10627*	.224588	.001	-1.63689	-.57566
RGL	WINTER	SUMMER	-.75404*	.224588	.003	-1.28466	-.22343
		MONSOON	1.10627*	.224588	.001	.57566	1.63689
	SUMMER	MONSOON	.09457*	.008233	.001	.07512	.11402
		WINTER	.04085*	.008233	.001	.02140	.06030
	MONSOON	SUMMER	-.09457*	.008233	.001	-.11402	-.07512
		WINTER	-.05372*	.008233	.001	-.07317	-.03427
	WINTER	SUMMER	-.04085*	.008233	.001	-.06030	-.02140
		MONSOON	.05372*	.008233	.001	.03427	.07317

Table 11. Continue.

Dependent Variable	(I) SEASON	(J) SEASON	Mean Difference (I-J)	Std. Error	Sig.	95% Confidence Interval	
						Lower Bound	Upper Bound
HSI	SUMMER	MONSOON	2.3427*	.31321	.001	1.6027	3.0827
		WINTER	1.8739*	.31321	.001	1.1339	2.6139
	MONSOON	SUMMER	-2.3427*	.31321	.001	-3.0827	-1.6027
		WINTER	-.4687	.31321	.295	-1.2088	.2713
GiL	WINTER	SUMMER	-1.8739*	.31321	.001	-2.6139	-1.1339
		MONSOON	.4687	.31321	.295	-.2713	1.2088
	SUMMER	MONSOON	1.0719*	.15518	.001	.7052	1.4385
		WINTER	.8063*	.15518	.001	.4396	1.1729
GiWt	MONSOON	SUMMER	-1.0719*	.15518	.001	-1.4385	-.7052
		WINTER	-.2656	.15518	.204	-.6323	.1010
	WINTER	SUMMER	-.8063*	.15518	.001	-1.1729	-.4396
		MONSOON	.2656	.15518	.204	-.1010	.6323
	SUMMER	MONSOON	.0827*	.01573	.001	.0455	.1198
		WINTER	.1042*	.01573	.001	.0671	.1414
	MONSOON	SUMMER	-.0827*	.01573	.001	-.1198	-.0455
		WINTER	.0216	.01573	.358	-.0156	.0587
	WINTER	SUMMER	-.1042*	.01573	.001	-.1414	-.0671
		MONSOON	-.0216	.01573	.358	-.0587	.0156
	SUMMER	MONSOON	-.5797	.39240	.304	-1.5068	.3474
		WINTER	-.0180	.39240	.999	-.9451	.9091
LvWt	MONSOON	SUMMER	.5797	.39240	.304	-.3474	1.5068
		WINTER	.5617	.39240	.327	-.3654	1.4888
	WINTER	SUMMER	.0180	.39240	.999	-.9091	.9451
		MONSOON	-.5617	.39240	.327	-1.4888	.3654

N=256; *. The mean difference is significant at 0.05 level.

captive propagation. The knowledge from the current study may be applied to better manage *P. atherinoides* in India and to future conservation strategies and adoption of these species as possible candidates for commercial aquaculture.

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Revealing Mucilage Event-Linked Community Composition in the Sea of Marmara from eDNA Metabarcoding Data

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ABSTRACT

Mucilage events are among the most conspicuous phenomena in marine ecosystems and present numerous challenges in determining the composition of communities associated with them. To overcome this problem, we used environmental DNA (eDNA) metabarcoding approaches to reveal the species-level resolution of community composition. Mucilaginous aggregates were sampled at six collecting sites during a novel mucilage event (autumn 2021–summer 2022) in the Sea of Marmara, Türkiye. A wide range of plankton community compositions was detected in mucilage samples. eDNA metabarcoding was effective in predicting the community composition of mucilage, which is composed of a wide variety of organisms from mucilaginous aggregates.

Keywords: eDNA, metabarcoding, mucilage, the Sea of Marmara, molecular ecology, biomonitoring

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INTRODUCTION

Marine ecosystems have been facing a greater impact from human activities than that at any other time in history (Berry et al., 2019), driven by a combination of natural factors and hydrological conditions (Mecozzi et al., 2012). These activities, particularly industrialization (Balint et al., 2018), have had negative consequences such as exploitation, pollution, and habitat loss in marine ecosystems (Lotze et al., 2006). Furthermore, intense industrial activities have caused unprecedented changes in ecosystem functions (Cardinale et al., 2012; Smith, 2003) across the world in this era of anthropogenic degradation (Boussarie et al., 2018; Danovaro, Umani & Pusceddu, 2009). As a result of these impacts, one remarkable event is the occurrence of marine mucilage formation that has been reported over the past 200 years, with an increasing trend in frequency for over the past 40 years (Kovač et al., 2023, Faganeli et al., 2010; Turk et al., 2010; Danovaro et al., 2009). Recently in Türkiye, a eutrophication-linked event was also captured by the Landsat 8 – OLI

satellite and shared on the official website of NASA with the title “Blooms in the Sea of Marmara” in the “images day” category in May 2015 (NASA Earth Observatory, 2015) in the Sea of Marmara (SoM). Subsequently, between autumn 2020 and summer 2021, a significant event of mucilage formation was captured by the satellites of Sentinel-2 and Worldview-3 (Tuzcu Kokal, Olgun, & Musaoglu, 2022), which was unprecedented in terms of the amount and duration of observation for the SoM.

Several studies (Fuks et al., 2005; MacKenzie, Sims, Beuzenberg & Gillespie, 2002) have indicated that mucilage phenomenon often begins in regions with stratified water columns. The SoM has also a two-layered water column, where the upper layer is low-salinity Black Sea water, and the lower layer is high-salinity Mediterranean Sea water (Ünlülata, Oğuz, Latif & Özsoy, 1990) with a boundary between them known as a pycnocline (Beşiktepe et al., 1994). The ecosystem of the SoM, which is composed of biological components from these seas, is unique (İşinibilir Okyar, Üstün & Orun, 2015).



Subsequently, the Istanbul (Bosporus) and Çanakkale (Dardanelles) straits have been established as a result of bidirectional dual water-mass exchange events, and studies have indicated that the last occurrence of this event was during the Holocene period at the end of the Würm Glaciation (Çağatay et al., 2009).

In addition to the stratified water columns of the SoM (Çağatay et al., 2015), the effects of anthropogenic activities on the coastal cities of the SoM are of significant concern (Aksu, Balkis, Taşkın & Erşan, 2011) and should be thoroughly investigated (Burak et al., 2009) in terms of mucilage formation. As a nearly enclosed intercontinental basin, the SoM (size ~ 70 × 250 km) (Albayrak, Balkis, Zenetos, Kurun & Erşan, 2006) contains straits that act as biological corridors (Demirel et al., 2023), making it an important region in several manners. To understand the key elements of the SoM that initiate mucilage formations, it is crucial to explore the following points, among others: (i) the SoM is an inland sea between the Anatolian and Thrace peninsulas (Wong, Lüdmann, Ulug & Görür, 1995); (ii) Istanbul, situated on the coast of SoM, is one of the most populous cities in the world with a significant impact on the region's anthropogenic activities (Karaca, 2013); (iii) the region is home to 20% of Türkiye's population and 87% of Türkiye's coastal population (Algan, Balkis, Çağatay, & Sarı, 2004); (iv) industrial activities in the region have caused significant environmental harm, primarily affecting the coastal and shelf areas of the SoM (Korkmaz et al., 2022); (v) the historical peninsula of Istanbul, including the Yenikapı port and the Golden Horn Estuary, as well as well-delineated polluted coastal inlets such as Erdek (Balkis & Çağatay, 2001) and Izmit Bay have typically been significant in human settlement (Lotze et al., 2006) throughout history (Onar et al., 2013; Algan et al., 2004); (vi) intense industrial activities responsible for the majority of anthropogenic discharges to the SoM continue to operate in these areas (Demirel et al., 2023); (vii) a high volume of tanker traffic carrying oil, thousands of vessels per day, poses a constant threat to the SoM ecosystem (Albayrak et al., 2006); and (viii) the metropolitan city of Istanbul, with a population of 16 million in 2023 (Türkiye İstatistik Kurumu, 2023), has been an attraction for people for centuries and is facing ongoing challenges related to waste management and pollution (Güneralp et al., 2021). Moreover, Yaşar (2001) reported that Izmit Bay receives untreated domestic waste from two million inhabitants living around its shores, combined with solid and liquid waste discharge from 300 large industrial plants, contributing to pollution in the bay. Furthermore, as emphasized in studies conducted by Okay et al. (Okay et al., 2001; Okay et al., 1998; Okay et al., 1996), Izmit Bay faces challenges in dealing with toxicity, heavy ship traffic, and petroleum refineries, which supply >30% of Turkey's demand, located on the north-eastern coast of the bay.

Marine biofilms are colonized primarily by surface-associated (Salta et al., 2013) marine organisms (Dang & Lovell, 2016), which are determined by source type, planktonic activities (Gram et al., 2002), and competition between organisms (Bosch, 2013). Extracellular particles, such as DNA, are released by marine organisms, and can be utilized as a supply of nutrients (including carbon, nitrogen, and phosphorus), which are essential for growth and biofilm development (Finkel & Kolter, 2001) of bacterial communities (Das et al., 2013). Surface-associated/attached commu-

nities of microorganisms (Muhammad et al., 2020), zooplankton feces, or feeding structures contain oil compounds, and mucus-rich particles formed by bacteria, which contribute to the formation of marine mucilage (Burd et al., 2020). Zooplankton (e.g., copepods and amphipods) as consumers of marine mucilage or oil provide for the sinking of oil into the seafloor (Almeda, Connelly, & Buskey, 2016; Schwing et al., 2015; Almeda et al., 2014; Fisher et al., 2014; Montagna et al., 2013; Mitra et al., 2012; Conover 1971) and also contribute to the production of marine oil snow with their fecal pellets (Burd et al., 2020). The dynamic and heterogeneous community structure of marine biofilms makes it challenging to model and investigate. Fungi secrete extracellular polymeric substances (Metzger et al., 2009), similar to those secreted by bacteria and phytoplankton, and particularly diatoms (Wotton, 2004). These substances act as a "glue" that holds the different components of marine mucilage together (Burd et al., 2020). Furthermore, the reflection of key environmental factors of a substratum by biofilms has vital implications for larval settlement of marine invertebrates (Dobretsov, 2010).

Despite numerous studies having emphasized the presence of mucilage in the SoM through traditional marine surveillance programs (Toklu-Alıclı, Özdelice & Durmus et al., 2021; Balkis-Ozdelice, Durmuş, & Balci, 2021; Toklu-Alıclı, Polat & Balkis-Ozdelice et al., 2020; Tas, Kus & Yılmaz, 2020; Ergul et al., 2021, İşinbilir-Okyar et al., 2015; Yılmaz, 2015; Altıok & Kayışoğlu, 2015; Balkis et al., 2011; Tüfekçi et al., 2010) such as "continuous plankton recording" and/or "trawl," (Zingone et al., 2021), these methods are not always easy to use or cost-effective (Zaiko et al., 2015; Barbour, 1999; Jennings & Kaiser, 1998) to monitor the biodiversity (Thomsen et al., 2016). Nevertheless, emerging DNA-based tools, such as environmental DNA (eDNA) metabarcoding, are cost-effective, nature-friendly, and widely adopted and promising for future applications (Zaiko, Samuiloviene & Ardura, 2015; Pochon et al., 2013; Wood et al., 2013; Darling & Mahon, 2011). Mucilage phenomena have been one of the most conspicuous in terms of characterization (Pompei et al., 2003), and their characterization involves several difficulties. However, eDNA analysis (Ogram et al., 1987) is a molecular tool that can overcome some of the abovementioned difficulties (Genitsaris, Stefanidou & Sommer, 2019; Zaiko et al., 2015; Del Campo et al., 2014; Mächler et al., 2014; Bik et al., 2012) to ensure effective biomonitoring. The emerging science of eDNA (Kelly et al., 2014) refers to the genetic remnants of life that can be obtained from a wide range of environmental samples (Taberlet, 2012). This method can provide detailed taxonomic resolution even when the samples are bulk mixtures of organisms obtained via plankton tows (Berry et al., 2019; Taberlet et al., 2018; Thomsen & Willerslev, 2015).

In this study, we applied eDNA metabarcoding approaches to explore the poorly studied biodiversity of marine mucilage/snow using a molecular approach. We built on the study of Doğan et al. (in press) and going to the species-level characterization, we attempt to help understand the possible linkages between the components, including eukaryotic organisms (algae, fungi, and animalia), and community composition of mucilage samples, using the CO1 gene, collected from selected stations in the SoM during the novel 2020–2021 mucilage event, as mentioned earlier.

MATERIALS AND METHODS

During the Marmara Sea and North Aegean Sea Expedition (in June 2021), three replicates of mucilage samples were obtained from six stations (M1–M6) (Figure 1), targeting 0–1 m of the surface seawater layer in the water column, via Yunus-S Research/Vessel (R/V). We followed and modified the protocol outlined by Buxton et al. (2021) and collected three samples of mucilage using negative field controls, according to Keskin (2014), at each station.

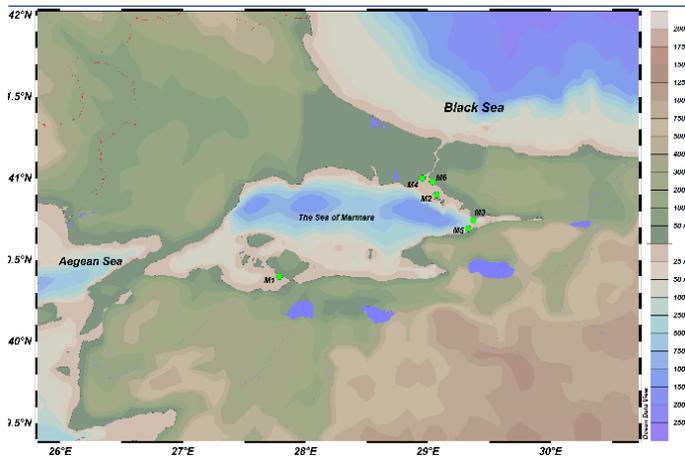


Figure 1. Study area in the Sea of Marmara, (M) indicating the location of the six sampling sites: M1, M2, M3, M4, M5, and M6, as the Bay of Erdek, Kinalıada, Bay of İzmit, Yenikapı, Çınarcık, and Kalamış, respectively.

