



Reconstructing Holocene hydroclimate variability and coastal dynamics of the Nile Delta: A diatom perspective

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ABSTRACT

This study analyses diatom assemblages from a Nile Delta core (B-1) to probe Holocene hydroclimate changes and their influence on the ecological habitats of the delta coast, with a further focus on the effects of relative sea-level rise. We found that the freshwater diatom *Aulacoseira granulata* varied in tandem with hydroclimate pulses in the Nile watershed, driven by the shift of the Intertropical Convergence Zone (ITCZ), therefore serving as a proxy for palaeo-Nile flow. Based on the ecological affinities of diatom taxa, we defined 5 diatom assemblages (I-V). Assemblage I (>10.0-8.5 ka BP) shows high *A. granulata* abundance (60–80%), implying peak Nile flow during the African Humid Period (AHP), at least 3 times greater than that of the recent past. Assemblage II (8.5-7.5 ka BP) sees decreased *A. granulata* (20–40%) and emergence of freshwater diatoms with benthic-oligotrophic characteristics (e.g. *Epithemia gibba*), suggesting reduced Nile flow and a delta estuary where the habitat became shallower with lower nutrient content. Assemblage III (7.5-6.0 ka BP) shows *A. granulata* resurgence (50–80%), reflecting Nile hydroclimate variability post-AHP due to the migration of the Intertropical Convergence Zone (ITCZ). The high-resolution diatom spectra of B-1 also revealed major habitat changes, from saline to fluvial-dominated environments, around 6.0 ka BP. Assemblage IV (6.0-3.2 ka BP) indicates a notable Nile flow decline and freshwater community proliferation, coinciding with coastal habitat expansion and delta progradation due to RSL stabilisation and basin-wide aridification. Assemblage V (3.2-2.0 ka BP) is marked by drought-tolerant diatoms (*Nitzschia amphibia*, *Hantzschia amphioxys* and *Cavinula cocconeiformis*), indicating intensified aridification.

1. Introduction

In the context of investigating human occupation and basin-wide hydroclimate changes due to relative sea-level rise, the Nile Delta and its associated sediment archives are of significant scientific interest (Fig. 1). During the Holocene, the region is characterised by not only the arid climatic conditions prevalent in the lower basin but also by significant hydrodynamic changes observed in the upper basin, particularly in the context of interactions between the Indian monsoon, the African monsoon and the Congo Air Boundary (CAB) (Butzer, 1976; Said, 1981). Huge amounts of water and sediment have been washed downstream to

feed the delta coast, mainly from the African Highlands and the Ethiopian Plateau where heavy rainfall attained ca. 1400–2000 mm per year (Vizy and Cook, 2003; Slingso et al., 2005; Wolman and Giegengack, 2007).

In the early Holocene, the northward migration of the Intertropical Convergence Zone (ITCZ) caused a prolonged humid period over North Africa, known as the African Humid Period (AHP, generally, ca. 14.5–5.0 ka BP) (Woodward et al., 2007; Macklin et al., 2012; Shanahan et al., 2015; Tierney et al., 2017; Pausata et al., 2020). During this period, the monsoon rain belt expanded northwards turning the Sahara into the 'Green Sahara' and creating conditions that supported early

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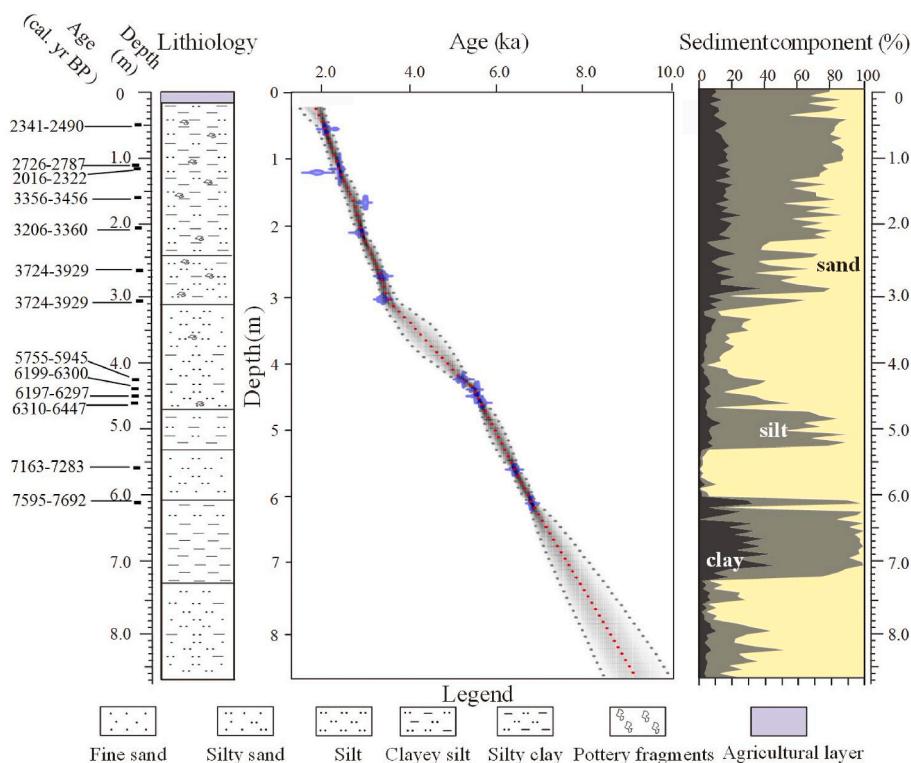


