

Large-Scale Geographic Size Variability of *Cyprideis torosa* (Ostracoda) and Its Taxonomic and Ecologic Implications

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Wrozyna C, Mischke S, Hoehle M, Gross M and Piller WE (2022) Large-Scale Geographic Size Variability of Cyprideis torosa (Ostracoda) and Its Taxonomic and Ecologic Implications. Front. Ecol. Evol. 10:857499. doi: 10.3389/fevo.2022.857499 Body-size variability results from a variety of extrinsic and intrinsic factors (environmental and biological influences) underpinned by phylogeny. In ostracodes it is assumed that body size is predominantly controlled by ecological conditions, but investigations have mostly focused on local or regional study areas. In this study, we investigate the geographical size variability (length, height, and width) of Holocene and Recent valves of the salinity-tolerant ostracode species Cyprideis torosa within a large geographical area (31°-51° latitude, and 12°-96° longitude). It is shown that distant local size clusters of Cyprideis torosa are framed within two large-scale geographical patterns. One pattern describes the separation of two different size classes (i.e., morphotypes) at around \sim 42° N. The co-occurrence of both size morphotypes in the same habitats excludes an environmental control on the distribution of the morphotypes but rather could point to the existence of two differentiated lineages. Generally, correlations between valve size and environmental parameters (salinity, geographical positions) strongly depend on the taxonomic resolution. While latitude explains the overall size variability of C. torosa sensu lato (i.e., undifferentiated for morphotypes), salinity-size correlations are restricted to the morphotype scale. Another large-scale pattern represents a continuous increase in valve size of C. torosa with latitude according to the macroecological pattern referred as Bergmann trend. Existing explanations for Bergmann trends insufficiently clarify the size cline of C. torosa which might be because these models are restricted to intraspecific levels. The observed size-latitude relationship of C. torosa may, therefore, result from interspecific divergence (i.e., size ordered spatially may result from interspecific divergence sorting) while environmental influence is of minor importance. Our results imply that geographical body-size patterns of ostracodes are not straightforward and are probably not caused by universal mechanisms. Consideration of phylogenetic relationships of ostracodes is therefore necessary before attempting to identify the role of environmental controls on body size variability.

Keywords: body size, geographical variability, Bergmann's rule, ostracodes, Cyprideis, morphometrics

INTRODUCTION

Body size is a master trait of an organism, because it influences nearly every aspect of the biology of an organism (Verberk et al., 2021). It is determined by and/or responds to extrinsic environmental factors (abiotic and biotic) and intrinsic (biological) influences that reflect direct functional and/or evolutionary effects, all further underpinned by phylogeny (Hart and Bychek, 2011). These common underlying causes and processes result in consistent large-scale patterns across temporal, spatial or taxonomical scales. One of the longest and best-known ecological generalizations is that large body size is typically found at higher latitudes, or in colder climates as first described by Bergmann (1847)) for mammals. Bergmann's rule has also been demonstrated to apply to some groups of ectotherms, but observed patterns and possibly underlying mechanisms remain elusive (Ashton and Feldman, 2003; Blanckenhorn et al., 2006; Tesche and Hodges, 2015; Maszczyk and Brzeziński, 2018).

Within the crustaceans, ostracodes are exceptionally wellsuited for the investigation of Bergmann clines due to their high taxonomic diversity (ca. 2000 extant non-marine species; Martens et al., 2008), an extensive fossil record (Horne, 2005; Martens et al., 2008), and an extraordinary diversity of life-history strategies (Chaplin et al., 1994; Butlin et al., 1998; Geiger et al., 1998). This, in combination with their relatively short generation time, denote ostracodes as valuable model organisms to study fundamental evolutionary questions (Fernandes Martins et al., 2009). Moreover, the ubiquitous distribution of ostracodes in nearly all freshwater and marine habitats, but also interstitial and semi-terrestrial environments (Horne et al., 2002) allows to investigate geographical variation of body-size traits in various environments (e.g., marine vs. freshwater; tropical vs. temperate).

However, ostracodes (and crustaceans in general) were neglected in research of macroecological patterns so far. Therefore, there is poor knowledge on effects of geographic variation on overall ostracode size. Actually, ostracode-body size was mainly investigated with respect to ecophenotypical responses which are mostly based on local or small regional scales and/or related to single environmental parameters (e.g., Yin et al., 2001). These studies have shown that ostracode size is influenced by salinity (Boomer et al., 2017), temperature (Martens, 1985), as well as pH and CaCO₃ concentration (Kim et al., 2015). Hitherto, Bergmann trends were detected only in deep-sea species (Hunt and Roy, 2006; Hunt et al., 2010).

In this study, we investigate geographical size variation of an ostracode taxon generally assigned to the species *Cyprideis torosa*. *Cyprideis torosa* was chosen because it has a large geographical range which covers the Eurasian realm and Africa (Wouters, 2017; **Figure 1**) where it occurs in freshwater to hypersaline habitats and is characterized by tolerance of a wide temperature range and various oxygen and substrate conditions (De Deckker and Lord, 2017). In addition, high abundance of valves in Quaternary deposits (e.g., West et al., 1994; Pint et al., 2012) enables examination of body-size traits on different temporal scales.

We investigate valve-size variability of C. torosa along a 20° latitudinal range to address questions related to evolution

of body-size clines in non-marine ostracodes. In order to understand if and how specific these relationships are, we compare the new information with a dataset of the Neotropical species *Cytheridella ilosvayi* (Wrozyna et al., 2016, 2018, 2019). Questions addressed include: (1) How does size vary within *C. torosa*'s geographical range and the Holocene timescale? (2) Is there a clinal variation in body size (valve length, height, and width) of *C. torosa* across latitude? (3) Is there a size variation in the species according to salinity?