DNA extraction was performed using the DNeasy Plant Pro Kit-Qiagen (Germany) according to the manufacturer's instructions. The primer pairs of CO1, as described elsewhere (Leray, et al., 2013), were used for the initial PCR assay. The index primers were incorporated into the second PCR assay based on their specific locations and genes. The PCR products were purified using AMPure XP beads (Beckman Coulter, Brea, CA, USA). The concentration of the pool was determined and validated using the KAPA Library Quantification qPCR kit (Roche, Germany). The pool was then sequenced on an Illumina platform (Illumina, USA) with paired reads (2×150 bp) (Gen Era Diagnostics Inc., Türkiye).

Following the sequencing of NovaSeq, S4, 2×150 bp read length service, the raw sequences were received as demultiplexed fastq files and processed using The Advanced Pipeline for Simple yet Comprehensive Analyses of DNA metabarcoding data (the APSCALE graphical user interface) "apscale_gui" pipeline v1.2.0 (https://github.com/TillMacher/apscale_gui) (Buchner, Macher & Lesse 2022), which is based on VSEARCH (Rognes, Flouri, Nichols, Quince & Mahe, 2016) and cutadapt (Martin, 2011). This module consisted of demultiplexing, paired-end merging, primer trimming, quality & length filtering, dereplication & pooling, Operational Taxonomic Units (OTU) clustering, denoising (ESVs), chimera removal, LULU filtering, and remapping steps.

The analysis commenced with the adapter primer trimming stage, as the raw data had already been demultiplexed by the sequencing company. The early stages of the analysis pipeline comprised the deletion of primer sequences and tags by cutadapt (Martin, 2011) and the evaluation of the quality of each read based on specific per-base quality and read length thresholds. The pooled and dereplicated reads were clustered into OTUs based on the similarity threshold (97%) and denoised into "Exact Sequence Variants" (ESVs). Chimeras were automatically detected and removed from the OTUs and ESVs using the vsearch -uchime_denovo command. OTUs and ESVs were mapped against the dereplicated files. The LULU filtering algorithm (Frøslev et al., 2017) was used to reduce the number of erroneous OTUs/ESVs and achieve more realistic biodiversity metrics. Finally, the OTUs and ESVs were remapped to the sequences of each sample, and read tables were generated. A taxonomic assignment was performed using the BOLD system with the BOLDigger module (Buchner & Leese, 2020) and Midori2 (MIDORI2_UNIQ_NUC_GB257_CO1_BLAST) in the local blast module. The final taxonomy table was generated using the "JAMP filter" option (Elbrecht, 2022). During this process, ambiguous records were flagged along with their respective situation.

The TaxonTableTools (TTT) v1.4.8 (Macher, Beermann & Leese, 2021) was used for downstream analyses. A taxon table was generated using the taxon table converter module, with the taxonomy and read tables generated in previous analyses serving as input. The replicates were merged, considering of the consistency of the OTUs present in each replicate. Subsequently, progress was made in the taxon table filtering module, where taxa exhibiting <85% similarity were eliminated. The read threshold was set at 0.01%. Negative control subtraction was performed as the final step.

The assigned taxonomy of each OTU in the taxon table was validated against the Global Biodiversity Information Facility (GBIF) database (<https://www.gbif.org/>). Consequently, any spelling errors were corrected, and the synonyms for each respective taxon were examined and updated automatically. The reads were processed by normalization and metadata were generated to facilitate the execution of downstream analyses. Basic statistics (read count, OTU number, taxonomic resolution, and richness) were generated. Venn diagrams were generated by comparing the BOLD and Midori taxon tables at each taxonomic level, ranging from OTU numbers to phyla. Rarefaction analyses were performed using both all-in-one and per-taxon approaches with the aim of determining the number of species and/or OTUs that separated among each respective taxonomic group. The species-level site occupancy was calculated using a heatmap representation of presence/absence of data. Per-taxon statistical analyses were conducted to retrieve the count of reads, number of OTUs, and species. Parallel categorical analyses were performed to determine the samples that were represented by species and/or higher taxonomic levels. The distribution of designated taxa was confirmed by cross-referencing with the GBIF using an Application Programming Interface (API). At the phylum and species levels, read proportions were calculated and plotted using heat maps, bar charts, and pie charts. Krona charts were gener-

ated to display both single sample and combined data points (Supplementary Data).

Diversity analyses were conducted by computing both the alpha and beta diversity measures. Specifically, we determined alpha diversity by calculating the number of OTUs per sample, and for beta diversity, we computed Jaccard distances between samples. The taxon table was used to generate a taxon list that incorporated all recognized taxa and diminished redundancy. In cases where numerous OTUs were assigned to the species level, the maximum, average, and minimum genetic distances between OTUs were calculated to discern highly diverse or cryptic species.

RESULTS AND DISCUSSION

Analyzing the CO1-based community composition within mucilage samples obtained from collection sites

Following the sequencing process, the raw reads underwent processing, resulting in the trimming of 3,078,342 M reads, with an average of 1,669,203 and 1,284,649 reads that successfully passed the quality-filtering step. Subsequently, the preprocessing steps were clustered into 1657 OTUs for further analysis. Of these, 429 records were found as “No Match,” 172 records were found as “blank,” and the remaining 1056 records were assigned to OTUs. After curation (merging of replicates and subtraction of negative controls), 431 OTUs (629,535 reads) with a similarity of $\geq 85\%$ to the reference sequence remained and were selected for downstream analyses. Among the records of the 431 OTUs, 22 were ambiguous. The average sequence length was 118 bp, with a minimum length of 100 bp and a maximum length of 147 bp. A total of 34 phyla, 78 classes, 157 orders, 106 families, 65 genera, and 57 species, all of which were unique to each taxonomic level, were observed against the Midori2 database.

When we compared our basic statistics in both Midori2 and BOLD databases, 22 phyla were detected as shared between the two databases, including Amoebozoa, Arthropoda, Ascomycota, Bacillariophyta, Bryozoa, Cnidaria, Echinodermata, Mollusca, Porifera, Rhodophyta, and Rotifera. In addition, Midori2 identified 10 phyla, including Bigyra, Cercozoa, Choanoflagellata, and Eusea, and BOLD identified 5 phyla, including Heterokontophyta, Onychophora, and Zygomycota. At the class level, the 39 taxonomic groups that were detected by both Midori2 and BOLD databases and were the most abundant based on OTUs and species were Anthozoa, Bivalvia, Bacillariophyceae, Copepoda, Dinophyceae, Gastropoda, Hydrozoa, Rhodophyta, Scyphozoa, and Staurozoa. The two databases under consideration shared 214 OTUs (Figure 2a) and 23 species (Figure 2b), whereas Midori2 detected an additional 32 species not found in BOLD, and BOLD detected 16 species not found in Midori2. Moreover, Midori2 detected 217 OTUs not present in BOLD, whereas BOLD detected 143 OTUs not present in Midori2 (Figure 2). The analysis of the two databases revealed that a significant overlap exists between the two, with approximately 20% of OTUs (Figure 3a) and species (Figure 3b) being shared between them.



Figure 2. A comparison of Midori2 and BOLD databases focusing on (a) OTU and (b) species in each database.

The relative abundance and distribution proportions of the orders across different samples were investigated by comparing the BOLD and Midori2 databases. Amphipoda, Caenogastropoda, Ceramiales, Cheliostomatida, Corallinales, Diplostraca, Gastrochaenida, Perciformes, Sessilia, Stauromedusae, and Thalassiosirales correlated in both databases (Figure 4).

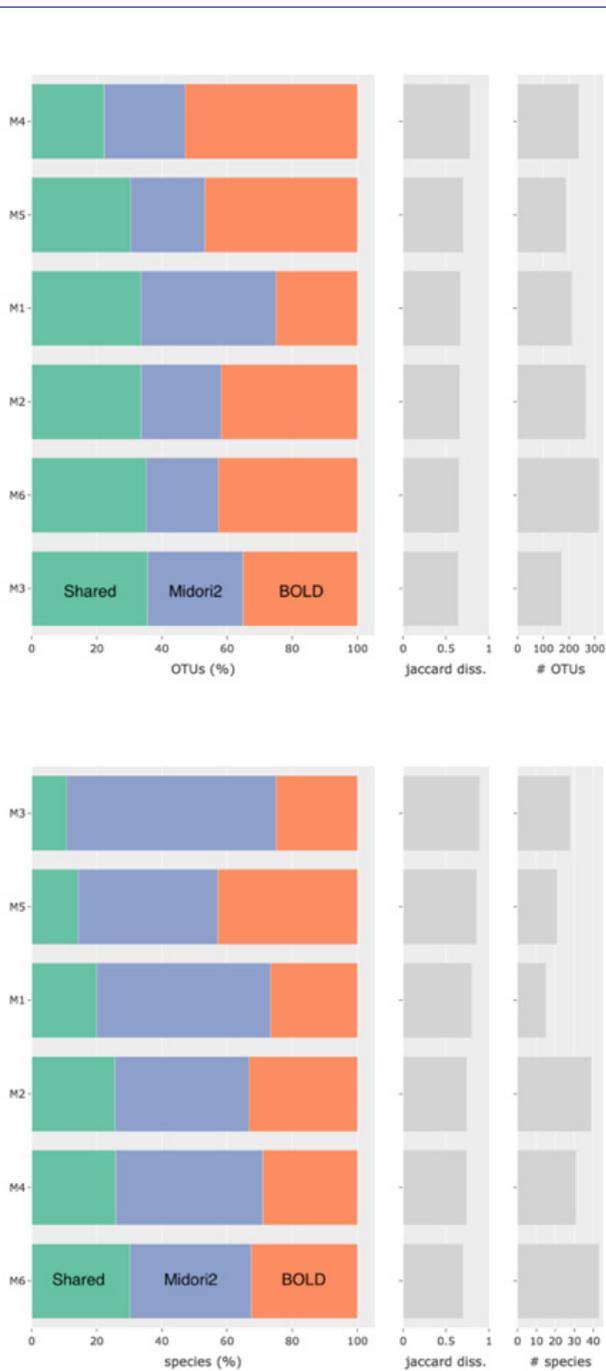


Figure 3. (a) OTUs and (b) species sources between Midori2 and BOLD databases.

At the species level, taxonomic richness retrieved from BOLD and Midori2 databases correlated per phylum. Arthropoda, Rhodophyta, Mollusca, and Ochrophyta were the most prominent phyla (Figure 5).

By selecting the top 10 phyla containing the most OTUs, the sample-based rarefaction analysis demonstrated that 15/52 species observed in the first sample subsequently increased to 41/52 in the all-in-one method (Figure 6a). Testing the species and

OTUs to separate among phyla (Figure 6b) revealed that the phylum Ochrophyta reached eight species (8/8) in the second sample, which flattened in subsequent samples. The phylum Mollusca reached five species in the third sample and ended with seven (7/7) species in the second sample. The phylum Cnidaria reached two species (2/2) in the third sample and remained constant throughout the samples (Figure 6b).

The M6 sample exhibited the greatest degree of alpha diversity among all samples, whereas the M3 and M5 samples were limited to fewer than 115 OTUs/30 species each (Figure 7). The Kalamış (M6) sample had 181 OTUs (Figure 7a) (29 OTUs at species level) and 23 species (Figure 7b), following the M1 sample with 159 OTUs. The M3 and M5 samples exhibited the lowest degree of alpha diversity among all samples, as evidenced by the presence of 111 and 100 OTUs, respectively. Despite the comparatively small number of OTUs in M1 and M5 samples, the number of reads at these samples was substantial, with 140 k reads recorded in the M1 sample and 108 k reads in the M5 sample. Compared with the M1 sample, which had a low number of OTUs at the total (111) and species level (13) and 33 k reads, the M2 sample exhibited a higher number of OTUs of 26 at the species level and 121 k reads (Figure 7).

A total of 87 OTUs were identified in the phylum Arthropoda, 73 OTUs were identified in the phylum Ochrophyta, 46 OTUs were identified in the in the phylum Rhodophyta, and 41 OTUs were identified in the in the phylum Mollusca. Conversely, the remaining 27 phyla were represented by fewer than three species each. Various taxonomic groups such as Amoebozoa (27), Chlorophyta (7), Dinoflagellata (4) from Protozoa, Cnidaria (14), Porifera (6), and Echinodermata (3) from Animalia showed the presence of OTUs. Overall, 87 OTUs of the phylum Arthropoda were assigned to 15 distinct species, 73 OTUs of the phylum Ochrophyta were assigned to species, 41 OTUs of the phylum Mollusca were assigned to six species, and 46 OTUs of the phylum Rhodophyta were assigned to five species. Despite comprising the most numbers of OTUs, the phylum Arthropoda exhibited a relatively low proportion of reads by <4%. In contrast, members of the phylum Cnidaria (Figure 8a), including *Aurelia aurita*, exhibited a dominant presence, accounting for >40% of all reads (for samples with >97% similarity) (Figure 8b). Members of Ascomycota, including *Cladosporium allacinum*, accounted for 9% of the reads, followed by the members of Mollusca, such as *Bittium reticulatum*, accounting for 5% of the reads, whereas the coccolithophore *Emiliania huxleyi* (3%) was also represented.