Fig. 2. AMS ^{14}C -dated Holocene strata of core B-1 and associated sediment texture. The age model is based on 13 dates.

high sedimentation rates during the Holocene have been documented (Giaime et al., 2022).

Core samples were collected continuously using a COBRA vibrator equipped with 2.0-m long tubes with a diameter of 9.0 cm. Core samples were stored in wooden boxes and transported to the laboratory for further analysis. The core was split into two halves, one for logging and the other for archiving. Core sediment was described including sediment colour, water content, lithology and biogenesis. Core sediment was then sampled at regular 5-cm intervals and dried via lyophilisation before analysis. A thin agricultural layer (0–40 cm; Fig. 2) at the top of the core was not considered for diatom analysis to avoid impacts from recent human activities. Also, we reasoned that the linearly-dated Holocene sediments of B-1 (Fig. 2) provide a reliable decadal-to-centennial-scale record of diatom communities, even though the influence of taphonomic processes along the delta coast cannot be entirely excluded (Espinosa et al., 2022).

2.2. Radiocarbon dating, age-depth model, and Holocene sea-level reconstruction

In total, thirteen sediment samples (12 organic muds and one shell fragment) from B-1 were sent to Beta Analytic in Florida, USA for AMS dating (Table 1). The ^{14}C dates were calibrated into calendar years before present (cal. yr BP) at confidence intervals of 95.4% (2σ), using the "rBacon" package for (Blaauw and Christen, 2011) and the IntCal 20 curve and Marine 20 curve (Blaauw and Christen, 2011; Heaton et al., 2020; Reimer et al., 2020) and a marine reservoir age of $\Delta R = 400 \pm 79$ yr. Age-depth calculations for all dates, expressed as "ka BP", were performed using the method described by Blaauw and Christen (2011). The Holocene sea-level curve of the study area was reconstructed using 20 dated lagoon-peats, including 3 from B-1, 4 from BR-1 (Fig. 3a) (Giaime et al., 2022) and 13 from Arbouille and Stanley (1991). The topographic elevation of the core sites was benchmarked relative to msl (see Giaime et al., 2022 for detailed methodology). We utilised the Errors-In-Variables Integrated Gaussian Process (EIV-IGP) model (Cahill et al., 2016) to calculate rates of Relative Sea Level (RSL) change from

Table 1
Radiocarbon dates of core B-1.

Core name	Depth (m)	Laboratory No.	Materials	Conventional age (yr BP)	Calibrated age (cal. yr BP)	$\delta^{13}\text{C}$ (‰)
B-1	0.50	Beta-571272	Organic mud sediment	2380 ± 30 BP	2341–2490	−21.7
B-1	1.10	Beta-571274	Organic mud sediment	2360 ± 30 BP	2726–2787	−21.9
B-1	1.15	Beta-600306	Shell fragment	2650 ± 30 BP	2016–2322	−11.9
B-1	1.60	Beta-560433	Organic mud sediment	3180 ± 30 BP	3356–3456	−18.5
B-1	2.05	Beta-571275	Organic mud sediment	3060 ± 30 BP	3206–3360	−21.3
B-1	2.70	Beta-560434	Organic mud sediment	3560 ± 30 BP	3724–3929	−19.2
B-1	3.05	Beta-571276	Organic mud sediment	3560 ± 30 BP	3724–3929	−17.8
B-1	4.25	Beta-560435	Organic mud sediment	5140 ± 30 BP	5755–5945	−22.6
B-1	4.40	Beta-600306	Organic mud sediment	5450 ± 30 BP	6199–6300	−17.0
B-1	4.50	Beta-600307	Organic mud sediment	5440 ± 30 BP	6197–6297	−18.4
B-1	4.60	Beta-571277	Organic mud sediment	5610 ± 30 BP	6310–6447	−19.7
B-1	5.60	Beta-600986	Organic mud sediment	6310 ± 30 BP	7163–7283	−20.4
B-1	6.10	Beta-548856	Organic mud sediment	6820 ± 30 BP	7595–7692	−16.4