MATERIALS AND METHODS

Sampling Sites

Our study area was chosen with respect to cover inland occurrences and combination of fossil and Recent material. The study area includes nine locations and represents a wide geographical range from ca. 31° to 51° N and 12° to 96° E (Germany, Romania, Turkey, Israel, Kazakhstan, China), coinciding roughly with the known geographical range of the species in Europe and Asia (Wouters, 2017; **Figure 1**).

The included ostracod material covers a stratigraphic range from Middle Pleistocene to Recent (**Table 1**). Majority of the localities represents distinct inland occurrences. Only coastal lakes [Hersek lagoon (HK) and Danube Delta (BS)] probably have at least a temporary connection to marine conditions. Some localities are represented by more than one sample either from different water bodies within a small geographical region or from different stratigraphic levels of a sedimentary sequence (**Table 1**). Since the foremost question of this study addresses the variation between populations and/or localities, all samples from one locality or nearby locations where habitat connectivity is theoretically possible (e.g., the ISR samples from the wetland Enot Zuqim, Israel; **Table 1**) were pooled. Recent *Cyprideis* valves are derived from bio- and taphocenoses.

Salinity data of modern populations of Mansfeld Lakes (MS) and Enot Zuqim (ISR) were measured in situ in the field with hand-held probes. Estimation of paleo-specific conductivity for the fossil localities of Lake Balkhash (BKA), Sea of Galilee (SOG), and Tarim Basin (YDK) were derived by the application of ostracode-based transfer functions established for the Near East and the Tibetan Plateau. These transfer functions are calibrated with modern quantitative relationships between species compositions and related conductivity values. Applied by different statistical approaches (e.g., weighted-averaging partial least squares regression) to fossil species assemblages, reliable estimates of past conductivity can be derived (Kalanke, 2015; Mischke et al., 2019, 2020). Since conductivity values are calculated for each stratigraphic layer based on relative abundances of ostracode species, assemblage variations in a sedimentary sequence of a specific locality typically result in variations of the calculated conductivity values. Variation ranges are, however, small varying between 0.5 and 1.3 PSU for samples of Sea of Galilee (SOG) and 1.5 and 1.6 PSU for Lake Balkhash (BKA).

Hersek lagoon (HK) is located at the eastern Marmara Sea. The fossil record continuously indicates polyhaline conditions.



Samples were derived from the upper part of a sediment core dating back to \sim 1200 years BP. Since the composition of the ostracode fossils did not change during the considered time a salinity value of today (29 PSU; Mischke et al., 2012) was assumed as reliable estimate of the paleo-salinity. A salinity estimate for the modern samples from the Danube Delta (BS) was taken from the literature (van de Velde et al., 2019). Samples from Lake Bosten (XB) were excluded from size-salinity regression analyses since no salinity data were available. Recorded specific conductivity data were converted to salinity for comparison with published salinity data using the factor of 0.65 (Hem, 1982; **Table 1**).

Morphological Variables

All adult specimens were sorted under a binocular microscope and photographed with a digital Keyence VHX-950F microscope. Photographs of left and right valves of females and males representatives for each locality are provided in the **Supplementary Figure 1**. Measurements of maximum length, height, and width were completed directly *via* the Keyence software (VHX-7000 Series Software) on screen (**Figures 1A,B**). Valve-width measurements were obtained as "highest points" on focus-stacked photographs which were automatically merged by the software of the microscope. This approach for the measurement of valve width was not yet used for ostracodes. Due to dimorphism, both sexes and both left and right valves were investigated.

Statistical Analyses

Prior to statistical analyses, size, salinity, and geographical data were log-transformed and all analyses were done separately for the four sub-data sets of female and male left and right valves. Site-specific size variability was tested for normal distribution by Shapiro Wilk's tests separately for length, height and width. Multivariate normality was tested on overall data sets (length, height, width) by Mardia test of multivariate skewness and kurtosis followed by Doornik and Hansen omnibus tests. Principal component analyses were used to explore the overall size variability in relation to morphotypes distribution. Classification of the specimens according to a morphotype group was based on histograms of length. Threshold values for the differentiation of the morphotypes were derived from the areas (i.e., bins) with minimal overlapping of both size groups in the histograms. As a Permutational Multivariate Analysis of Variance (PerMANOVA) assumes no (normal) distribution it was selected for testing of significant differences between groups (i.e., morphotypes, sites) based on all size traits (length, height, width). Pairwise comparisons (i.e., post hoc analyses) were done by pairwise Hoteling's tests of each site. Relationships between salinity and size traits were investigated by linear regressions on log-transformed salinity and size data. Redundancy analyses (RDA) were performed to assess which of the investigated potential explanatory variables, geographical region (i.e., latitude and longitude) or salinity, provide largest explanatory power on

TABLE 1 Sample-locality information for the investigated valves of Cyprideis torosa. Classification of the climate follows the Köppen system.

ID	Locality	Country	Latitude	Longitude	Altitude [m. a. s. l.]	Climate	Catchment area [km ^{-2]}	freshwater biome	Salinity (measured or estimated values in PSU)	Age	Age range of the sample	Number of valves [left females/left males/right females/right males]	Reference	Remarks
YDK	Lop Nur, Tarim Basin	China	40.20806	90.2997	780	today Bwk/past probably wetter	unknown	lake	mesohaline (11.5)	Early Holocene		29/27/30/28	Mischke et al., 2017, 2019	
XB	Lake Bosten	China	41.94900	86.775	1048	today Bwk/past probably wetter	today ~ 56,000	lake	oligohaline to possibly mesohaline (-)	Middle Holocene		12/21/24/19	Mischke and Wünnemann, 2006	
MS	Mansfeld Lakes	Germany	51.48498	11.7405	90	Cfb	411	lake	oligohaline (0.8-4.7)	Recent		33/16/18/18	Wrozyna, own data	*
SOG	Sea of Galilee	Israel	32.83861	35.5337	-212	today Bsh/past probably	2730	lake	oligohaline (0.5-1.3)	Late Holocene	~100 - 1600 years BP	29/22/35/20	Kalanke, 2015	**
ISR	Shore of the Dead See	Israel	31.7106	35.4537		Bsh	40,000	wetland	oligohaline (3.2-4)	Recent		25/27/46/25	Mischke et al., 2014	***
BKA	Lake Balkhash	Kazakhstan	46.28922	74.0085	314	Dfb	501,000	lake	oligohaline (1-3-1.6)	Late Holocene		80/45/81/53	Mischke et al., 2020	ŧ
BS	Danube Delta	Romania	44.61537	28.9296	0	Cfb	Black Sea	delta lake	oligohaline (1)	Recent		13/13/12/9	Opreanu, own data	†
ΗK	Hersek lagoon	Turkey	40.724	29.519	0	Csa	Sea of Marmara	coastal lake	polyhaline (29)	Late Holocene		20/16/17/12	Mischke et al., 2012	

* Samples pooled from three lakes including Lake Süß, Lake Binder and Lake Kerner.