Among the identified Arthropoda OTUs, the class Insecta consisted of the most abundant OTUs, with 56 OTUs comprising >60% of the OTUs. Despite this, only seven species were assigned to this class. In contrast, the class Copepoda that included five OTUs was well represented by three distinct species and comprised >80% of Arthropoda reads. The calanoids *Paracalanus parvus* (83% reads of Copepoda) and *Pseudocalanus elongatus* (13% reads of Copepoda) were dominant among copepod species, with low read representation by the cyclopoid *Oithona similis* (1% reads of Copepoda and 56% of Cyclopoida), with 100% similarity. Furthermore, the maxillopods *Amphibalanus amphitrite* (96% reads of Maxillopoda) and *Amphibalanus impro-*

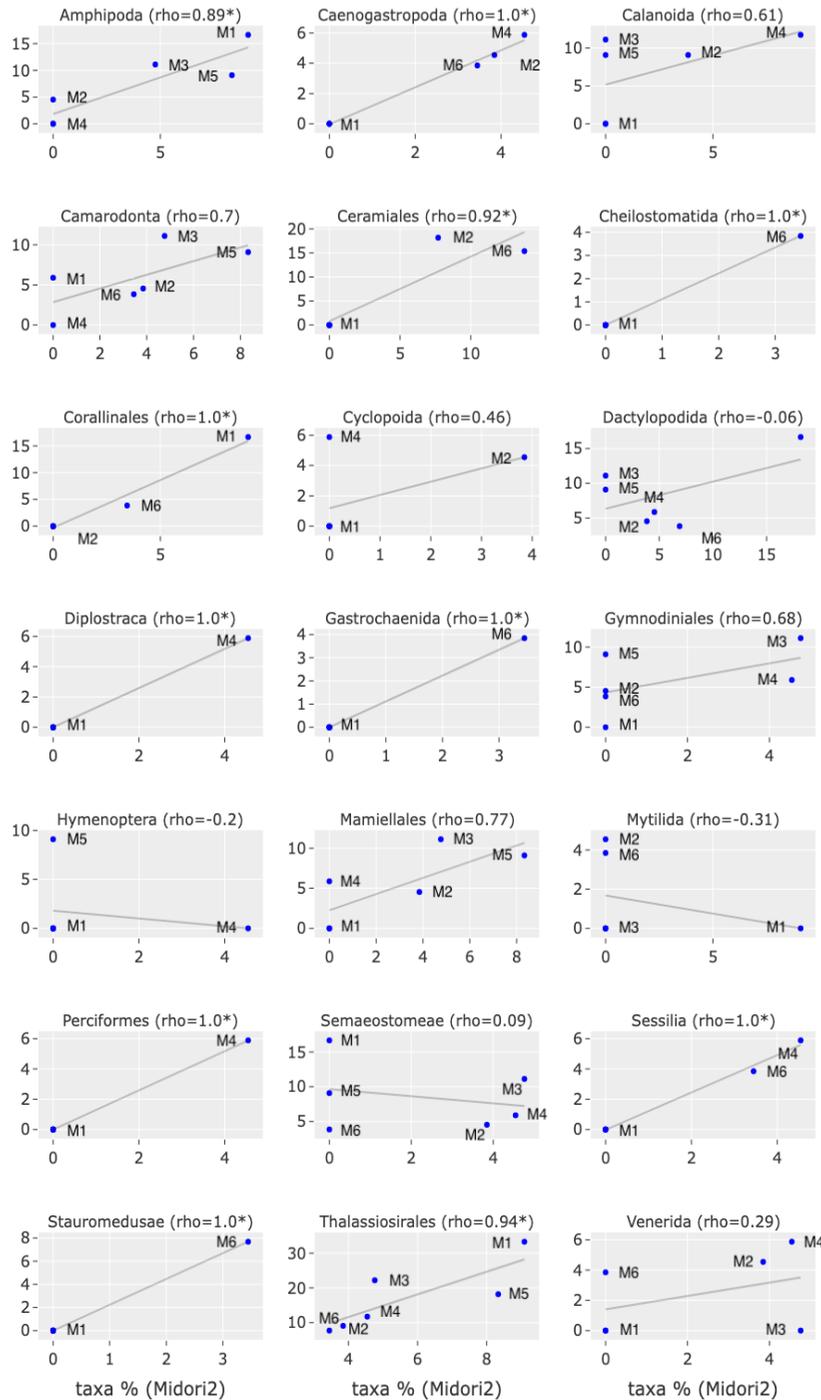


Figure 4. Taxon proportions, pairwise analysis of samples (including >98% similarity): rho (Spearman's rank correlation coefficient), samples marked with an asterisk indicate a positive monotonic relationship (they display consistency in the same direction).

visus (4% reads of Maxillopoda) and the branchiopod *Pleopsis polyphemoides* were represented by three OTUs with 100% similarity and comprised 10% reads of arthropods. Remarkably, nine OTUs were classified as ambiguous and subsequently assigned to the same species, viz., *Lucilia caesar*, which belongs to the order Diptera.

A total of 73 OTUs were detected within the phylum Ochrophyta. In particular, 12 OTUs belonged to the order Ectocarpales, and 16 OTUs were assigned to the order Naviculales within the phylum Ochrophyta. Among the phylum Rhodophyta, 25 OTUs were identified within the order Ceramiales, and the species-level match was well represented by the following species: *Apoglos-*

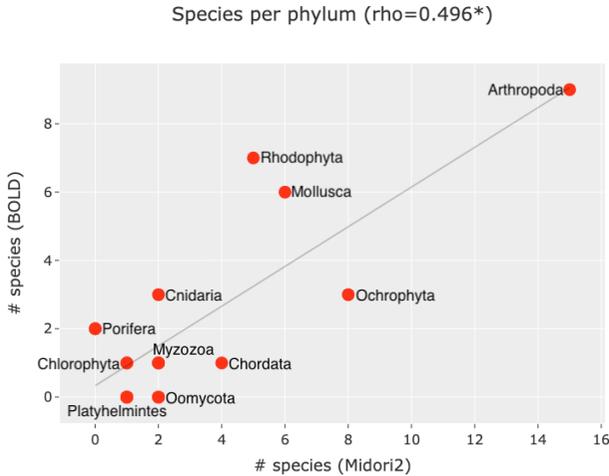


Figure 5. Comparisons of taxonomic richness across each database (BOLD and Midori2) based on phylum level.

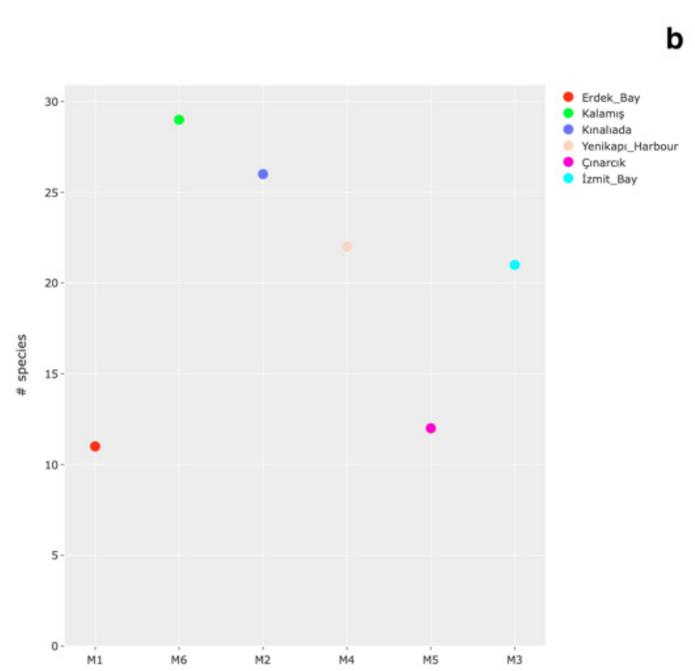
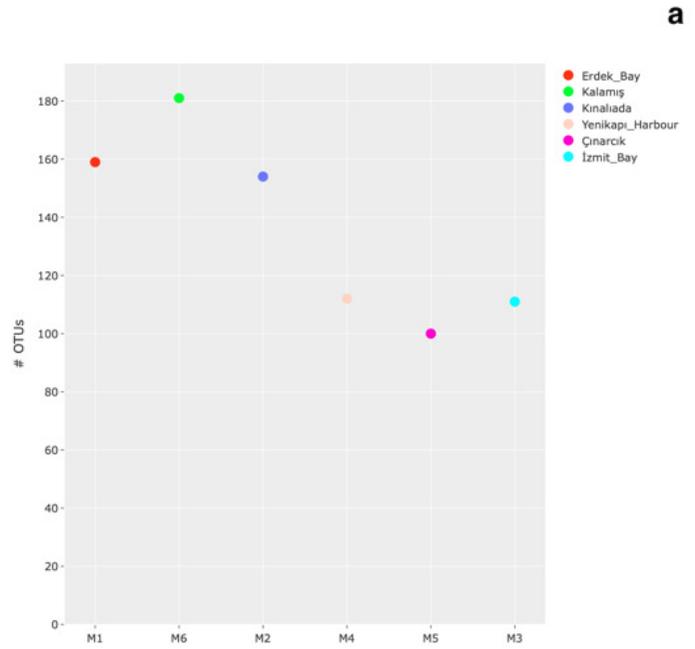


Figure 7. Alpha diversity, comprising (a) the number of OTUs and (b) the number of species per sample.

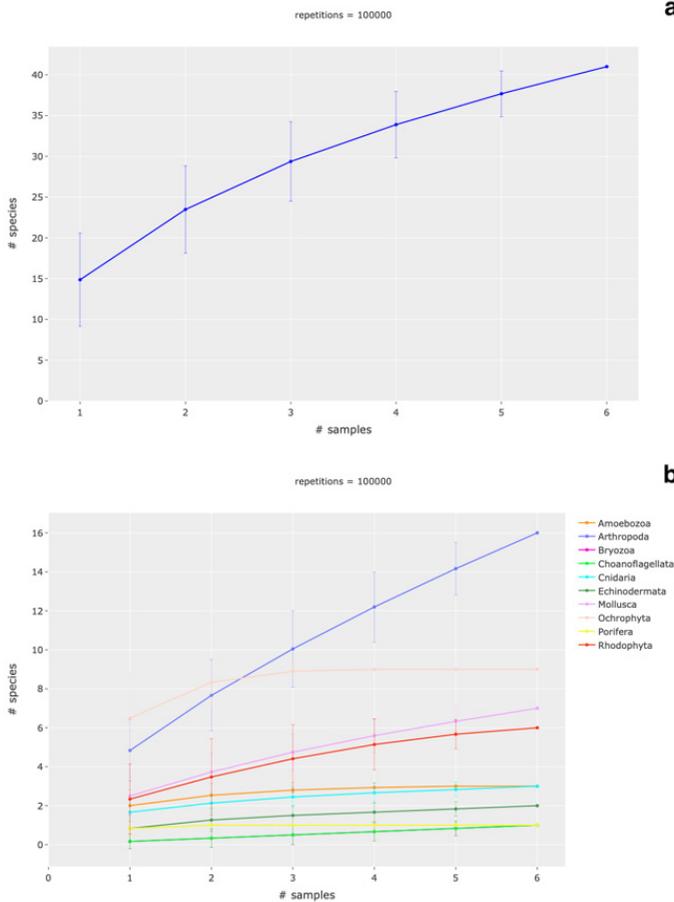


Figure 6. The use of sample-based rarefaction was examined through the application of both all-in-one (a) and per-taxon (b) approaches (100,000 repetitions) in top 10 phyla.

sum ruscifolium (19% of Rhodophyta reads), *Polysiphonia morrowii*, *Dasysiphonia japonica*, and *Pterothamnion crispum*, all of which exhibited 100% similarity. Furthermore, 11 OTUs were identified within the order Bacillariales that exhibited a remarkable abundance of reads within the phylum Ochrophyta, which comprised three OTUs identified for *Cylindrotheca* and two OTUs identified for the genus *Nitzschia*. The order Thalassiosirales comprised three OTUs, which included a two-species-level representation of *Skeletonema pseudocostatum* (1% of all reads)



Figure 8. The proportions of reads for (a) each phylum (including all data) and (b) species (for samples with >97% similarity) in a heat map and bar graph among samples.

(observed among all mucilage samples) and *Skeletonema* sp., with 100% similarity. Although taxonomic assignment was 100% similar to the genus *Skeletonema* due to the multiple species hits (*Skeletonema dohrnii*, *Skeletonema marinoi*, and *Skeletonema costatum*), these OTUs corresponded to a single genus (*Skeletonema*). Nevertheless, despite the presence of 12 OTUs in the

order Ectocarpales, no OTUs were assigned to species-level representation. The order Lithodesmiales was represented by *Ditylum brightwellii*. The order Chaetocerotales comprised two OTUs that were assigned to *Chaetoceros socialis* with 100% similarity. The order Rhizosoleniales was represented by two OTUs, both of which were assigned to the family Rhizosoleniaceae.

Among the protists, 27 OTUs were identified within the phylum Amoebozoa, exhibiting a high similarity of 85%–93% to the class Discosea. In addition, two OTUs and one taxa *Squamamoeba japonica* were well represented at the species level, with 100% similarity. Four OTUs and two species were identified within Dinoflagellata as follows: *Pfiesteria piscicida* belonged to the order Peridiniales, and *Cochlodinium polykrikoides* and *Gyrodinium instriatum* belonged to the order Gymnodiniales. Although five OTUs were assigned to the Bigyra class, no species were identified within this assignment. The coccolithophore *Emiliana huxleyi* (3% of all reads) that belonged to the phylum Haptophyta was represented by a single OTU. A total of 49 OTUs were identified within Fungi, of which 42 were present in both Oomycota and Basidiomycota, with 22 and 18, respectively, and 9 OTUs were exclusive to Ascomycota. Although Fungi exhibited a relatively high representation, only three OTUs could match at the species level, with their similarity exceeding 100%. These OTUs belonged to the following species: *Cladosporium allacinum* (9% of all reads and 95% of Ascomycota reads), *Aspergillus puulaausensis*, *Globisporangium spinosum* (Oomycota), and *Anisopolidium ectocarpii* (Basidiomycota).