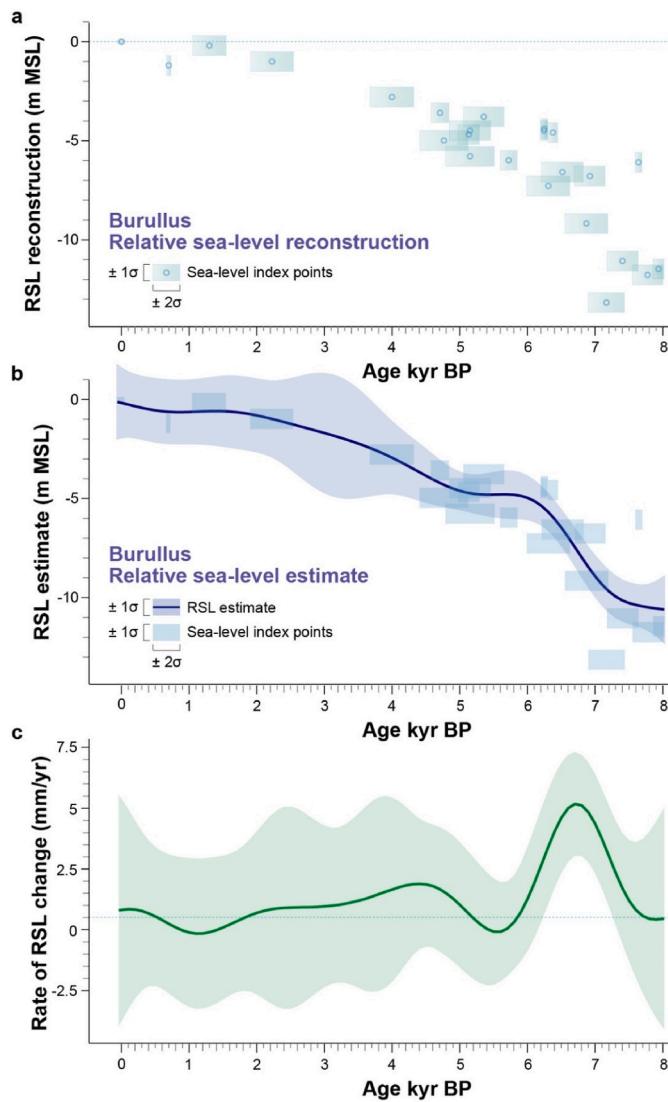


Fig. 3. Standardised RSL reconstruction for the Buto coast from 8000 ka BP to present. a) The RSL history is based on 20 SLIPs. The blue boxes represent the SLIPs from the lagoons and salt marshes. The dimensions of the boxes denote the $\pm 1\sigma$ altitudinal and $\pm 2\sigma$ chronological errors associated with each SLIP. b) Errors-In-Variables Integrated Gaussian Process model (EIV-IGP; Cahill et al., 2016) for RSL estimates (with the 1σ altitudinal error). c) Holocene RSL rising rates denoted in mm/yr.

Sea-Level Index Points (SLIPs) (Fig. 3b), applying a local Glacial Isostatic Adjustment (GIA) correction of -0.0512 mm^{-1} (Marriner et al., 2023). The EIV-IGP model employs a Gaussian Process prior on RSL rate estimates and integrates this process to derive estimations of the underlying RSL trend (Cahill et al., 2016). To tackle temporal uncertainty, this method is nested within an Errors-In-Variables framework. RSL uncertainties associated with the SLIPs are accommodated within the statistical model by introducing observation-specific error terms in the RSL data model. This modeling strategy offers a comprehensive assessment of uncertainty in both RSL estimates (Fig. 3) and the inferred rates of RSL change over time (Cahill et al., 2016).

2.3. Grain size

A total of 154 sediment samples was taken from B-1 for grain size analysis using ultrasonic methods (S Table 1). Approximately 0.1-g of pre-homogenized sediment was pretreated with a 30% H_2O_2 solution to oxidize organic matter and a 10% HCl solution to remove carbonates.

Before analysis, all of the sample was dispersed and homogenized using a solution of sodium hexametaphosphate. Grain size was measured using a Beckman Coulter LS13 320 Particle Size Analyzer.