** Samples were derived from different depths of a sediment core. *** Samples pooled from 6 different localities in a wetland at the shore of the Dead Sea.

\$Samples from a sediment core and surface sediments pooled together.

+Specimens collected alive from two adjacent lakes.

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the size data (length, height, width) of *C. torosa*. All statistical analyses were done with the software PAST (version 4.09; Hammer et al., 2001).

RESULTS

Size Variability

In total 875 valves composed of 504 females and 371 males were investigated. Minimum sample sizes of the different sites range from 12 and 13 valves of females (left and right valves, respectively) and 12 and 9 male valves (left and right, respectively) to maximum samples sizes of 80 and 81 female valves (left and right, respectively) and 46 and 53 male valves (left and right, respectively). The mean number of investigated valves per sample is 109 (left and right males and females together). With a total of 259 valves, Lake Balkhash provided largest total sample size (161 female valves and 98 male valves). Lowest valve numbers were provided by the sample of the Danube Delta (BS) with a total of 47 valves (25 females and 22 males; **Table 1**).

The valve length of adult left and right valves of female *C. torosa* varies from 742 to 1087 μ m and 723 to 1049 μ m, respectively. Male specimens show ranges from 861 to 1182 μ m and 795 to 1155 μ m for left and right valves, respectively. The width of left and right valves of female specimens is in a range from 166 to 307 μ m and 140 to 287 μ m, respectively. Male valves vary between 121 and 299 μ m (left valves) and 127 to 302 μ m (right valves). Generally, right valves of both males and females are smaller and less vaulted outside (**Figures 2A,B**).

According to a Shapiro-Wilk test most traits and samples exhibit normal distribution ($\alpha = 0.05$; Table 2). Size traits differ with respect to their normal distribution. While length and height display normal distributions for the majority of samples the number of normal distributed samples is lower for valve width. Not normally distributed samples are majorly from Enot Zuqim (ISR) and Lake Balkhash (BKA; Table 2). Mardia tests for multivariate normality indicate that the overall data sets of left and right females and left and right males are not multivariate normally distributed (female LV: skewness: 2.683, $p \le 0.001$; kurtosis: 29.86, p = 0; Doornik and Hansen omnibus E = 128.2, $p \le 0.001$; female RV: skewness: 1.935, $p \le 0.001$; kurtosis: 18.24, $p \le 0.001$; Doornik and Hansen omnibus E = 26.17, $p \le 0.001$; male LV: skewness: 3.835, p < 0.001; kurtosis: 21.4, p < 0.001; Doornik and Hansen omnibus E = 40.37, $p \le 0.001$; male RV: skewness: 1.935, *p* < 0.001; kurtosis: 18.24, *p* < 0.001; Doornik and Hansen omnibus E = 26.17, $p \le 0.001$).

While there is a distinct linear relationship between length and height (*r*-values for all relationships ≥ 0.96 , $p \leq 0.001$), the regression relationship is slightly reduced between length and width (*r*-values between 0.63 and 0.75; $p \leq 0.001$).

Although variance becomes generally smaller with larger sample sizes, there are exceptions as for instance in samples of Enot Zuqim (ISR) in which large variation ranges are caused by presence of two unevenly distributed size morphotypes (**Figure 2F**). In both length-height and length-width plots, it is indicated that *C. torosa* represents two size classes or morphotypes (**Figure 2**). The existence of two morphotypes is also expressed in bimodal distributions of length and height data (Figure 2). Means of the morphotypes' length differ significantly with the smaller morphotype I providing for females left and right valves 829 and 803 µm, and for male left and right valves 929 and 897 µm (also Figure 2; Supplementary Tables 1-4). The larger morphotype II is characterized by mean length of female left and right valves of 952 and 937 µm, and male left and right valves display mean lengths of 1040 and 1023 µm. Valve width generally does not support a separation in two or more size classes (Figure 2E). Samples from Enot Zuqim (ISR) and Sea of Galilee (SOG) show in the width data an ambiguous pattern slightly following a bimodal distribution as length and height. Regression lines of length-height relationships are similar for both morphotypes. Divergence occurs by proportional differences in valve width-length relationships. Morphotype I is, thus, characterized by slightly larger width relative to the valve length if compared to the larger morphotype II. Also, correlation coefficients for the width-length relationship are higher for morphotype II (Table 3).

The first two axes of the PCAs performed on all size traits explain in all sub-data sets > 98% of the total variance (Figure 3). PC 1 explains between 80.8% (female right valves) and 87.9% (female left valves; 83.2% male left valves; 81.9% male right valves) and is in all analyses evenly high correlated with length, height, and width (r-values range for length between 0.87 and 0.94, for height between 0.88 and 0.95, and for width between 0.89 and 0.93). Scatter plots disclose the disjunct size variability of the morphotypes with morphotype I displaying negative scores on PC 1 and morphotype II having positive scores on PC 1. As indicated by histograms and length-height plots valves from Hersek lagoon (HK) provide an intermediate position in the scatter plots caused by co-occurrence of both morphotypes. PerMANOVA results showed that morphotypes are significantly different (females RV: total sum of squares = 1.419, within-group sum of squares = 0.695, F = 257.0, p < 0.001; females LV: total sum of squares = 1.375, within-group sum of squares = 0.534, F = 391.4, p < 0.001; males RV: total sum of squares = 0.978, within-group sum of squares = 0.551, F = 138.6, p < 0.001; males LV: total sum of squares = 0.844, within-group sum of squares = 0.404, F = 198.7, p < 0.001).