An analysis of the phylum Porifera revealed six OTUs, with five of these belonging to the Demospongiae class and the remaining one belonging to the Homoscleromorpha class. The Demospongiae class was represented by four different orders in Midori2 and *Halisarca desqueyrouxae* with 99% similarity, which belongs to the Halisarcidae family in the BOLD database. A total of 12 OTUs were identified in the phylum Cnidaria, of which 5 were assigned to the Hydrozoa class, and two species were identified. One of the OTUs matched with the Hydractiniidae family with 93% similarity, and three of them were assigned to the Staurozoa class, with one matched with the Lucernariidae family at 91% similarity and another matched with the order Stauromedusae at

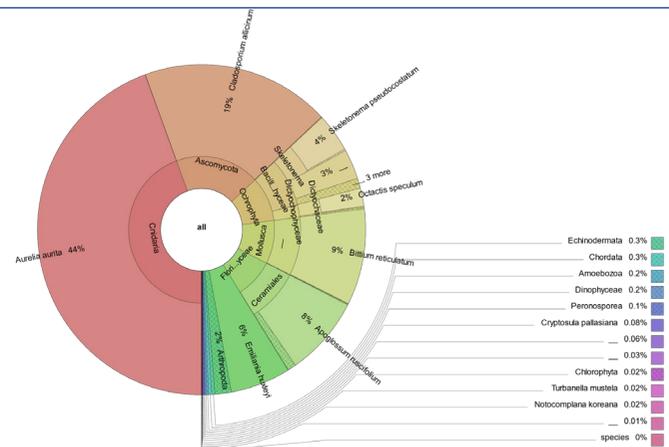


Figure 9. Krona charts based on read proportions (>97% similarity) when sampling locations were merged.

In contrast to the representative phyla (Figure 12a), the absence/presence of species was not uniform across all regions (NM and EM), except for a few cosmopolitan organisms such as *Aurelia aurita*, *Aureococcus anophagefferens*, *Cladosporium allcinum*, *Corallina caespitosa*, *Emiliana huxleyi*, *Micromonas pusilla*, *Neoparamoeba aestuarina*, *Octactis speculum*, *Paracentrotus lividus*, *Cochlodinium polykrikoides*, *Gyrodinium instriatum*, *Pfiesteria piscicida*, *Pseudochattonella farcimen*, *Skeletonema pseudocostatum*, and *Vicicitus globosus* (Figure 12b).

This study represents one of the first extensive investigations to utilize metabarcoding approaches to analyze community composition, including protists, fungi, and animalia, associated with the mucilage events that occurred during the novel 2020–2021 mucilage event in the SoM. Results revealed the community composition of the mucilage and identified the predominant microorganisms thriving among samples.

Taxonomic assignment revealed that the mucilage samples were categorized into three orders of the main category and were rep-

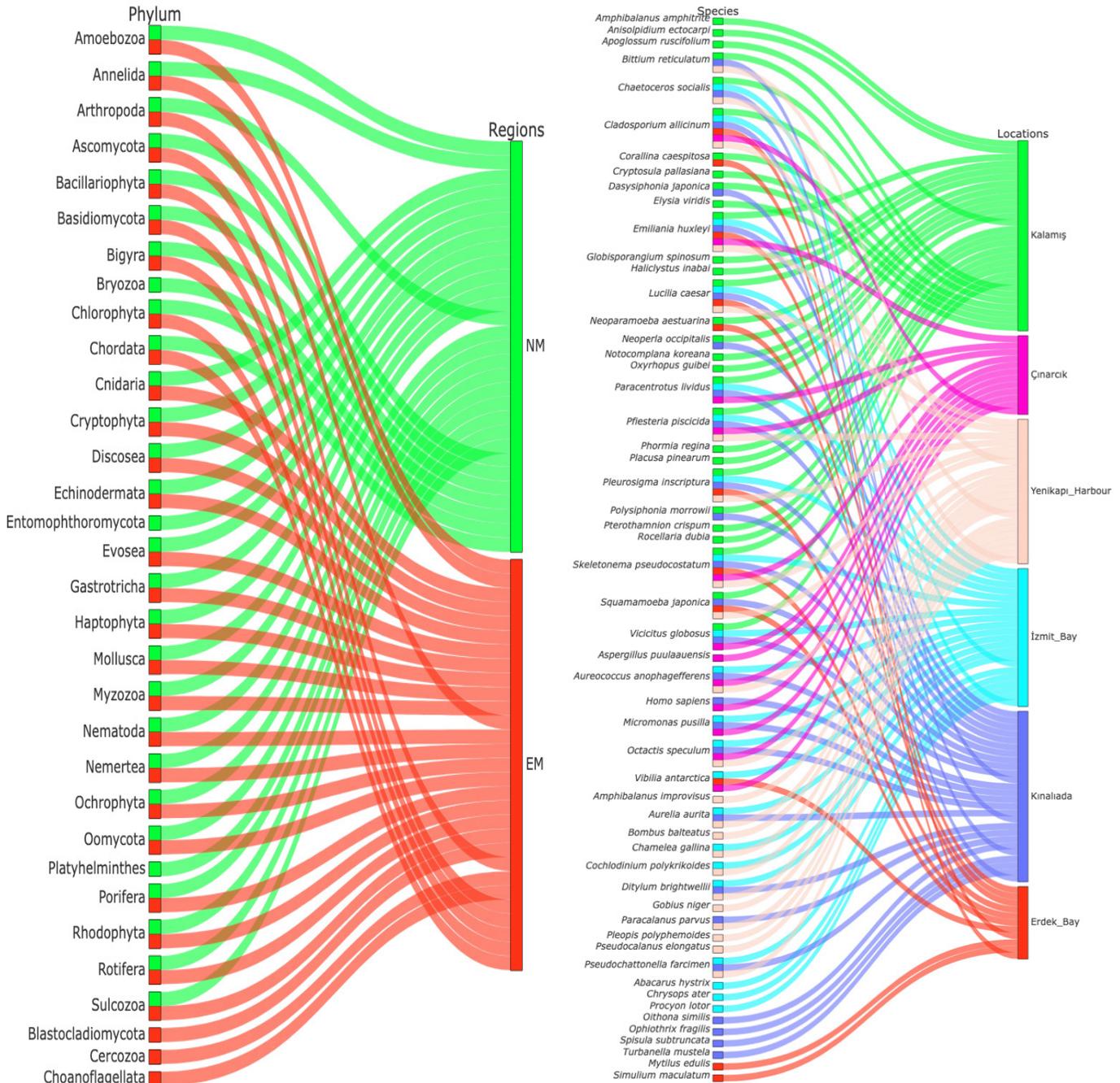


Figure 12. An examination of categories, particularly the taxonomic structure of mucilage samples, exploring the phyla (a) and (b) species compositions.

resented by unicellular eukaryotes (protists), fungi, and other multicellular eukaryotes. We detected a variety of groups, including harmful dinoflagellates such as *Cochlodinium polykrioides* (Gobler, 2008), *Gyrodinium instriatum* (Nagasoe et al., 2006), and *Pfiesteria piscicida* (Burkholder & Glasgow Jr, 1997) that are implicated in the formation of red tides, which have the potential to cause fish death and fishery losses. Within the class Bacillariophyceae, some important groups that could have also contributed to the 2007–2008 mucilage event in the SoM, such as the genera *Cylindrotheca*, *Ditylum* (*Ditylum brightwellii*), and *Skeletonema* (*Skeletonema pseudocostatum*), were also detected.

The coccolithophores *Emiliania huxleyi* and *Dictyocha speculum*, which contribute to carbon and silica cycles, especially in fundamental mineral fluxes within the global ecosystems (Turley, 1991), were observed in our analyses. In the early 2000s, researchers recorded the occurrence of some *D. speculum* species for the first time in the SoM at relatively close stations and at depths with our sampling regime (Deniz, Taş & Koray, 2006), corroborating the findings of the two studies. The dataset has also revealed the presence of filamentous fungal species such as *Cladosporium allicinum* and *Aspergillus puulaauensis* as well as *Globisporangium spinosum* and *Anisopidium ectocarpii* within the realm of Fungi, which release extracellular polymeric compounds similar to those released by bacteria and phytoplankton that serve as a glue in mucilage (Burd et al., 2020).

Some red algal species, including *Apoglossum ruscifolium*, *Coralina caespitosa*, and *Polysiphonia morrowii*, known for having orders that produce polysaccharides (Duarte, et al., 2004; Usov, 2011), were also observed. We also observed the presence of the bryozoan *Cryptosula* that depends on phytoplankton as a major food source, as well as some microspecies such as bivalves (*Chamelea gallina*, *Mytilus edulis*, *Rocellaria dubia*, and *Spisula subtruncata*) as consumers of mesozooplankton (Davenport, Smith, & Packer, 2000), gastropods (*Bittium reticulatum* and *Elysia viridis*), copepods (Calanoids: *Paracalanus parvus*, *Pseudocalanus elongatus*, and Cyclopoid: *Oithona similis*), the cladoceran *Pleopsis polyphemoides*, and the moon jellyfish *Aurelia aurita*. Among these, some taxa, including the abovementioned ones, were the most abundant during the 2018 mucilage event in the area (Okyar et al., 2015), and their (copepods and cladocerans) prevalence and dominance among zooplankton have been recorded in the northeastern SoM (İsinibilir et al., 2008). Chitinous zooplankton, such as copepods and their fecal pellets are recognized as hotspots for microbial activity. Specifically, copepods, such as *Paracalanus parvus*, *Pseudocalanus elongatus*, and *Oithona similis*, which play vital roles in the pelagic food web (Turner, 2004), and their documented mucilage-consuming habits, were also part of the animal composition. Moreover, the scyphozoan *Aurelia aurita*, a species associated with the major bacterial groups (Kos Kramar et al., 2019) and widely acknowledged as a significant player in marine ecosystems (Weiland-Bräuer et al., 2015), was observed. Studies (Brodeur et al., 2002; Sommer & Lengfellner, 2008) have demonstrated that *Aurelia aurita* can significantly influence ecosystem dynamics by affecting planktonic food web structure. By consuming ichthyoplankton, jellyfish exhibit predatory behavior and can potentially compete with fish (Purcell, 2005).

The phylum Mollusca has been well represented in animals that comprised certain groups such as Gastropoda (*Bittium reticulatum*) and Bivalvia, which are commonly documented in marine snow (Shanks & Walters, 1997). Specifically, the gastropod species *B. reticulatum* (12% reads of Animalia) was also previously demonstrated to be associated with eutrophication (Gacia et al., 2009) and biofilm formation (Castejón-Silvo & Terrados, 2017) (D'alelio et al., 2011). Moreover, some gastropod species are affected by the toxicity of algal species (Díaz, 2006) that also comprise nutrient sources (e.g., *Apoglossum ruscifolium* and *Dasysiphonia japonica*); these two algal species were also detected in our dataset.

CONCLUSION

The application of eDNA metabarcoding tools may provide a snapshot of the community composition of the factors that trigger mucilage formation events. Continuous attempts to gather data at a larger and more continuous timescale are crucial to improve our understanding of this phenomenon.

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Author contributions: Mİ, OD, and RB were responsible for devising the study's design. OD conducted the environmental DNA (eDNA) sampling, while ADÖ and OD managed the laboratory workflow and wet-lab process, respectively. OD performed bioinformatic analyses, visualization, and writing, whereas RB supervised the dry laboratory process. Mİ, OD, and RB composed the initial draft of the manuscript. Mİ supervised the project, administered it, and secured funding, while all authors contributed to the manuscript and approved the final version for publication.

Ethical approval: No animal testing was performed during this study.

Conflict of interest: The authors have no conflicts of interest to declare.

Sampling and field studies: All necessary permits for sampling has been obtained by the authors from the competent authorities.

Data availability: The raw sequencing data stored at TRUBA resources, as mentioned in the acknowledgments, are available upon request.

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A Review on the Impact of Thermal Stress on Fish Biochemistry

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ABSTRACT

Fish are an important resource for humans, providing food, economic support, and ecological services. However, rising global temperatures and subsequent increases in their habitat water temperature, pose a significant challenge. We conducted a systematic review to understand the biochemical responses of thermal stress on fish. Stress can be acute (rapid exposure for a short duration) or chronic (repetitive long-term exposure). Stress responses occur at neurotransmitter and hormonal levels, progressing to peripheral and organism-wide effects. Prolonged stress leads to reduced growth, reproductive impairments, heightened infection susceptibility, and mortality. Elevated temperatures serve as abiotic stressors, triggering biotic stress responses. Fish employ strategies to cope with thermal stress, including altering gene expression, metabolite profiles, cellular signaling, and enzyme activity. Cumulative effects of thermal stress induce oxidative stress, causing cell death, organ failure, and mortality. Stressors increase the energy demand, prompting changes in hormonal, enzymatic, and biomolecular responses. Cortisol alters gene expression, stimulating glucose synthesis (gluconeogenesis). Other hormones (thyroid hormones, epinephrine, norepinephrine, insulin, glucagon) also play roles in the thermal stress response. Enzymes involved in metabolic pathways have optimal temperature and pH ranges altered by thermal stress. Heat shock proteins and warm acclimation proteins act as protective mechanisms by preserving the structural integrity of proteins, which is crucial for maintaining proper functionality and cellular responses. Further research is needed to expand on these molecular mechanisms to evaluate proper mitigation strategies.