2.4. Diatom analysis

Diatom fossil analysis was performed on 54 samples from B-1, selected at sampling intervals of 5–20 cm. Samples with few frustules were not taken into account (S Table 1). Diatom frustules were isolated using the flotation method and identified following the procedures outlined in Renberg (1990) and Battarbee et al. (2001). Diatom taxa were identified and counted under a microscope at magnifications of $400\times$ and $1000\times$.

In B-1, siliceous microfossils appeared well preserved: several fragile taxa including *Nitzschia* spp., *Tryblionella* spp. and *Pinnularia* spp. could be identified to species level with relatively intact frustules. *Aulacoseira granulata* could often be seen in chains with over two cells. In order to further verify the efficiency of diatom preservation in B-1, five random slides were examined to estimate the ratio of intact frustules at $400\times$ according to the method of McQuoid and Hobson (2001). The ratios of intact frustules on all the five slides were calculated to be over 60%.

Given the relatively constant sedimentation rates for core B-1, the absolute abundance (D_{abs} , valves/g DW) of diatom frustules can reflect the changes in diatom assemblages throughout the core sample. The relative abundance of each taxon (D_{re} , %) was calculated as a percentage (%) based on the total number of valves in each sample. Planktonic and benthic taxa were separately summed to show their temporal change along with other proxies.

Following Muylaert et al. (2009) and Baselga (2010), alpha diversity (D_α) and beta diversity (β_{sim}) were also computed to further determine the gradual compositional turnovers of assemblages in chronological order.

$$D_\alpha = 1 - \sum D_{re}^2 \quad (1)$$

$$\beta_{sim} = \frac{\min(b, c)}{a + \min(b, c)} \quad (2)$$

In the context of two adjacent samples, the variables a , b and c denote, respectively, the abundance of shared species, the abundance of species exclusive to one sample, and the abundance of species exclusive to the other.

Multiple statistical methods were applied to identify significant ecological changes in the diatom assemblages. Before testing, only taxa with a relative abundance of $\geq 0.5\%$ and appearing in no less than 3 samples were selected for the analysis. This was mainly to avoid the influence of rare species on the significance tests (Kato, 1993). D_{re} matrix data of diatoms were firstly run for their ecological affinity using the two-way indicator species analysis (TWINSpan) in WinTWINS Version 2.3 (Hill and Šmilauer, 2005), which helps classify different taxa groups according to their similarities. From this base, the diatom dataset was further analysed by CONISS using the software Tilia (Grimm, 1987) to establish a chrono-ecological zonation of B-1. The principal component analysis (PCA) in the CANOCO Version 4.5 (ter Braak and Šmilauer, 2002) was also applied to distinguish the typical species and potential controlling factors. In CANOCO, to further determine the waxing and waning of taxa in various time periods, the variations of taxa in chronological order were modeled using general additive modelling-GAM (Hastie and Tibshirani, 1990). Stepwise selection using the Akaike information criterion applied to the complexity control of the GAM model (Hastie and Tibshirani, 1990), and the maximum number of degrees of freedom for the smoother term was set to 2.

Sediment core BR-1 from our previous study, and further cores from the study area, were incorporated into our sedimentological database (Fig. 1) (Arbouille and Stanley, 1991; Giaime et al., 2021).

3. Results

3.1. Holocene strata and relative sea-level curve

Sediment core B-1 revealed a series of Holocene strata typical of lagoonal-coastal environments in the central Nile Delta (Fig. 2) (Stanley et al., 1996). The dark grayish lagoonal muds, rich in organic matter occurred in the Holocene sediments. The yellowish sand sections with brackish shell fragments and diatoms prevailed in the lower-middle part (below core depth of 3.8 m) of the sediment core, which was interlayered with lagoonal muds (Fig. 2). Silt and clay with brackish-freshwater shell fragments and diatoms gradually became predominant (>50%) in the upper core sediment (Fig. 2). A thin agricultural soil (0.15–0 m) occurred on the core top. The age-depth model revealed the full Holocene strata of B-1, corresponding to the core's bottom at 8.7 m depth (Fig. 2). Based on the age model, relatively stable sedimentation rates were observed, ranging between 0.6 and 1.2 mm/year (Fig. 2).

The reconstructed Holocene relative sea-level curve of the study area showed that RSL was 12.0 m below msl ca. 8.0 ka BP. RSL rose rapidly between 8.0 ka and 6.0 ka BP, before plateauing at around 4 m below MSL between 6.0 ka and 5.0 ka BP. RSL then gradually rose over the next 5000 years, reaching present levels (Fig. 3). The interaction between sea-level rise and paleo-Nile flow will be explored in the following sections, describing their collective impact on the ecological habitat of the Nile Delta.