Latitudinal Variability of Size

Generally, largest valves occur in Mansfeld Lakes (MS) followed by Lake Bosten (XB), Lake Balkhash (BKA), Black Sea (BS), and Hersek lagoon (HK). Smallest valves occur in Enot Zuqim (ISR), Tarim Basin (YDK), and Sea of Galilee (SOG; **Supplementary Figures 2**, **3**). Thereby, the cluster of the smaller valve sizes (i.e., morphotype I) includes all valves from Sea of Galilee (SOG) and Tarim Basin (YDK). Valves of Hersek lagoon and Enot Zuqim (ISR) contribute to both clusters. While valve sizes for Hersek lagoon can be considered intermediate, probably including both morphotypes, the data for Enot Zuqim represent two distinctly separated size classes in which only one locality contains solely valves of the larger size cluster while the other localities include only the valves sizes according to the smaller cluster (**Figure 2F**).



FIGURE 2 | Length-height (A) and length-width (B) plots for female and male left and right valves (LV and RV, respectively). Distribution of size data presented as histograms of valve length (C), valve height (D), valve width (E), and length-height plot in which populations are indicated (F). Population abbreviations are: YDK, Tarim Basin, China; XB, Lake Bosten, China; BKA, Lake Balkhash, Kazakhstan; ISR, Dead Sea, Israel; SOG, Sea of Galilee, Israel; BS, Danube Delta, Romania; HK, Hersek lagoon, Turkey; MS, Mansfeld Lakes, Germany. Details about the localities are provided in Table 1. Displayed are data of female left valves, other datasets are available in the Supplementary Material.

Within-sample variability of length shows similar ranges for males and females between 59 and 178 μm and 45 to 220 μm , respectively. Small sample sizes generally provide lower

differences (e.g., XB). Maximum length differences are recorded for HK (males) and XB (females). Absolute differences for valve width range for males and females between 59 and 178 μm and 32

		Normal distribution according to Shapiro Wilk's test ($\alpha > 0.05$)	Non-normal distribution according to Shapiro Wilk's test ($\alpha < 0.05$)
Length	Female LV	BKA, SOG, MS, YDK, HK, BS	XB, ISR
	Female RV	BKA, SOG, MS, YDK, HK, XB, BS	ISR
	Male LV	SOG, MS, XB, BS, HK, YDK	BKA, ISR
	Male RV	BKA, SOG, MS, YDK, HK, ISR, XB, BS	-
Height	Female LV	BKA, SOG, MS, YDK, HK, XB, BS	ISR
	Female RV	BKA, SOG, MS, XB, BS, YDK	ISR, HK
	Male LV	SOG, MS, XB, ISR, HK, YDK	BKA, BS
	Male RV	BKA, SOG, MS, YDK, HK, ISR, XB, BS	-
Width	Female LV	SOG, MS, YDK, HK, XB, BS	BKA, ISR
	Female RV	MS, BS, HK	BKA, SOG, ISR, XB, YDK
	Male LV	MS, ISR, BS, YDK	BKA, SOG, XB, HK
	Male RV	BKA, SOG, MS, HK, ISR, XB	YDK, BS

TABLE 2 Overview of Shapiro Wilk's tests for normal distribution separately for length, height, and width of female and male left (LV) and right (RV) valves.

Abbreviations of localities are as following: YDK, Tarim Basin, China; XB, Lake Bosten, China; BKA, Lake Balkhash, Kazakhstan; ISR, Dead Sea, Israel; SOG, Sea of Galilee, Israel; BS, Danube Delta, Romania; HK, Hersek lagoon, Turkey; MS, Mansfeld Lakes, Germany.

TABLE 3 Differences of length-height regression coefficients and slopes of both Cyprideis morphotypes indicated for both sexes and left (LV) and right (RV) valves.

			n	Length	n-height		Length-width			
				Equation	<i>r</i> -value	<i>p</i> -value	Equation	<i>r</i> -value	p-value	
LV	Female	Morphotype I	100	Y = 1.035 * X - 0.346	0.94	≤0.001	Y = 0.288 * X + 2.25	0.46	≤0.001	
		Morphotype II	136	Y = 0.934 * X - 0.04	0.85	≤0.001	Y = 0.211 * X + 2.48	0.49	≤0.001	
RV	Female	Morphotype I	98	Y = 1.066 * X - 0.45	0.97	≤0.001	Y = 1.381 * X - 1.72	0.69	≤0.001	
		Morphotype II	148	Y = 0.995 * X - 0.24	0.89	≤0.001	Y = 1.681 * X - 2.66	0.55	≤0.001	
LV	Male	Morphotype I	74	Y = 1.018 * X - 0.34	0.92	≤0.001	Y = 0.710 * X + 0.203	0.38	≤0.001	
		Morphotype II	36	Y = 1.224 * X - 0.964	0.87	≤0.001	Y = 1.062* X - 0.82	0.48	0.003	
RV	Male	Morphotype I	73	Y = 1.034 * X - 0.40	0.84	≤0.001	Y = 0.90 * X - 0.40	0.48	≤0.001	
		Morphotype II	46	Y = 1.076 * X - 0.52	0.92	≤0.001	Y = 1.31 * X - 1.59	0.64	≤0.001	

The asterisk is a common symbol to indicate multiplication.



FIGURE 3 | Principal component analysis of the variability of size traits (length, height, width) of females left valves of *Cyprideis torosa*. Indicated is the small morphotype I (orange) and the larger morphotype II (gold). Valves of Hersek lagoon (HK) where both morphotypes co-occur are highlighted in gray.

and 121 μ m, respectively. ISR provides the largest within-sample variability for width of male and female valves (**Supplementary Figures 2B, 3**).