Keywords: Fish stress, Global warming, Thermal stress, Biochemical responses, Molecular mechanism

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INTRODUCTION

Importance of fishes

Fish are very important aquatic animals to humans as they provide different health benefits, economic support, and ecological services (FAO, 2016). As per the www.fishbase.de website, there are a total of 33,230 fish species in the world (Fishbase, 2020). Direct or indirect beneficial impacts of fish on the human population are numerous and have prevailed from ancient ages; only a few of these are discussed here. Fishes are important nutritional food for humans as well as for other animals. Fish are a low-cost protein source for humans and nearly

3.1 billion humans depend on fish to the tune of at least 20% of their total animal protein intake. Fish are the source of different essential amino acids such as lysine, valine, and functional amino acids, such as leucine and arginine (Ganguly et al., 2018). Fish are also a source of important vitamins, minerals, and other nutrients such as A, D, E, K, B12, sodium, potassium, magnesium, zinc, calcium, iron, copper, iodine, selenium, and fish oils (http://www.fao.org/fileadmin/user_upload/newsroom/docs/BlueGrowthNutritionRev2.pdf). Fish oils are rich in important PUFAs, such as linoleic acid, α -linolenic acid, arachidonic acid, EPA and DHA. Fish oils have different physiological benefits in several patho-



logical conditions such as heart problems, atherosclerosis, rheumatoid arthritis, bipolar disorder, osteoporosis, asthma, etc. (Mohanty et al., 2019). Ornamental fish also are used in pet therapy to reduce mental depression (Gardiánová & Hejrová, 2015). Fish spas are also very popular nowadays (Riyaz & Arakkal, 2011). Fish are also used in the production of feed for other animals including other fish (Tacon & Metian, 2008).

Increment in water temperature

'Climate change' is a nightmare for the whole world and one of the most challenging hazards to be faced on the Earth in the coming future. Climate change results in a rise of the global surface temperature, known as 'Global warming'. The mean temperature of Earth's surface so far has risen by 1.4 °F over the last century and is expected to rise more (2 to 11.5 °F) over the coming hundred years (<http://www.epa.gov/climatechange/basics/>). Global warming is not only an increase in the temperature but also an alteration of different parameters of water bodies (Ficke et al., 2007, Brander, 2010). The surface temperature of the Earth has already increased (nearly +0.93 °C in the past 150 years) and is predicted to increase a few degrees more (1–4 °C) up to the end of this century (IPCC, 2007). Other than the increment in the Earth's surface temperature, experts have predicted some regional variations in temperatures (IPCC editor, 2012). Thus, impacts will vary in different parts of the globe. More events of extreme, abrupt, and frequent changes in temperature are documented in past decades (1991–2000) compared to earlier decades (1971–1990) (Dash & Mamgain, 2011). Different researchers (Diffenbaugh et al., 2007; Ray et al., 2012) also have found extreme events earlier. A heatwave was observed in Ahmedabad, India when the temperature rose to 46.8 °C during May 2010 (Azhar et al., 2014). Increments in the air temperature also led to a rise in the water temperature. Jurgelėnaitė and Jakimavičius (2014) have established a good positive correlation between air temperature and water temperature. The IPCC (2013) in their fifth assessment report estimated ocean warming by 0.09 to 0.13 °C per decade over the past 40 years. There will be an increment in the temperature of water bodies that will alter the habitat of different aquatic animals. Fishes are temperature-sensitive animals (poikilotherms); therefore, increments in the habitat temperature in the coming future beyond the adaptation capacity may exert abiotic stress among them.

STRESS BIOLOGY

General stress in animals

Cannon (1929) was the pioneer researcher who proposed the concept of stress as an "emotionally stimulation situation". Later, Selye (1936) explained the stress in the biological point of view. He demonstrated stress as a "non-specific pathological" response to different "noxious agents". However, several researchers encountered the concept of "the non-specific nature" of stress and proposed it is as highly specific (Mason, 1971; Pacak & Palkovitis, 2001). Stress is the discomfort state of the body under a threat where coordinated responses of physiological, biochemical, and behavioral processes cumulatively help the organism to overcome the situation (Chrousos & Gold, 1992). The initial response is to adapt to the situation by changing different biological processes of the body to a new normal situation in response

to external as well as internal stimuli, which are cumulatively known as "allostatic load" (Schreck, 2001). The term "allostatic" denotes the body's capacity to attain stability amid change, with "load" signifying the toll or expense incurred by the body in adapting to stress (McEwen, 2013). Conceptually, stress can be defined as an alteration of physiological, biochemical (in the lower and higher organism), and mental (only in higher organisms) states due to different external (environmental factors), internal (body imbalance), or emotional stimuli that in the long-run effect performance and create physiological disorders, which every organism tries to avoid although it is nearly impossible.

Stress responses

Under stress, fish initially respond at the neurotransmitter and hormonal level (primary response) then at the peripheral level (secondary response) (Mazeaud et al., 1977), and finally to the whole organism or population level (tertiary responses) (Wodemeyer & McLeay, 1981). The primary responses are the "alarming stages" and two hormonal axes are involved in any kind of stress response in fish. The sympathetic-chromaffin (SC) axis and the hypothalamic-pituitary-interrenal (HPI) axis regulate oxygen uptake, transfer, and energy metabolisms (Wendelaar, 1997, Roychowdhury et al., 2020b). The secondary responses are the stages of "resistance". During that stage, fish try to engage all the available mechanisms to cope with the situation. The tertiary response of an organism is the "stage of exhaustion" where fish are unable to fight against the stress, reflected in their performance, and finally may cause even death (Roychowdhury et al., 2020a, Roychowdhury et al., 2020b). Therefore, the primary responses occur at a systemic level whereas secondary responses occur at the peripheral level, and the tertiary response affects the whole organism (Wodemeyer & McLeay, 1981). Once a fish receives stress stimuli, its sensory neurons in the brain activate the hypothalamus to release different hormones and neurotransmitters. Releasing hormones (RHs) such as CRH and TRH are the most important. These RHs act on the pituitary to release different hormones such as ACTH and α -MSH. ACTH further helps to release cortisol, an important hormone under stress. Cortisol is produced from an internal gland whereas catecholamines are produced in a chromaffin gland and interestingly both of them are found in the kidneys of fish (Milano et al., 1997). These hormones further alter the biochemistry, immunology, and physiology of fish under stress and induce the secondary responses (Weyts et al., 1999). These effects, in the long term, are reflected in fish performance as a reduction in growth, reproduction, susceptibility to infection, and ultimately mortality (Schreck et al., 2001).

Acute stress and chronic stress

Stress responses in fish depend on the types of stressors, the duration of exposure, the rate of introduction to stressors, and the amount of the stress (Barandica & Tort, 2008; Wedemeyer, 1997). Stresses are of two types, acute and chronic, based on the duration of exposure time. Acute stress involves rapid exposure for a short duration of exposure time (hours). Chronic stress is due to constant or repetitive exposure to the stressor for long periods (Barandica & Tort, 2008).

Acute stress is mainly the "fight or flight" response and is initiated mostly in the primary response. Acute stress in Black Sea trout

(*Salmo trutta labrax*) due to thermal exposure may involve an alteration in blood glucose, cortisol, total protein, serum ion concentrations, and lysozyme activity (Dengiz Balta et al., 2017). Acute stress is sometimes helpful for an organism. However, chronic stress can be viewed as repetitive exposure to stressors and the central nervous system (CNS) does not get sufficient time to recover (Sapolsky, 1996). It generally causes secondary and even tertiary responses (health issues). Under any kind of chronic stress, the body changes to 'a new normal' state. Chronic stress impacts nearly all physiological processes, and experimental conditions have been devised to induce such stress, revealing varying durations in different fish species. For instance, it necessitated 15 days in zebrafish (Chakravarty et al., 2013) and Nile tilapia (Volpato & Barreto, 2001), 28 days in rainbow trout (Moltesen et al., 2016), and 8 days in zebrafish (Golla et al., 2020). Intriguingly, chronic stress elicits a more intricate set of responses compared to acute stress. While acute stress predominantly triggers primary and secondary responses, chronic stress encompasses all three types of stress responses. This complexity underscores the multifaceted nature of physiological adaptations in animals under prolonged stress conditions (McEwen & Gianaros, 2011).

THERMAL STRESS IN FISH

In the earlier section, it was discussed that fishes are temperature-sensitive animals, therefore, an increment in the habitat temperature beyond the thermal tolerance capacity causes fish to feel abiotic stress. Thermal tolerance is the range of temperatures encompassing minimum to maximum levels where fish can survive and perform all their physiological functions. The upper-temperature tolerance limit (UTTL) of fish is the highest temperature up to which fish can survive (Daniel et al., 2008, Roychowdhury et al., 2019). The UTTL of fish depends on geographical location (Sorte et al., 2011), previous thermal history (Beitinger et al., 2000), transgenerational acclimatization (Donelson et al., 2012), the type of feed consumed (Kumar et al., 2014), and the rate of temperature increments (Camilo & Maria, 2006). The most commonly employed tools for studying the UTTL in fish are the lethal temperature maximum (LT_{max}) and critical temperature maximum (CT_{max}) (Beitinger et al., 2000). Thermal plasticity in fish refers to the capacity of fish species to adjust their physiological and behavioral traits in response to temperature changes (Comte & Olden, 2017). Fish exhibit thermal plasticity through alterations in metabolism, enzyme activity, gill function remodeling, changes in spawning time, and behavioral adaptations (Angiulli et al., 2020; Das et al., 2012; Farrell, 2009; Roychowdhury et al., 2020b). Parameters like warming temperature (WT) and thermal safety margin (TSM) are crucial for studying thermal plasticity in ectothermic animals (Becker & Genoway, 1979; Madeira et al., 2007). In fish physiology, aerobic space refers to the volume of oxygen available for aerobic metabolism, crucial for energy production (Jobling, 1995). Factors influencing this space include dissolved oxygen levels, gill efficiency, and metabolic rate (Farrell, 2009). Elevated water temperature often results in decreased dissolved oxygen (DO), leading to hypoxic conditions. Understanding aerobic space helps assess fish adaptation to environmental conditions and stressors (Eliason et al., 2013). The Tem-

perature Coefficient (Q₁₀) measures the temperature sensitivity of physiological or biochemical processes, indicating the rate of change with a 10-degree Celsius temperature increase. In the context of oxygen consumption in fish, the Q₁₀ value describes how oxygen consumption rates vary with temperature changes. Researchers use Q₁₀ values to understand thermal dependence in fish and other ectothermic organisms, providing insights into the temperature's influence on metabolic processes and aiding predictions of climate change impacts on fish physiology and ecology. Variations are observed among fish species based on their adaptations to specific environmental conditions (Claireaux & Lefrançois, 2007; Farrell, 2009; Pörtner & Farrell, 2008).

'Fight or flight' is the initial strategy for most organisms in a hostile environment. Fish try to move toward the more preferable environment but if they fail to do so, they will feel stress (Kovach et al., 2012). A warmer temperature is abiotic stress that causes different biotic stress in organisms (Nakano et al., 2014). However, data on the impacts of thermal stress on different biochemical pathways are limited in carp fish. Therefore, previous literature on thermal stresses on fish and other organisms needs to be reviewed to understand the thermal stress biology of an organism. Besides, this information can help to design proper methodology and selection of different biomolecules expected to be altered under thermal stress in fish. The process of adaption and stress are difficult to distinguish at the initial stages as fish try to fight the stress by altering their biochemical pathways in higher temperatures. Patterns of the expression of different biomolecules are changed initially due to adaptation and later on due to stress (Bijlsma & Loeschcke, 2005). The consequence of an event depends on the extent of stress, the genetic makeup of the organisms, and previous history of exposure (Beitinger et al., 2000). A higher temperature is known to increase the kinetic properties of molecules causing a higher diffusion rate for micro-molecules (Bag et al., 2014) and denaturation of macromolecules (Wu, 1995). Maintenance of the native three-dimensional structure of a protein is very much essential for proper functioning, and interaction with other biomolecules involved in cellular response, and any randomness beyond certain limits may lead to cellular death (Nakamura & Lipton, 2009). Fish try to cope with the situation by changing gene expression patterns and are reflected in alteration in metabolites (biomolecules), cellular signaling systems (hormones and others), and enzyme activity (Lancaster et al., 2016). The stress is tolerable to a certain extent after the performance of the species drops (La & Cooke, 2011; Pörtner & Knust, 2007). Due to the cumulative impacts of the different effects of thermal stress, oxidative stress begins to occur and results in cellular death, organ failure and ultimately death of the organism (Figure 1).