3.2. Ecological affinity

A total of 27 taxa was used for multiple statistical analyses after removing the rare species (S. Table 1). TWINSpan was firstly applied with 9 levels (0, 1%, 2%, 5%, 10%, 15%, 20%, 40%, 60%) to classify taxa into 5 groups (Groups 1–5) according to their similarities (Fig. 4). These groups were differentiated on the basis of marine to fluvial habitats, to support the temporal zonation in CONISS (Fig. 5). Group 1 included mostly marine and brackish taxa. Group 2 consists primarily of brackish to freshwater taxa. Group 3 was dominated by freshwater taxa, and Groups 4 and 5 included freshwater and drought-tolerant species. It is worth noting that some species in Group 1 (including *Staurosira construens*, *Stephanodiscus hantzschii*, and *Campylodiscus clypeus*) and Groups 4 and 5 (including *Staurosira venter*, *Nitzschia umbonata*, *Surirella ovalis*, and *Nitzschia vitrea*) (Fig. 4) are referred to as marine-freshwater species because they have been found to live in both freshwater and seawater (https://www.algaebase.org/). In the present study, we attributed species in Group 1 to the marine domain and those in Groups 4–5 as having freshwater-brackish affinities based on the similarities to their contiguous taxa (Fig. 4).

3.3. Holocene diatom assemblages

Based on the characteristics of diatom assemblages identified in the CONISS analysis, there were 5 Holocene diatom assemblages (I-V) established in chronological order (older to younger) in B-1. Three sub-units (A-C) were assigned to assemblage IV (Fig. 5).

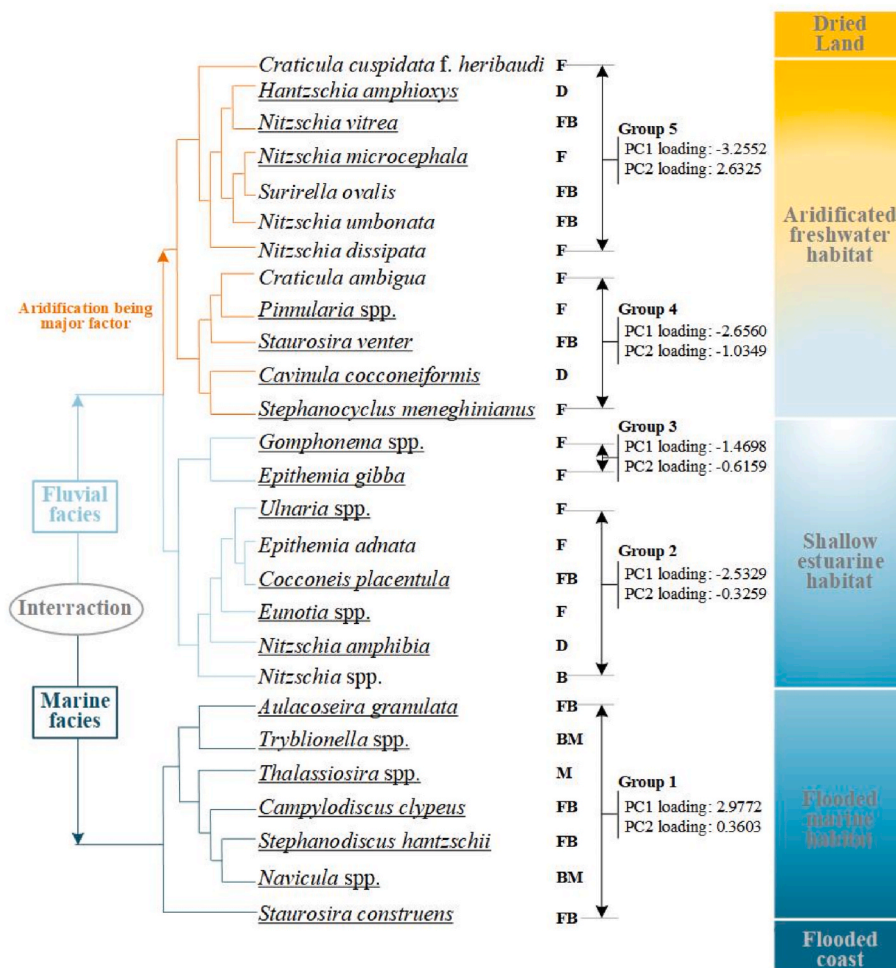


Fig. 4. Ecological affinities of diatom taxa from core B-1, based on TWINSpan analysis. F-freshwater taxa; D-drought tolerant species; FB- freshwater-brackish species; B-brackish taxa; BM-brackish and marine taxa; M-marine taxa.