PerMANOVA results showed that the sites are significantly different (females RV: total sum of squares = 1.419, withingroup sum of squares = 0.570, F = 51.24, p < 0.001; females LV: total sum of squares = 1.375, within-group sum of squares = 0.388, F = 87.86, p < 0.001; males RV: total sum of squares = 0.997, within-group sum of squares = 0.284, F = 62.60, p < 0.001; males LV: total sum of squares = 0.844, within-group sum of squares = 0.243, F = 62.24, p < 0.001). Pairwise comparisons (i.e., *post hoc* tests) confirmed that valve size is different between the sites. Exceptions represent sample pairs BKA and BS, and ISR and YDK, which are

TABLE 4 | Regression coefficients of log-transformed size-latitude relationships for *Cyprideis torosa* and *Cytheridella ilosvayi* separated for sexes and left and right valves (LV and RV, respectively).

		Cypride	is torosa		Cytheridella ilosvayi		
		Female LV	Male LV		Female LV	Male LV	
	Latitude-log length	0.81 (p < 0.001)	0.82 (p < 0.001)	Latitude-log length	0.10 (p = 0.291)	0.02 (p = 0.895)	
		Female RV	Male RV		Female RV	Male RV	
		0.77 (p < 0.001)	0.81 (p < 0.001)		0.09 (p = 0.317)	0.21 (p = 0.170)	
		Female LV	Male LV		Female LV	Male LV	
	Latitude-log height	0.79 (p < 0.001)	0.85 (0.001)	Latitude-log height	0.26 (p = 0.007)	0.31 (p = 0.017)	
		Female RV	Male RV		Female RV	Male RV	
		0.77 (p < 0.001)	0.83 (p < 0.001)		0.13 (p = 0.206)	0.03 (p = 0.823)	
		Female LV	Male LV				
	Latitude-log width	0.57 (p < 0.001)	0.49 (p < 0.001)				
		Female RV	Male RV				
		0.34 (p < 0.001)	0.43 (p < 0.001)				
		Female LV	Male LV				
Latitude-log length	Morphotype I	0.09 (p = 0.333)	0.25 (p = 0.029)				
	Morphotype II	0.32 (p < 0.001)	0.29 (p = 0.077)				
		Female RV	Male RV				
	Morphotype I	0.46 (p < 0.001)	0.13 (p = 0.276)				
	Morphotype II	0.56 (p < 0.001)	0.66 (p < 0.001)				
		Female LV	Male LV				
Latitude-log height	Morphotype I	0.38 (p < 0.001)	0.34 (p = 0.003)				
	Morphotype II	0.32 (p < 0.001)	0.64 (p < 0.001)				
		Female RV	Male RV				
	Morphotype I	0.58 (p = 0.001)	0.44 (p < 0.001)				
	Morphotype II	0.62 (p < 0.001)	0.63 (p < 0.001)				
Latitude-log width		Female LV	Male LV				
	Morphotype I	0.09 (p = 0.374)	0.16 (p = 0.161)				
	Morphotype II	0.39 (p < 0.001)	0.02 (p = 0.239)				
		Female RV	Male RV				
	Morphotype I	0.24 (p = 0.022)	0.05 (p = 0.692)				
	Morphotype II	0.39 (p < 0.001)	0.39 (p < 0.001)				

Regression analyses of C. torosa are presented undifferentiated and morphotype-specific.

not significantly different in terms of their size characteristics (Supplementary Figure 2).

Size in terms of length, height and width of morphotypes is moderately to not correlated to latitude. The *r*-values are very variable ranging between r = 0.02 and r = 0.64 with a mean of r = 0.25 (**Table 4**). In contrast, considering length and height of *C. torosa* valves as one group they are correlated with latitude (**Figure 4**), with correlation coefficients of $r \ge 0.77$ for all data sets (female and male left and right valves; **Table 4**). Width displays lower and more heterogeneous *r*-values between males and females. While r-values of latitude-width for males are almost identical (r = 0.43 and r = 0.49, respectively) females provide a relatively large range around the values of the males. Thus, the highest correlation is exhibited by left female valves (r = 0.57) in contrast to a relatively low *r*-value shown by right female valves (r = 0.34).

Size-Salinity Relationships

Variation of valve size in relation to salinity reveals an opposite relationship to that observed in the latitude-size relationship,

but this also depends on whether morphotypes are considered or not. Correlation coefficients between valve size traits and salinity are low if *C. torosa* is considered undifferentiated for morphotypes (*r*-values range between 0 and 0.16) of both female and male valves. Correlation coefficients are distinctly higher if the relationships are considered separately for both morphotypes. Thus, size traits of both morphotypes of *C. torosa* are positively correlated with salinity. This is particularly true for length and height. Valve width displays only for morphotype II a weak positive relationship with salinity (**Figure 5** and **Table 5**).

The RDA explains between 37.5% (female right valves) and 49.9% (female left valves; 44.4% male left valves; 46.4% male right valves) of the variability in the size data. Variabilities explained by the RDA axes 1–3 were highly significant at p = 0.001. The explanatory percentages of the axes differ greatly with very low contributions of the second and third axes to the overall variability (<0.6% and <0.8% of left female and male valves; and 2 and 2.2% of right female and male valves), respectively. According to the RDA biplots, latitude has the strongest influence



on valve length and height while salinity and longitude are not relevant for the overall variability in size data. Valve width is not controlled by geographical area and salinity (**Figure 6**).

DISCUSSION

Valve-Size Variability

Cyprideis torosa s.s. occurs at the beginning of the Calabrian Stage (middle Early Pleistocene) (Gliozzi et al., 2017) and persists until present day without known speciation as documented from African *Cyprideis* (Wouters and Martens, 1994, Wouters and Martens, 2001; Schön et al., 2000). Therefore, all fossil and Recent records are considered the same species and it is not reasonable to treat them differently. Additionally, our results show no pattern in valve size related to Holocene temperature conditions (see below), so the combination of Recent and fossil records is not expected to promote any of the patterns found in the dataset.