BIOCHEMICAL MECHANISM

Any kind of stressor increases the requirement of physiological energy that can be fulfilled by alteration of hormonal, enzymes and bimolecular responses. In general, glucose, triglycerides and protein act as energy molecules depending upon the nature and quantity of energy needed (Mergenthaler et al., 2013). Glucose serves as an immediate energy source, while triglycerides provide major energy, and protein provides energy during hun-

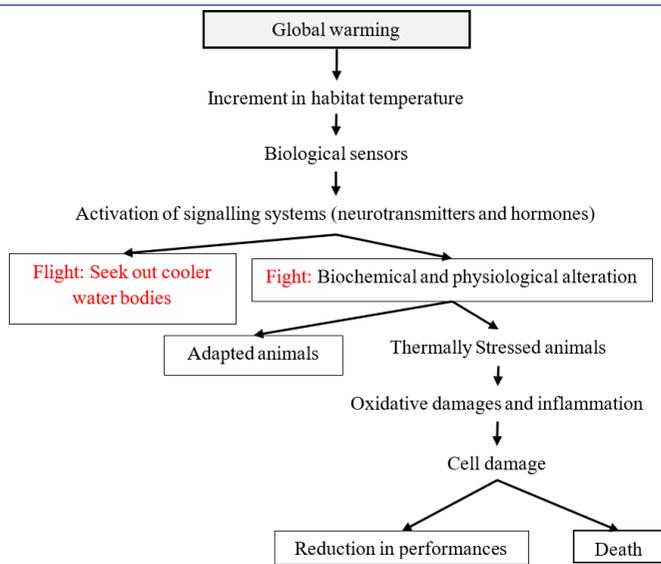


Figure 1. Conceptual representation of the consequences of thermal stress in fish.

ger or in some physiological disorders and stress. In stressful situations, fishes try to increase blood glucose to provide more energy to vital organs including the brain by changing the metabolism of biomolecules under the control of hormones (Mergenthaler et al., 2013). The steroid hormone, cortisol alters the pattern of gene expression (transcription level) and increases the expression of the different enzymes required for the synthesis of new glucose molecules (gluconeogenesis) (Babitha & Peter, 2010; Vijayan et al., 1997) and breakdown of the stored glycogen (glycogenolysis) (Vijayan et al., 1993 and 1996) during stress to increase blood glucose concentration in serum. Cortisol converts the non-essential proteins of muscle to amino acids and transports them through the blood to the liver for gluconeogenesis (Vijayan et al., 1993; Vijayan & Moon, 1994; Freeman & Ilder, 1973). Cortisol is also documented to mobilize stored lipids in teleost species (Dave et al., 1979). Thyroid hormones, T3 and T4 are produced in the thyroid gland and normally maintain a positive nitrogen balance, but T3 inhibits protein synthesis in higher concentrations (Murray et al., 1996). Like cortisol, thyroid hormones are also known to act at the transcriptional level and modify its function or 'fine-tune' the actions of cortisol and adrenaline in a stressful situation (Peter, 2011). Epinephrine and norepinephrine, released from chromaffin cells, are recognized by their immediate elevation of plasma glucose levels (Arends et al., 1999; Ruane et al., 2001). Their roles encompass enhancing glucagon secretion, facilitating glycogen breakdown (glycogenolysis), initiating new glucose synthesis (gluconeogenesis), mobilizing fatty acids (beta oxidation), and concurrently suppressing insulin production, glycogen synthesis (glycogenesis), and glucose breakdown (glycolysis) (Nelson & Cox, 2001). Their primary targets are muscle, adipose tissue, and liver, and increase the supply of oxygen to tissue by increasing heart rate, blood pressure, and respiratory passage (Nelson & Cox, 2001). Cortisol plays a role mostly in chronic stress, while epinephrine and norepinephrine act as emergency hormones and are involved in immediate responses. Insulin and glucagon maintain glucose homeostasis in blood in

normal conditions but their direct involvement in heat stress is limited. Insulin increases the utilization of glucose by increasing glucose transportation in cells and increasing the rate of glycolysis in extrahepatic tissues and glycogenesis in hepatic tissue but glucagon plays the counter role of insulin by producing glucose through gluconeogenesis and glycogenolysis thus increasing blood glucose (Nelson & Cox, 2001).

Blood glucose is used to monitor general stress for different species. Energy-rich molecules like ATP, GTP, phosphocreatine, succinate, and ketone bodies are also known to play an important role during thermal stress and can be used as possible markers for thermal stress (Dijk et al., 1999). Metabolism of biomolecules is under hormonal control necessitating monitoring of the hormones cortisol, T3, T4, epinephrine, norepinephrine, insulin, and glucagon during stress.

Every enzyme has an optimum temperature and pH and thermal stress is known to change these parameters for cold-blooded fish. Enzymes involved in the metabolism of energy-rich biomolecules alter their activity by changing their concentration (expression) or affinity (modification) towards the substrate under a stressed condition. Enzymes catalyzing the irreversible steps of carbohydrate metabolism (hexokinase, phosphofructokinase1, pyruvate kinase, citrate synthase, glucose 6 phosphatase, fructose-1,6 bis-phosphatase, etc.), protein metabolism (glutamate-pyruvate transaminase and glutamate-oxaloacetate transaminase), lipid metabolism (lipase), and creatinine metabolism (creatinine kinase) are important from this perspective (Dhanasiri et al., 2013, Abbaraju & Rees, 2012; Ton et al., 2003).

Thermodynamics explains life as the maintenance of organized cellular structures through energy generated by oxidizing energy-rich nutrients by enzyme(s) that act at an optimum temperature under the control of hormones. Temperature changes directly affect these activities. Proteins are the mediators of diverse types of cellular functions (Alberts et al., 2002) including enzymatic action to scale formation, body growth to reproduction, and digestion to excretion. The function of a protein depends on its native structure that gets denatured due to the rise in temperature causing more vibration in molecules and related disturbance. Heat shock proteins (Hsps), which are highly conserved, are the molecular chaperones required for the proper folding of the protein and are also involved in protein transport to cellular compartments (Alberts et al., 2002; Shi & Thomas, 1992). At higher temperatures Hsps refold the denatured protein to its original native structure and the heat shock factor (Hsf) induces the expression of various Hsps in the process (Wu, 1995). There are various types of Hsps within the cell such as Hsp10, HspB group (Hsp27, hspB1, hspB6), HspA group (Hsp70, Hsp71, Hsp72, Hsp78), HspC group (Hsp90, hsp94), Hsp40, Hsp60, Hsp104, and Hsp110 (Antonova et al., 2007; Benjamin & McMillan, 1998; Li & Srivastava, 2004; McLemore et al., 2005; Salinthon et al., 2008; Schlesinger, 1990). Hsp30, Hsp27, and Hsp47 are identified in goldfish (Wang et al., 2007) while Hsp78 and Hsp100 are found in *Poeciliopsis lucida* (Norris et al., 1995). Warm acclimation protein (wap) is another candidate that plays an important role during heat stress (Kikuchi et al., 1997) (Figure 2).

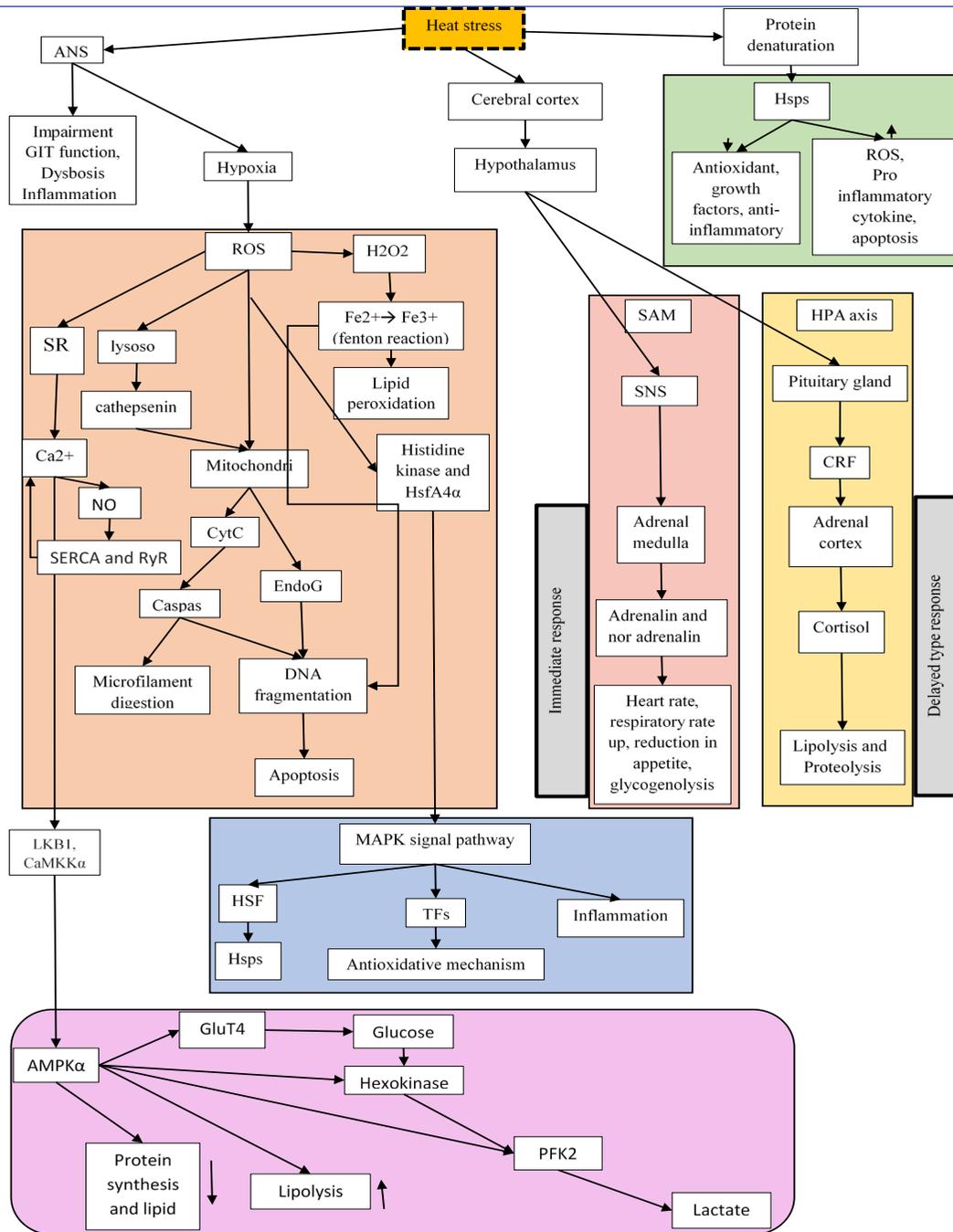


Figure 2. Impact of thermal stress at the molecular biology level (Antoun et al., 2017; Cui et al., 2016; Kourtis and Tavernarakis, 2011; Kültz, 2015; Xing et al., 2019).

Numerous experiments have been conducted to study the impacts of heat stress on fish and a few of them are cited here: rahu (Akhtar et al., 2013; Kumar et al., 2015; Mohapatra et al., 2014), common carp (Ouellet, et al., 2013), rainbow trout (Recsetar et al., 2012), blenny species (Camilo and Maria, 2006), salmon (Dengiz Balta et al., 2017; Nakano et al., 2014), *Channa punctatus* (Kaur et al., 2005), pacific sardine (Kaur et al., 2005), and *Danio dangila* and *Brachydanio rerio* (Majhi & Das, 2013). The selected temperatures for treatments were mostly sub-lethal with no fish death (Das et al., 2002; Das et al., 2005; Nakano et al., 2014).

These experiments were mainly focused on the recovery responses (Kumar et al., 2015), adaptation strategies (Das, 2002), gene expression pattern (Ouelle et al., 2013), oxidation-reduction state (Nakano et al., 2014), thermal tolerance, growth and oxygen consumption (Das et al., 2005), and search for bio-markers (Purohit et al., 2014). Different experimental conditions were used for warmer water exposure to induce heat stress in fish. The three main determinants to introduce thermal stress to fish are the amount of temperature, duration of exposure, and rates of temperature increment, and all of them varied in the experi-

ments (Akhtar et al., 2013; Das et al., 2005; Dengiz Balta et al., 2017; Kaur et al., 2005; Kumar et al., 2015; Majhi & Das, 2013; Mohapatra et al., 2014; Nakano et al., 2014; Recsetar et al., 2012). Fish may feel both acute stress and chronic stress based on events that will occur in future (IPCC, 2014). Different researchers have studied the impacts of both acute and chronic thermal stress among other animals including fish (Hernández-López et al., 2018) but data is very limited for IMC. Thermal stress is expected to alter the different physiological and biochemical processes of the body and that may be reflected in different pathways and performance of fish (Figure 3).

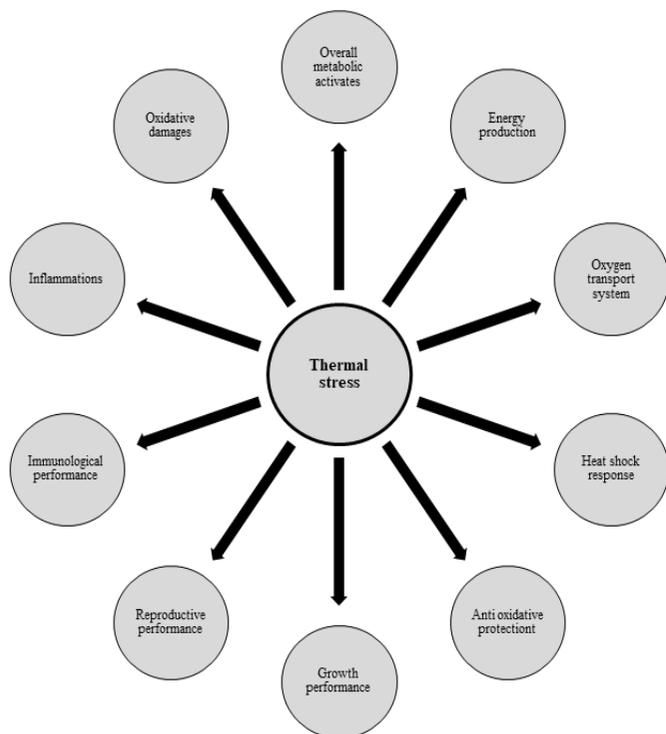


Figure 3. Abiotic thermal stress affects different physiological pathways (Aftabuddin et al., 2017; Aftabuddin and Roychowdhury, 2019).

DISCUSSION

The review revealed that thermal stress has substantial effects on fish, leading to reduced growth, reproductive impairments, increased susceptibility to infections, and even mortality. These consequences can have profound ecological and economic implications, considering the vital role of fish in providing food, supporting economies, and maintaining ecosystem balance. The study highlights the distinction between acute and chronic stressors, with acute stress resulting from short-term rapid exposure and chronic stress arising from long-term repetitive exposure. Understanding this differentiation is crucial for developing effective strategies to mitigate the impacts of thermal stress on fish populations. The elevated water temperatures associated with global warming act as abiotic stressors, triggering various biotic stress re-

sponses in fish. It was observed that fish employ a range of mechanisms to cope with thermal stress, including alterations in gene expression patterns, metabolite profiles, cellular signaling pathways, and enzyme activity. These adaptive responses enable fish to maintain their physiological balance and attempt to counteract the detrimental effects of thermal stress. However, the cumulative impact of thermal stress can induce oxidative stress, leading to cellular death, organ failure, and ultimately mortality. This highlights the urgent need to address the challenges posed by rising temperatures and the associated thermal stress on fish populations. The study emphasizes the role of hormonal, enzymatic, and biomolecular responses in the fish's attempt to meet the increased energy demands caused by stressors. Hormones such as cortisol play a significant role in altering gene expression and stimulating glucose synthesis through gluconeogenesis. Other hormones, including thyroid hormones, epinephrine, norepinephrine, insulin, and glucagon, are also involved in the fish's response to thermal stress. Enzymes associated with metabolic pathways in fish have optimal temperature and pH ranges that can be altered by thermal stress. Maintaining the proper structure and functionality of proteins is crucial for cellular responses and overall fish health. Heat shock proteins and warm acclimation proteins act as protective mechanisms, helping to preserve protein integrity and mitigate the adverse effects of thermal stress.