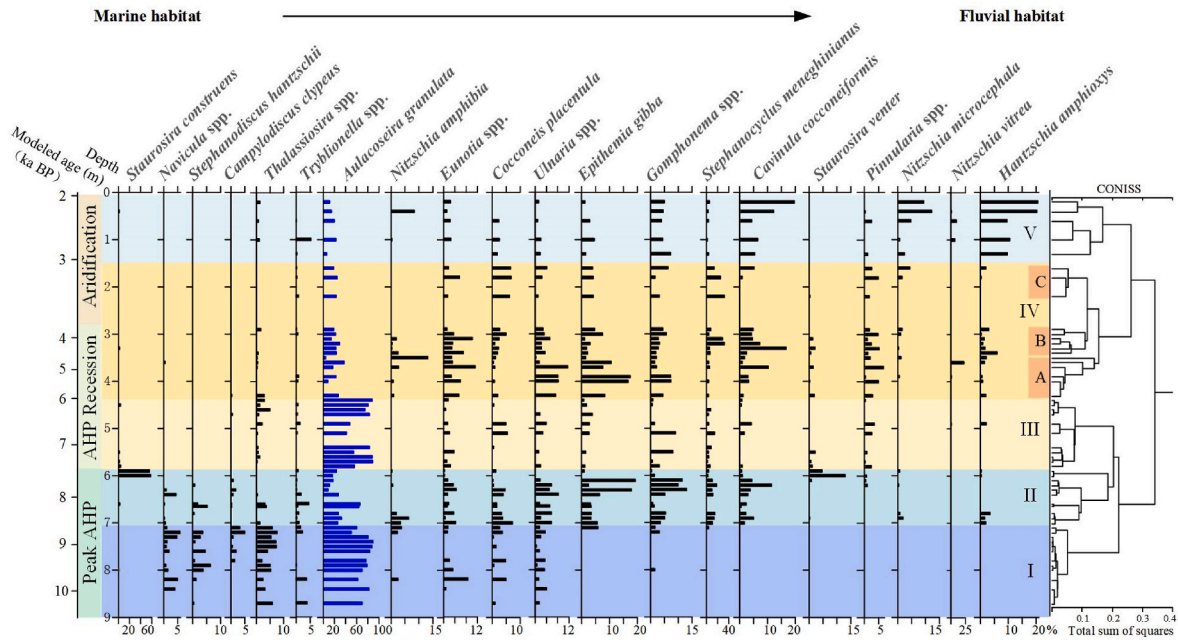


Fig. 5. CONISS-zoned diatom spectrum of B-1.

Assemblage I (below 7.0 m core depth, ca. >10.0-8.5 ka BP) was characterised by dominant fresh-brackish species *A. granulata*, as well as marine and brackish taxa, including *C. clypeus*, *Navicula* spp., *S. hantzschii*, *Thalassiosira* spp., and *Tryblionella* spp. Some low abundances of freshwater species such as *Ulnaria* spp. and *Eunotia* spp. were also observed (Fig. 5).

Assemblage II (7.0–5.9 m core depth, ca. 8.5-7.5 ka BP) witnessed a dramatic decrease in marine and brackish species, but an obvious increase in freshwater species in terms of both species' richness and abundance. Representative species included *E. gibba*, *S. meneghinianus*, *Gomphonema* spp., and *C. coccoeiformis* (Fig. 5).

Assemblage III (5.9–4.3 m core depth, ca. 7.5-6.0 ka BP) saw an increasing abundance of *A. granulata*. Comparatively, both marine and freshwater species became scarce (Fig. 5).

Assemblage IV (4.3–2.9 m core depth, 6.0-3.2 ka BP) persisted for a longer period, ca. 2000 years (Fig. 5). Freshwater species were dominant in both sub-units A and B, with a diverse range of species that alternately held the predominant position. When entering sub-unit C, the marine and brackish species almost disappeared, and freshwater-brackish taxa comprised only a small proportion with a low number of species, such as *A. granulata* and *C. placentula*. There was a hiatus of diatoms between B and C, lasting ca. 500 years (2.9–2.2 m core depth).