The observed size-variation ranges within samples as well as the overall data set of *C. torosa* fits very well to known sizevariability patterns of modern *C. torosa* (Boomer et al., 2017) and fossil *Cyprideis* species (Gitter et al., 2015). It is somewhat surprising that the length variation detected by Berndt et al. (2019) of 770–1036 μ m for females and 903–1106 μ m for males, respectively, almost completely matches our data range although these data are derived from a single sediment-core location covering the last ~7800 years. This similarity highlights not only the large local variation in size over time, but also that (large) spatial and (long) temporal variations can have similar amplitudes. Actually, it is assumed that geographical and temporal changes in body size are manifestations of the same drivers (Yom-Tov and Geffen, 2011).

As shown by the length-width plot (**Figure 2B**), morphotypes are differentiated not only by length but also by proportions in valve length and width expressed by differences in the regression slopes (**Table 3**). Morphotype I displays, thus, slightly larger width relative to the valve length compared to morphotype II. This indicates that valve shape could be also different.

Size-Latitude Relationships

So far, data sets of geographical size variability of ostracode species are mostly restricted to relatively small (i.e., regional) scales. Our size data revealed two large-scale geographical patterns. The first is the almost complete separation in two size classes at around \sim 42° N (**Figures 5A–C**), and the second is that clinal variation of body size of *C. torosa* is consistent to the Bergmann trend.

While the Bergmann trend describes a successive increase of valve length with latitude, *Cyprideis* also shows a differentiation between smaller and larger size populations (i.e., morphotypes I and II) (**Figures 2,3,6**). This could be an additional size cline representing, e.g., the adaptation to different environmental conditions as observed in other organisms (Bai et al., 2016). However, localities Enot Zuqim (ISR) and Hersek lagoon (HK) contain both morphotypes, which refuses an ecological control.

Co-occurrence of two size groups can result from different generations per year which is in turn affected by length of the warm season. Thus, populations in warmer areas with a longer summer period produce two generations per year, and oppositely, only one generation per year occurs due to a shorter warm period in colder regions. This seasonal differentiation is often accompanied by different body sizes related to differences in the developmental time (Aguilar-Alberola and Mesquita-Joanes, 2014; Wesławski et al., 2020). Actually, life history and/or seasonal population structure of C. torosa is ambiguous. While Mediterranean populations clearly show a bivoltine (Mezquita et al., 2000) reproduction pattern, northern occurrences provide evidence for both univoltine (Belgium; Heip, 1976) and bivoltine (United Kingdom; Roberts et al., 2020) patterns. Nonetheless, a bivoltine reproduction would result in a slightly overlapping but double peaked distribution of size data at local scales. Enot Zugim (ISR) samples show such a pattern. Samples were derived from different but possibly connected sites within a wetland. This indicates that morphotypes have a disjunct distribution even within a small geographical area. Only Hersek lagoon (HK) provides both morphotypes. The co-occurrence of different size groups as a result from bivoltine reproduction is therefore only for Hersek lagoon (HK) possible.





Although body-size evolution results from diverse factors, divergence of size can be consistent with phylogenetical patterns (Suatoni et al., 2006; Haye et al., 2019). It has been shown that interspecific size distribution is not random but instead probably organized through interspecific interactions. Two processes are known that lead to population-wide patterns of interspecific body-size variations. The first is adjustment (i.e., character displacement) which is an evolutionary response of species in sympatry and allopatry. The other one represents size assortment according to ecological interaction among species in which different-sized species can successfully colonize the same locality together (Sota et al., 2000).

In contrast, intraspecific morphological divergence may result from phenotypical plasticity, where the influence of abiotic or biotic selective forces could lead to the expression of alternative phenotypes between populations from a single genotype, or divergent selection drives local adaptation and heritable evolutionary differences in phenotypes on account of genetic differentiation (Bower et al., 2018). The bimodal sizepattern of *C. torosa* (**Figures 2C,D**) contradicts assumptions on the expression of phenotypical plasticity, which is usually associated with gradual successions in morphological traits (Emlen et al., 2007; Lafuente and Beldade, 2019). The observed body-size patterns may, therefore, reflect interspecific size differentiation. TABLE 5 | Regression coefficients of log-transformed size and salinity data for females and males left (LV) and right valves (RV).

			n	Length	n-salinity	Height	-salinity	Width-salinity	
				<i>r</i> -value	<i>p</i> -value	<i>r</i> -value	<i>p</i> -value	<i>r</i> -value	<i>p</i> -value
LV	Female	Morphotype I	94	0.57	≤0.001	0.6	≤0.001	0.02	0.822
		Morphotype II	120	0.38	≤0.001	0.45	≤0.001	0.39	≤0.001
RV	Female	Morphotype I	98	0.56	≤0.001	0.63	≤0.001	0.24	0.238
		Morphotype II	123	0.47	≤0.001	0.58	≤0.001	0.39	≤0.001
LV	Male	Morphotype I	75	0.57	≤0.001	0.33	0.003	0.14	0.229
		Morphotype II	72	0.57	≤0.001	0.43	0.004	0.28	0.017
RV	Male	Morphotype I	77	0.43	0.001	0.74	≤0.001	0.11	0.362
		Morphotype II	46	0.72	≤0.001	0.69	≤0.001	0.84	≤0.001

Number of investigated specimens (n) is given for each analysis. Locality HK was excluded from the regressions due to co-occurrence of both morphotypes.



Although extensive investigations of morphological and/or genetical divergence of *C. torosa* are missing, there are hints pointing to spatial variations in morphology (Wouters, 2017) and genetics (Sywula et al., 1995). In our dataset, the occurrence of distinct morphotypes with little geographical overlap could point to the existence of at least two different species. Nonetheless, integrated approaches combining examinations of morphological traits, preferably by quantitative and genetic data, are therefore needed to evaluate not only the importance of divergence in size of *Cyprideis* but also to achieve a more complete understanding of spatial morphological variability and underlying causes.