The findings of this systematic review emphasize the importance of further research to enhance our understanding of the molecular mechanisms underlying these responses. Such knowledge can guide the development of targeted conservation and management strategies to safeguard fish populations in the face of a rapidly changing environment.

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Declaration of Generative Ai in Scientific Writing: During the preparation of this work the author(s) used [https://chat.openai.com/] in order to improve English language such as checking grammar, spelling and for finding some references. After using this tool/service, the author(s) reviewed and edited the content as needed and take(s) full responsibility for the content of the publication.

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The Death of the Beluga Sturgeon (*Huso huso*) Rediscovered in the Küçükçekmece Lagoon: Let's Focus on Causes

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ABSTRACT

Sturgeon, a key species in fisheries and aquaculture due to their high economic and ecological value, face numerous challenges in their natural habitats. The spawning grounds in rivers, essential for sturgeon reproduction, are often compromised by the blockage of migration routes caused by various obstacles. Lagoons, providing favorable conditions for fish growth, are also hunting grounds for predatory fish species. The Küçükçekmece Lagoon, located within İstanbul's borders in Türkiye, stands as one of the most significant lagoons in the region, especially noted for its position on the bird migration route. However, it has been suffering from unplanned urbanization, leading to a notable decline in water quality due to improper urban planning and industrial activities over the years. This decline is aggravated by waste drainage into the lagoon's wetland area. Notably, the last sturgeon sighting in the Küçükçekmece Lagoon was in 1986. After approximately 35 years, the discovery of two deceased beluga sturgeon (*Huso huso*) in the lagoon, which connects to the Marmara Sea in Türkiye, marked a significant event. This study aims to investigate the causes behind the death of these beluga sturgeons and also proposes potential solutions for enhancing the ecosystem's vitality of the lagoon. These solutions include addressing waste drainage, ensuring shoreline compliance, improving the sediment structure laden with organic matter, and artificially introducing fresh water. Such measures are crucial for the restoration and maintenance of the ecological balance in the Küçükçekmece Lagoon, which is vital for the survival of significant species like the sturgeon.

Keywords: Acipenseridae, brackish water, eutrophication fish migration, water pollution

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INTRODUCTION

Over recent decades, there has been a continuous and significant decrease in natural sturgeon populations (Pikitch, et al., 2005; Zhang, et al., 2019). This trend, combined with the ongoing high demand for caviar on the market, has paved the way for the growth of sturgeon farming, with a primary focus on caviar production, as reported by Bronzi & Rosenthal (2014). These fish are typically recognized for their distinctive appearance, massive size, and their renowned product, caviar. They are an ancient species that inhabit only the northern hemisphere (Ustaoglu & Okumus, 2004). Currently, all Eurasian sturgeon species are facing a signif-

icant threat, while the situation is somewhat better in America (Williot et al., 2011).

Memiş (2014) reported that the status of all sturgeon species in Türkiye has undergone a complete transformation in the past 40 years, and many are now on the brink of extinction. This change is believed to be due to various factors, including human-made modifications to rivers, such as the construction of embankments, dams, regulators, and hydroelectric power plants, which have prevented upstream migration to spawning sites (Ustaoglu Tırıl & Memiş, 2018). Pollution from industries, agriculture, and sewage, as well as overfishing, have also been identified as significant contributors to



the drastic decline in sturgeon populations, both along the Black Sea coast and in Türkiye's three largest rivers (Kızılırmak, Yeşilirmak, Sakarya) (Ustaoğlu Tırlı & Memiş, 2018; Memiş et al., 2019).

Due to the significant decline in sturgeon populations, this species has been under protection since the 1980s. Türkiye signed the CITES agreement on 22 December 1996 and the catching and sale of sturgeon was banned by the Turkish authorities on April 1, 1998 (Memiş et al., 2020). In an effort to preserve natural sturgeon populations, sturgeon are being bred through aquaculture methods and released into the wild, contributing to the conservation of these endangered species (Akbulut et al., 2011).

Wetlands are the most productive areas on earth in terms of biological production and are also very rich in biodiversity (Tırlı, 2004). Lagoons have a special place among wetlands due to their connection to the sea. They are formed by the complete or partial closure of river mouths that flow into the sea with a coastal spit and sometimes by the accumulation of sea sand transported by currents in front of bays or gulfs (Joyeux & Ward, 1998; Karataş, 2005). They cover 13% of the continental coastal zone and are of great importance for biochemical processes and high productivity. Also, because of differences in salinity values, they have special ecological conditions that contribute to the richness of lagoon areas in terms of biodiversity (Gilabert, 2001; Çevik et al., 2008; Acarlı et al., 2009). They provide suitable conditions for fish to spawn and grow, as well as being preferred places for predatory fish species to feed. With these features, lagoons are also attractive centers for waterbirds and important feeding and offspring care areas.

Küçükçekmece Lagoon is one of the most important lagoons in Türkiye. Because of urbanization and industrialization, deterioration in water quality affects the ecosystem significantly (Albay et al., 2005; Köker et al., 2021). In the 1980s, various species of commercial fish were detected in the lagoon, but the fish population decreased in the early 2000s (Topçuoğlu et al., 1999). In the study conducted by Meriç (1986) on Küçükçekmece Lagoon, it was reported that 31 fish species inhabit the lagoon. These were *Acipenser gueldenstaedtii*, *Alburnus istanbulensis*, *Alosa tanaica*, *Anguilla anguilla*, *Aphanius fasciatus*, *Atherina boyeri*, *Belone belone*, *Clupeonella cultriventris*, *Chelon ramada*, *C. aurata*, *C. saliens*, *Engraulis encrasicolus*, *Esox lucius*, *Gasterosteus aculeatus*, *Huso huso*, *Knipowitschia caucasica*, *Mugil cephalus*, *Mullus surmuletus*, *Neogobius melanostomus*, *Nerophis ophidion*, *Petroleuciscus borysthenicus*, *Platichthys flesus*, *Proterorhinus semilunaris*, *Sander lucioperca*, *Sardina pilchardus*, *Scardinius erythrophthalmus*, *Pomatomus saltatrix*, *Sprattus sprattus*, *Syngnathus abaster*, *Trachurus mediterraneus* and *Vimba vimba* (Meriç, 1986; Deveciyan, 2006). Because of water pollution, many fish species have moved away from this lagoon and its fish population and diversity have decreased in recent decades (Özuluğ & Saç, 2019).

Beluga sturgeon (*Huso huso* Linnaeus, 1758) has not been recorded in Küçükçekmece Lagoon since 1986 (Meriç, 1986). Recently, two beluga sturgeon were found in the lagoon after 35 years. While the original goal of this study was to investigate their existence and causes of death, it has become evident that a broader approach is required. Given the deteriorated condi-

tion of the fish, which precludes detailed biochemical or histopathological analysis, the study now emphasizes understanding the lagoon's ecosystem structure and identifying ways to improve it. This approach is crucial, considering the fragile state of species like the beluga sturgeon and the need for a comprehensive strategy to enhance the resilience and health of the entire ecosystem.

MATERIALS AND METHODS

Study area and environmental variables

Küçükçekmece Lagoon, located in the west of Istanbul and connected to the Marmara Sea by a narrow channel, is 18 m deep and has a surface area of 15.22 km² (Figure 1). Eşkinöz, Nakkaşdere and Sazlıdere creeks flow into the lagoon and these creeks show seasonal flow (Albay et al., 2005, Topçuoğlu et al., 1999). Sazlıdere Dam, which is a drinking water supply, is located on Sazlıdere, the most important freshwater source of the lagoon. Due to this dam, the freshwater inflow into the lagoon has decreased.

During our field studies, the discovery of two nearly decomposed *H. huso* individuals on July 1, 2020, in the Küçükçekmece Lagoon was initially reported by local residents who frequently visit the lagoon's shores for recreational and fishing activities. This encounter was later confirmed by public officials who monitor the lagoon. The discovery was facilitated by wave movements bringing the fish to the shore (Figure 2). One of the individuals was found to be completely deteriorated with body integrity lost.

The length of the other fish stranded on the shore was measured on the field with a 1 mm calibrated meter, and a sample was taken from the pectoral fin rays to determine its age. The fin ray was prepared for reading by being placed in bleach for 2 min, and all excess tissue and skin were removed using a scalpel and forceps (Parr et al., 2018). The fin ray was sectioned using a Metkon low-speed saw (Metkon Microcut, Bursa) equipped with two diamond-tipped blades (Metkon Microcut 150) and a 0.15 mm spacer between the blades (Deval et al., 2018). Three sections were cut serially, beginning at approximately 25% of the total length of the fin ray, starting at the condyle base (Parr et al., 2018). The method for determining the age of the beluga sturgeons involved analyzing the sectioned fin rays, specifically focusing on the number of translucent bands present. Following the approach outlined by Bakhshalizadeh et al. (2017), we identified both opaque and translucent bands as annual growth rings, or annuli, on the fin ray. This identification is based on the principle that these bands represent distinct periods of growth and development in the beluga sturgeon's life cycle. Fin ray sections were examined and photographed under the reflected light using an Olympus SZX16 stereo microscope (with 2 × 10 magnification) (Deval et al., 2018). The fish species identification was made according to Akşıray (1987) and Kottelat & Freyhof (2007). The current scientific names of the species were verified according to Fricke et al. (2022).

For water quality analysis, water samples were taken from the surface waters where the beluga sturgeon were found. The YSI multiparameter (YSI 650 MDS) was used to measure dissolved oxygen, salinity, pH, and temperature *in situ*. Chlorophyll-a (chl-a)

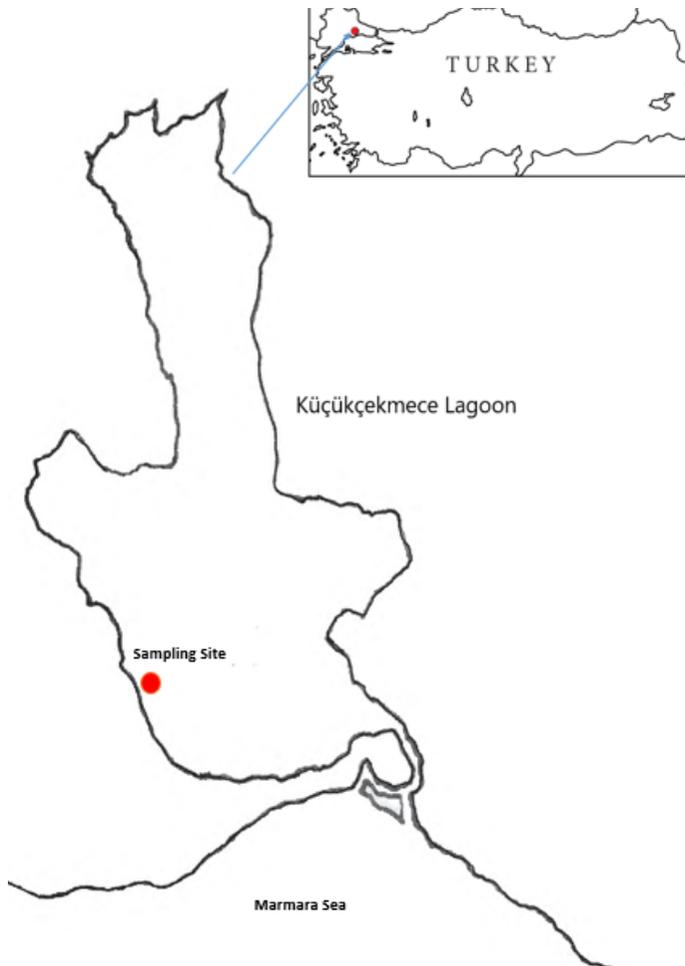


Figure 1. Location of Küçükçekmece Lagoon. (Modified from Köker et al., 2021).

was determined using the method of ISO 10260 (1992). Nutrient concentrations: total inorganic nitrogen (TIN: $\text{NO}_2\text{-N} + \text{NO}_3\text{-N} + \text{NH}_4\text{-N}$) and total phosphorus (TP) were measured according to APHA-AWWA WPCF (2005). The Secchi depth (SD) was measured *in situ*. The trophic state of the Küçükçekmece Lagoon, Carlson Index was used and total phosphorus (TP), chlorophyll-a (Chl-a), and Secchi depth (SD) measurements were used to calculate the index (Carlson, 1977).

RESULTS AND DISCUSSION

The subject of this present study is the beluga sturgeon-*Huso huso*, an anadromous fish species that migrates kilometers upstream to the source of rivers from the seas in order to spawn on deep and gravelly-stony river bottoms (Chebanov & Galich, 2013). It is known that beluga sturgeon entered the Küçükçekmece Lagoon in the past (Meriç, 1986). This species would enter the lagoon not for reproduction but for feeding purposes. The fish species known to inhabit the Küçükçekmece Lagoon are among the preferred food sources in the diet of beluga sturgeon.

The beluga sturgeon (*Huso huso*) can reach approximately 6 m in length and up to 1000 kg in weight, with records indicating lengths of 8 m and weights of 3200 kg. Individuals are typically

found to measure between 120-260 cm and weigh up to 363 kg (Berg, 1948). In the Küçükçekmece Lagoon, the age of the beluga sturgeon found in July 2020 was determined from the pectoral fin ray and it was around 20 years old. The specimen measured 250 cm in total length, indicating that the beluga sturgeon had reached the reproductive stage.