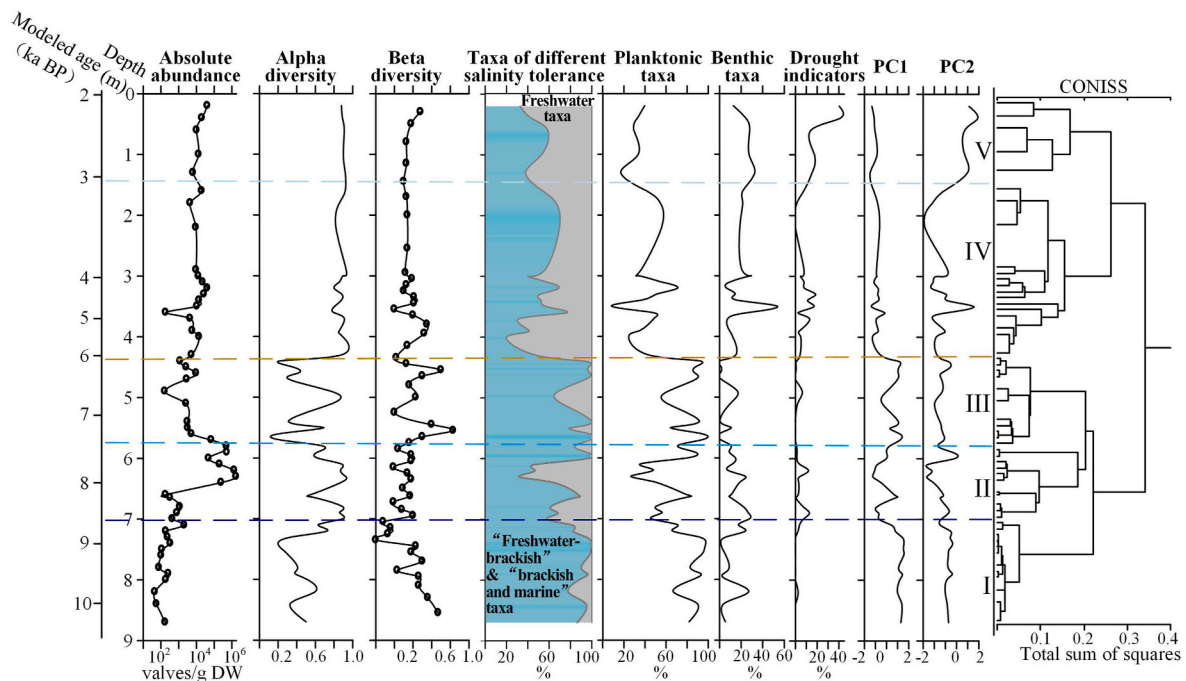


Fig. 6. CONISS diatom zonation of B-1, showing the temporal distribution of absolute abundances and species richness, Alpha/Beta diversities, freshwater and brackish-water sums and PC1 and PC2 (all discussed in the text).

Assemblage V (2.9–0.2 m core depth, 3.2–2.0 ka BP) showed an increasing dominance of freshwater species, such as *H. amphioxys* and *C. cocconeiformis*. Contemporaneously, a minor group of brackish and marine species coexisted (Fig. 5).

3.4. Distribution of diatom abundance and richness

The absolute abundance of diatom frustules ranged from 0.5×10^2 valves/g DW to 1.7×10^6 valves/g DW (Fig. 6), with the highest mean of 6.7×10^5 valves/g DW observed at a depth of 6.4–5.8 m (ca. 8.0–7.0 ka BP). By contrast, below the core depth of 6.6 m (ca. 8.0 ka BP), diatom frustule abundance was low, with a mean value of 4.0×10^2 valves/g DW.

There was a total of 149 diatom taxa identified, with the majority belonging to a wide variety of freshwater species (e.g. *Stephanocyclus meneghinianus*, *Cavinula cocconeiformis*, and *Epithemia gibba*) (S. Table 1 and Figs. 5 and 6). Due to the varying distribution of the predominant species *A. granulata*, accounting for >60% of the total, the richness pattern exhibited notable fluctuations (Figs. 5 and 6). Relatively low values, not exceeding 2.5, were mainly seen at core depths below 7.3 m, 5.4–6.1 m, and 4.4–4.7 m, corresponding to ca. 8.9 ka BP, ca. 7.7–7.1 ka BP, and ca. 6.5–6.1 ka BP, respectively. After ca. 6.0 ka BP (4.3 m core depth), the species composition of the diatom assemblage remained diverse. Variations in the alpha diversity were consistent with changes in the abundance of *A. granulata* (Fig. 6). According to pairwise dissimilarities detected by the beta diversity, when β_{sim} is closer to 1, the dissimilarity between the two assemblages becomes greater (Baselga, 2010; Socolar et al., 2016). Therefore, the apparent turnover reflected in B-1 appeared around 6.0 ka BP, when β_{sim} was >0.6. Also, planktonic communities remained remarkably high before ca. 6.0 ka BP, contrasting with a discernible moderate increase in benthic groups subsequent to this period (Fig. 6).