Body size of crustaceans (marine and freshwater) generally display Bergmann trends as reported in the review by Maszczyk and Brzeziński (2018). It is, however, not clear whether these patterns result from direct effects of temperature or from the interaction of temperature with other extrinsic factors. Actually, it is stated that temperaturedependent oxygen limitation (Huston and Wolverton, 2011; Horne et al., 2015; Rollinson and Rowe, 2018) is a major controlling factor for aquatic invertebrates beside a number of other factors (e.g., sexual selection, predation) that could cause correlations between body size and latitude (Verberk et al., 2011). It has been documented that rate of growth and size of ostracodes is affected by water temperature. Accordingly, valves of the same species will reach a larger size under lower temperature (Martens, 1985). Later, a strong temperature effect was noted on inter-molt time but not on growth factors (Liberto et al., 2014) indicating other or additional factors (e.g., genetical) that control the ultimate valve size. The explanatory power of these studies is, however, limited since they do not provide hints on side effects of temperature changes or related changes of other parameters such as oxygen availability. By this, a secondary dependence of temperature on the valve size could be related to the influence on the development time.

Hunt and Roy (2006) and Hunt et al. (2010) provide the only studies on ostracode size with respect to latitudinal patterns which showed that body size of deep-sea ostracode species has increased greatly over the past 40 million years mirroring the trajectory of bottom-water temperature that can be translated into a latitudinal cline of size according to the Bergmann trend.

Accepting that size clines are related to parameters connected to latitude such as temperature, climate change will lead to changes in body size (Teplitsky and Millien, 2014). Our data cover a relatively large temporal range from Early Holocene to modern day. Populations lived under warmer and colder periods relative to mean temperature variations of the late 20th century (**Figure 7**). If temperature is a major control on body size of *Cyprideis*, the distribution of size should be different from what we actually found with largest valves occurring under coldest temperatures and vice versa. However, there is no pattern in valve size in relation to Holocene temperature conditions revealed. This implies that size clines in non-marine ostracodes or at least (temperate) *Cyprideis* are probably not (directly) controlled by temperature. Another objective of this study was the question how species and latitude-specific size-latitude trends in non-marine ostracodes are expressed. A similar data set that enables the comparison of *C. torosa* with another species is available from the Neotropical ostracode species *Cytheridella ilosvayi* (Wrozyna et al., 2016, 2018, 2019). Both species belong to the same superfamily Cytheroidea, which is characterized by a great variation in size and valve shape and comprises marine, brackish, and freshwater species (Meisch, 2000). *Cytheridella ilosvayi* is a common freshwater species occurring in a variety of aquatic habitats in the entire Neotropical range. The size data (length, height) of *C. ilosvayi* were derived from modern populations sampled between 30°S and 26°N.

Despite relatively large gaps in the size data of *Cytheridella*, no or only a weak relationship between valve size and latitude is displayed by the available data (correlation coefficients of r < 0.3) (**Figures 5A,B** and **Table 4**). Assuming this as a general trend for the entire geographical range, it implies that tropical non-marine ostracodes might not develop latitudinal-size clines. However, more data from underrepresented regions and other species are needed to verify this hypothesis.

An explanation why Bergmann trends are possibly not exhibited by *Cytheridella* could be the missing or low temperature gradient and stable resources within its tropical to subtropical geographical range as it is suggested for some tropical vertebrates which provide little support for Bergmann trends (Reed, 2003; Read et al., 2018).





The preceding discussion has shown that existing models on causes and mechanisms of Bergmann trends only insufficiently explain latitude-size relationship of C. torosa. Actually, it is accepted that intra- and interspecific gradients in body size may represent different phenomena (Blackburn et al., 1999). Although Bergmann's rule was originally defined inter-specifically, it has been commonly applied to populations within a single species. The expected mechanisms involved in size-latitude trends, are primarily limited to intraspecific scales and no general mechanism accounting for interspecific size-latitude patterns has been suggested (Berke et al., 2013). Thus, the observed size clines in C. torosa may result from interspecific divergence (i.e., sizeordered spatial sorting; Muñoz et al., 2014). Approaches that parse out intra- and interspecific variations on size-environment relationships revealed that patterns apparent at the level of clades may be (almost) absent at the level of species, and vice versa. This could be also displayed by C. torosa where the strong correlation with latitude is displayed only when size data are used undifferentiated and are strongly diminished for both morphotypes (Table 4).

Meiri and Thomas (2007) recommended to consider phylogenetic structures within taxa in order to elucidate whether patterns in the geography of size result from turnover within or between taxonomic levels. From a growing number of organism groups inconsistent (interspecific) latitudinal-size clines are reported suggesting that there may be no universal interspecific trend in latitudinal variation of body size, even among closely related groups (Pallarés et al., 2019). This is reflected by *Cyprideis* and *Cytheridella*. Accordingly, their geographical body size patterns are neither straightforward nor universally transferable to ostracodes in general. The consideration of different phylogenetic scales and relationships is therefore important before any reliable conclusion on size-environment relationships can be drawn.