There are numerous reports that various hydraulic obstacles such as hydroelectric power stations, bridge abutments, and culverts built on rivers affect fish migration (DWA, 2005; Shortnose Sturgeon Status Review Team, 2010; Nyqvist et al., 2017; Nielsen and Szabo-Meszaros, 2022). Küçükçekmece Lagoon is a special wetland due to its location in the middle of Istanbul and it is an important habitat for living organisms from bacteria to fish. Since the 2000s, the lagoon has continued to be polluted, and with the construction of the Sazlıdere dam, the ecosystem has faced change and collapse due to the decrease in freshwater input (Köker et al, 2023). It has been reported that algal blooms have occurred in the Küçükçekmece Lagoon since the 1990s, resulting in the death of some fish (Albay et al. 2005; Köker et al. 2021).

Due to the increase in the salinity of the water, cyanobacterial blooms ceased to occur. The calculated $\text{TSI}_{(\text{Chl-a})}$ and $\text{TSI}_{(\text{TP})}$ mean values ($\text{TSI}_{(\text{mean})} = 58.21$) suggest that the Küçükçekmece Lagoon is in a nutrient-rich, eutrophic condition.

The physicochemical parameters measured during the investigation of the death of the beluga sturgeon individuals are shown in Table 1. The measured water temperature value was within the normal range for the summer season. Although it is known that the salinity level varies between 6-9 ppt in the area of Sazlıdere Dam where the freshwater inflow enters (Köker et al., 2021), the salinity level was determined as 13.5 ppt in the sampling area with currents coming from the channel where the lagoon connects to the Marmara Sea. The most important parameter was dissolved oxygen which measured 0.78 mg L^{-1} .

The trophic state of Küçükçekmece Lagoon was determined by following formulas based on total phosphorus (TP) and chlorophyll-a (Chl-a) (Carlson, 1977): $\text{TSI}_{(\text{Chl-a})} = 9.81 \ln(\text{Chl-a}) + 30.6$, and $\text{TSI}_{(\text{TP})} = 14.42 \ln(\text{TP}) + 4.15$. According to Table 1, it was determined that Küçükçekmece Lagoon is eutrophic.

Table 1. Some physical and chemical variables of the Küçükçekmece Lagoon during the sampling.

Environmental Parameters	June 2020
Temperature (°C)	26.35
Dissolved oxygen (mg L^{-1})	0.78
Salinity (ppt)	13.49
pH	8.20
Chl-a ($\mu\text{g L}^{-1}$)	1.11
TIN ($\mu\text{g L}^{-1}$)	5559.2
TP ($\mu\text{g L}^{-1}$)	270.1
$\text{TSI}_{(\text{TP})}$	84.88
$\text{TSI}_{(\text{Chl-a})}$	31.53
$\text{TSI}_{(\text{mean})}$	58.21

In July 2020, two individuals of *H. huso* who had died and whose bodies had washed ashore were found. One individual's tissues had completely decayed and its total length could not be measured. The total length of the other individual, which was in the process of decay, was 250 cm (Figure 2). Based on the overall condition of the fish bodies, it was clear that they had died a few days before our sampling.

The exact age of the *H. huso* individual shown in Figure 2-3, could not be determined due to the deterioration of the fin rays (Figure



Figure 2. Measurement of the length of the *Huso huso* individual.



Figure 3. Investigation of the species from the mouth and barbels.

4). The first ray of the pectoral fin taken from the 250 cm long fish and used for age determination was sectioned at the base. It was estimated to be approximately 20 years old (Figure 4).

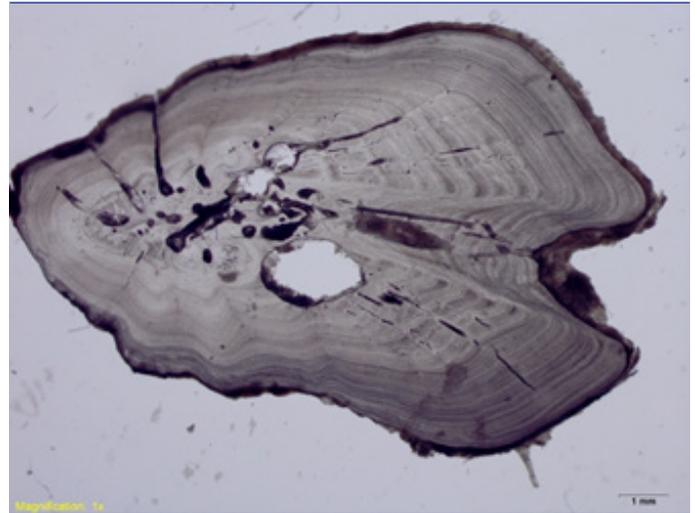


Figure 4. Pectoral fin barb beam section for age determination of beluga sturgeon.

Although the gonad examination of the fish suggested that it was female based on the consistency of the mass of the gonad and its large volume, there are signs similar to a male (M. Chebanov, personal communication, 7 July 2020).

Particularly during the summer season, factors such as increased water temperature and biological decomposition of waste within the lagoon lead to significant decreases in dissolved oxygen levels, even dropping to $0.78 \text{ mg O}_2 \text{ L}^{-1}$ (Table 1). The formation of oxygen-depleted zones throughout the water column, from the surface to the bottom of the lagoon, results in mass mortality events, especially among benthic invertebrates (e.g., crabs) and bottom-dwelling fish species (e.g., Gobiidae, Syngnathidae), as well as pelagic species like *A. boyeri* and *Chelon* spp., which have been observed during field observations. Atlantic sturgeon populations have been found to be declining in the Chesapeake Bay due to increased hypoxia (Secor & Gunderson, 1998; Secor & Niklitschek (2001). Even if the sturgeon has adapted physiologically to the environment, hypoxic conditions impair its respiratory metabolism, foraging activity, and growth (Wilson & Mckinley, 2004). These fish are typically sensitive to dissolved oxygen decreases. Cech & Doroshov (2004) reported that at moderate hypoxia ($3 \text{ mg O}_2 \text{ L}^{-1}$) at $15 \text{ }^\circ\text{C}$, the activity of Atlantic sturgeon did not change but gill ventilatory frequency increased by 50%. Also, Jenkins et al. (1993) report that regardless of the life stage, short-nose sturgeon died at oxygen concentrations of $2 \text{ mg O}_2 \text{ L}^{-1}$. For beluga hybrid sturgeon the critical concentration of dissolved oxygen was determined as $4 \text{ mg O}_2 \text{ L}^{-1}$ at $25 \text{ }^\circ\text{C}$ and the threshold concentration was $1.5 \text{ mg O}_2 \text{ L}^{-1}$ at $25 \text{ }^\circ\text{C}$ (Secor & Niklitschek, 2002). The natural stocks of this species are classified as Critically Endangered (CR) according to IUCN criteria (Gessner et al., 2022). The conservation of Küçükçekmece Lagoon, which serves

as a feeding ground for endangered species such as the beluga sturgeon, is crucial for their continued existence. To this end, improving water quality through appropriate measures is essential. In the short term, immediate action should focus on reducing pollution sources, such as enforcing stricter controls on industrial effluents and urban wastewater discharges into the lagoon. This would quickly alleviate some of the direct pressures on the water quality.

In the medium term, efforts should concentrate on restoring the natural flow and water levels of the lagoon by reevaluating the impact of the Sazlıdere Dam and exploring ways to increase freshwater input. This step will help in balancing the lagoon's ecosystem and reducing the salinity levels, which have been identified as a contributing factor to the deterioration of the habitat. Moreover, the employment of sturgeon culture facilities for conservation purposes, including the restocking of sturgeon into the lagoon or connected river systems, should be considered. This approach would not only aid in the direct replenishment of the sturgeon population but also serve as a vital tool for their conservation and genetic diversity.

For long-term sustainability, a comprehensive management plan should be developed, focusing on habitat restoration, including the revitalization of native vegetation along the shores and the creation of buffer zones to reduce nutrient runoff (Gessner et al., 2014). Additionally, establishing a continuous monitoring system to track water quality and biodiversity changes in the lagoon will provide valuable data for ongoing conservation efforts and help in adapting strategies as needed.

Implementing these measures in a phased approach will not only aid in the recovery and preservation of the Küçükçekmece Lagoon's ecosystem but also ensure the protection of species such as the beluga sturgeon, which rely on this habitat for survival.

CONCLUSIONS

In summary, the deaths of beluga sturgeon individuals and other fish species in the Küçükçekmece Lagoon, primarily attributed to pollution, algal blooms, and oxygen depletion, highlight the urgent need for conservation and rehabilitation efforts, especially during the summer season. Factors such as high temperature and salinity likely contributed to the stress on the beluga sturgeon, exacerbating their vulnerability. The broader challenges facing sturgeon species in Türkiye, including habitat degradation and migration barriers like hydroelectric power plants and bridges, further complicate their survival and reproduction.

While the sturgeon's current situation in Turkish waters is alarming, with their migration routes and spawning grounds severely impacted, there remains a significant opportunity to revitalize the Küçükçekmece Lagoon. This lagoon, situated within one of the world's major metropolises, Istanbul, still holds potential as a living wetland. To enhance the vitality of this critical habitat, specific measures are recommended:

1. Improving waste management and drainage systems to prevent pollution of the lagoon,

2. Ensuring compliance with and respect for the shoreline, thereby reducing habitat destruction and erosion,
3. Enhancing the organic-laden sediment structure, which is vital for the growth and sustenance of various aquatic species,
4. Replenishing Küçükçekmece lagoon with fresh water to balance the salinity levels and improve overall water quality.

Implementing these measures will not only contribute to the vitality of the Küçükçekmece Lagoon but also set a precedent for the conservation of other sturgeon habitats in Türkiye. It is imperative that a 'National Action Plan for the Conservation and Restoration of the Sturgeons of Turkey' be developed and executed by relevant authorities (Anon., 2015). This plan should focus on identifying and protecting the most suitable rivers for sturgeon survival and restoring their natural spawning grounds. Such efforts are crucial for the recovery of these ancient fish species and for preserving the biodiversity and ecological balance of Türkiye's aquatic ecosystems.

The findings of this study underscore the urgent need for a comprehensive and coordinated approach to conservation, highlighting the potential for recovery and revitalization of lagoon life, including the Küçükçekmece Lagoon. It is hoped that these insights will inform and inspire local, national, and international institutions to take decisive actions towards the sustainable management and preservation of these vital ecosystems.

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Table 1. Limitations for each manuscript type

Type of manuscript	Page	Abstract word limit	Reference limit
Original Article	≤20	250	40
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Figures, graphics, and photographs should be submitted as separate files (in TIFF or JPEG format) through the submission system. The files should not be embedded in a Word document or the main document. When there are figure subunits, the subunits should not be merged to form a single image. Each subunit should be submitted separately through the submission system. Images should not be labeled (a, b, c, etc.) to indicate figure subunits. Thick and thin arrows, arrowheads, stars, asterisks, and similar marks can be used on the images to support figure legends. Like the rest of the submission, the figures too should be blind. Any information within the images that may indicate an individual or institution should be blinded. The minimum resolution of each submitted figure should be 300 DPI. To prevent delays in the evaluation process, all submitted figures should be clear in

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Limitations, drawbacks, and the shortcomings of original articles should be mentioned in the Discussion section before the conclusion paragraph.

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While citing publications, preference should be given to the latest, most up-to-date publications. If an ahead-of-print publication is cited, the DOI number should be provided. Authors are responsible for the accuracy of references. List references in alphabetical order. Each listed reference should be cited in text, and each text citation should be listed in the References section. The reference styles for different types of publications are presented in the following examples.

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- American Psychological Association. (2010). Publication manual of the American Psychological Association (6th ed.). Washington, DC: APA.
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Citations must be indicated with the author surname and publication year within the parenthesis.

If more than one citation is made within the same paranthesis, separate them with (;).

Samples:

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Citation with one author;

(Akyolcu, 2007)

Citation with two authors;

(Sayiner & Demirci, 2007)

Citation with three, four, five authors;

First citation in the text: (Ailen, Ciambrene, & Welch, 2000)

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- **An eBook:** Millbower, L. (2003). *Show biz training: Fun and effective business training techniques from the worlds of stage, screen, and song*. Retrieved from <http://www.amacombooks.org/> (accessed 10.10.15)
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- **An article with DOI:** Gaudio, J. L. & Snowdon, C. T. (2008). Spatial cues more salient than color cues in cotton-top tamarins (*saguinus oedipus*) reversal learning. *Journal of Comparative Psychology*, <https://doi.org/10.1037/0735-7036.122.4.441>
- **Websites - professional or personal sites:** The World Famous Hot Dog Site. (1999, July 7). Retrieved January 5, 2008, from <http://www.xroads.com/~tcs/hotdog/hotdog.html> (accessed 10.10.15)

- **Websites - online government publications:** U.S. Department of Justice. (2006, September 10). Trends in violent victimization by age, 1973-2005. Retrieved from <http://www.ojp.usdoj.gov/bjs/glance/vage.htm> (accessed 10.10.15)
- **Photograph (from book, magazine or webpage):** Close, C. (2002). *Ronald*. [photograph]. Museum of Modern Art, New York, NY. Retrieved from http://www.moma.org/collection/object.php?object_id=108890 (accessed 10.10.15)
- **Artwork - from library database:** Clark, L. (c.a. 1960's). *Man with Baby*. [photograph]. George Eastman House, Rochester, NY. Retrieved from ARTstor
- **Artwork - from website:** Close, C. (2002). *Ronald*. [photograph]. Museum of Modern Art, New York. Retrieved from http://www.moma.org/collection/browse_results.php?object_id=108890 (accessed 10.10.15)

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