Control of Salinity on *Cyprideis torosa* Size

Boomer et al. (2017) investigated valve size of C. torosa in relation to salinity. They identified a two-fold pattern in which valve size is largest and relatively constant at salinities between 0.7 and 8% and gradually decreases under higher salinities. The present study not only reveals an opposite trend to that reported by Boomer et al. (2017) with smallest valves at lower and lowest salinities, but also provides implications for the scale dependency of the salinity influence. As clearly shown by the RDA (Figure 6), latitude contributes most explanatory power on the overall size variability of C. torosa superimposing the salinity control. This agrees well with observations that community patterns and intraspecific trait variability depend on the spatial and ecological scales of investigations (Evangelista et al., 2019), and are often structured by a hierarchy of environmental factors (Hirschfeld et al., 2021). Also, our data underline the importance of the taxonomic resolution (e.g., morphotypes) for inferences on the sensitivity of a species toward environmental (i.e., salinity) conditions. Disregarding the morphotypes and supposing all occurrences as one Cyprideis species no relationship between

size data and salinity is exhibited. In contrast, considering the morphotypes separately, significant (morphotype-specific) positive correlations between valve size (length and height) and salinity are revealed. Moreover, only morphotype II shows a weak correlation between width and salinity, while there is no correlation displayed by the other morphotype. These slight but obvious differences point to an ecological differentiation of the morphotypes possibly representing another indication for the presence of at least two species currently considered as *C. torosa*.

Thus, our investigations suggest that salinity is not a primary control on the size of *C. torosa* sensu lato as it is also assumed for other *Cyprideis* species (Gitter et al., 2015; Meyer et al., 2017). The discrepancies in salinity-size relationships of Boomer et al. (2017) and our data might result from integration of several unperceived *Cyprideis* species.

Nonetheless, considering the geographical distribution of the size data set of Boomer et al. (2017) the same latitudinal size trend as in our data is reflected and even extends the latitudinal size cline of *Cyprideis* (Figure 4).

Salinity values used for the investigations of salinitysize relationships were derived from different sources. Paleosalinity of the fossil occurrences of *C. torosa* for most sites was inferred by the application of ostracode-based transfer functions. These are based on the significant and independent contribution of electrical conductivity (i.e., salinity) to the total variance in ostracode species distribution and provide very reliable salinity estimates (Mischke et al., 2007, 2010, 2014). Although there have been probably no fundamental changes in salinity, sites where salinity was estimated due to similarity of fossil and Recent ostracode species distribution (Hersek lagoon), can provide only rough estimates of salinity. Verification of our findings and interpretations should be based on investigations of living specimens and *in situ* measurements of salinity conditions.

CONCLUSION

In conclusion, the investigation of distant Recent and Holocene populations of *C. torosa* shows that there is a large variation in valve-size traits (length, height, width). Distinct local size clusters are superimposed by two large-scale geographical patterns. One represents a differentiation in two significantly different size classes, which separates southern populations from northern counterparts at around ~42°N. The second large-scale pattern is the continuous increase of valve size of *C. torosa* with latitude according to a Bergmann trend. Contrary, the available size data of the tropical species *Cytheridella ilosvayi* do not display this trend.

The co-occurrence of both size morphotypes of *C. torosa* in a wetland close to the Dead Sea (Enot Zuqim) and in the Turkish Hersek lagoon precludes an environmental control on this geographical pattern. Additionally, size-salinity relationships are morphotype-specific. A possible explanation of this could be the existence of at least two different *Cyprideis* species.

Majority of explanations for Bergmann trends relate to ecological processes and are, thus, restricted to intraspecific

phylogenetic levels. These models only insufficiently contribute to the understanding of the observed size-cline of *C. torosa*. Our results imply that size of *Cyprideis* is predominantly controlled by phylogeny, rather than environmental factors (e.g., salinity).

By comparing both species (*Cyprideis torosa* and *Cytheridella ilosvayi*) it is indicated that size-clines in ostracodes might be taxon-specific and no general pattern or underlying causes for ostracodes exist. Therefore, it is important to consider phylogenetic relationships and possible existence of more species than anticipated before attempting to explain size variability solely by environmental influences.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author.

AUTHOR CONTRIBUTIONS

CW and WP designed and conceived the study and wrote the manuscript. SM, MG, and MH provided sample material and additional environmental data. MG and SM edited the manuscript. All authors commented and agreed on the final version of the manuscript.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fevo.2022. 857499/full#supplementary-material

Supplementary Figure 1 | External views of female and male left and right valves of *Cyprideis torosa* from the studied localities. YDK, Tarim Basin, China; XB, Lake Bosten, China; MS, Mansfeld Lakes, Germany; SOG, Sea of Galilee, Israel; ISR, Dead Sea, Israel; BKA, Lake Balkhash, Kazakhstan; BS, Danube Delta, Romania; HK, Hersek Iagoon, Turkey.

Supplementary Figure 2 | Variability of valve length (a) and width (b) of populations. Displayed are variability ranges of female valves. Boxplots of male valves are provided in the Supplementary Figure 3. Numbers indicate sample sizes. Population abbreviations are: YDK, Tarim Basin, China; XB, Lake Bosten, China; BKA, Lake Balkhash, Kazakhstan; ISR, Dead Sea, Israel; SOG, Sea of Galilee, Israel; BS, Danube Delta, Romania; HK, Hersek Iagoon, Turkey; MS, Mansfeld Lakes, Germany. Details about the localities are provided in **Table 1**. Tables display the pairwise comparisons of the *post hoc* test results of the PerMANOVAs for each data set [female right (RV; c) and left (LV; d) valves, and male right (e) and left valves (f).

Supplementary Figure 3 | Variability of valve length (a) and width (b) of populations. Displayed are variability ranges of male valves. Numbers indicate sample sizes. Population abbreviations are: YDK, Tarim Basin, China; XB, Lake Bosten, China; BKA, Lake Balkhash, Kazakhstan; ISR, Dead Sea, Israel; SOG, Sea of Galilee, Israel; BS, Danube Delta, Romania; HK, Hersek Iagoon, Turkey; MS, Mansfeld Lakes, Germany. Details about the localities are provided in Table 1.

Supplementary Table 1 | Size data of female left valves. Salinity data referred to the original publications. See Table 1 for details.

Supplementary Table 2 | Size data of female right valves. Salinity data referred to the original publications. See **Table 1** for details.

Supplementary Table 3 | Size data of male left valves. Salinity data referred to the original publications. See for Table 1 details.

Supplementary Table 4 | Size data of male right valves. Salinity data referred to the original publications. See **Table 1** for details.

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