3.5. Classification of diatom taxa

A PCA was applied to the diatom dataset to further refine taxonomic classification and discern the primary modes of assemblage shift. The first axis (PCA1), accounting for 41.45% of the total variation, was primarily influenced by *A. granulata* and *S. hantzschii* with high positive loading, and *C. cocconeiformis* with high negative loading (Fig. 7). The TWINSpan groups were also differentiated from each other in the PCA,

reflected in their loadings on PC 1 and PC 2 (Fig. 6). Group 1, including most marine and brackish taxa, were loaded on the positive PC 1 axis (2.9772) while all the other groups, mainly with the freshwater communities, on the negative PC 1 axis (Figs. 4, 6 and 7). The majority of taxa within Groups 2–4 exhibit negative scores on both PC 1 and PC 2, placing them within the third quadrant of the PCA bi-plot. Conversely, Group 5 is distinguished by the highest positive loading on PC 2 (2.6325), concurrently featuring all drought-tolerant species (*C. cocconeiformis*, *N. amphibia*, and *H. amphioxys*) within the second quadrant (Figs. 4, 6 and 7).

The GAM response curves organised chronologically, delineated the rise and decline of taxa into three distinct types: geometric growth (Fig. 8a), gradient descent (Fig. 8b) and normal curve (Fig. 8c). The response curves indicated that the diatom assemblages were composed of dominant taxa with their population optimum in different periods. In the group of normal curves, the population peak of a particular taxon appeared either before or after 6.0 ka BP (Fig. 8c).

3.6. Diatom indicative species and their ecological habitats

3.6.1. Marine and brackish taxa

The diatom assemblages in core B-1 consist of brackish and marine taxa adapted to saline water, most of which were classified into Group 1 and, to a lesser extent, Group 2 (Fig. 4). Relative dominance is attributable to *Thalassiosira* spp., including *T. decipiens* and *T. symmetrica*, both oceanic and coastal planktonic species (Zalat and Vildary, 2007; Hammad and Ibrahim, 2012). The marine species *S. hantzschii* and *C. clypeus* are recorded in thalassic environments of the Nile lagoonal area (Zalat and Vildary, 2007). The former prefers relatively low temperature (Shafik et al., 1997) and the latter is adapted to living in shallow water as a large benthic species (Pankow, 1976). Other dominant saline taxa are *Navicula* spp. (such as *N. Lyrata* and *N. aegyptiaca*) and *Tryblionella* spp. (such as *T. scalaris*, *T. granulata*, and *T. punctata*), both commonly found in coastal lagoons and estuaries (Cholnoky, 1968; Foged, 1979; Zalat and Vildary, 2007).

3.6.2. Freshwater-brackish taxa

There are nine species in this group, mostly assigned to Group 5 (Fig. 4). Such species frequently live in freshwater (Fig. 4), although they are also found in brackish-water settings, such as estuaries, lagoons and coastal waters (Hällfors, 2004; Hassan et al., 2005; Zong et al., 2010;

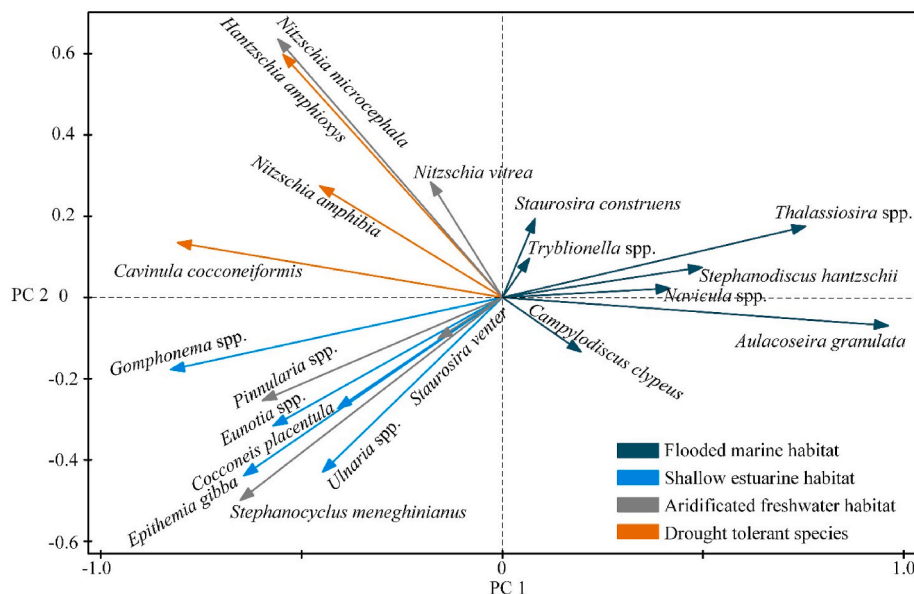


Fig. 7. PCA analysis of B-1.

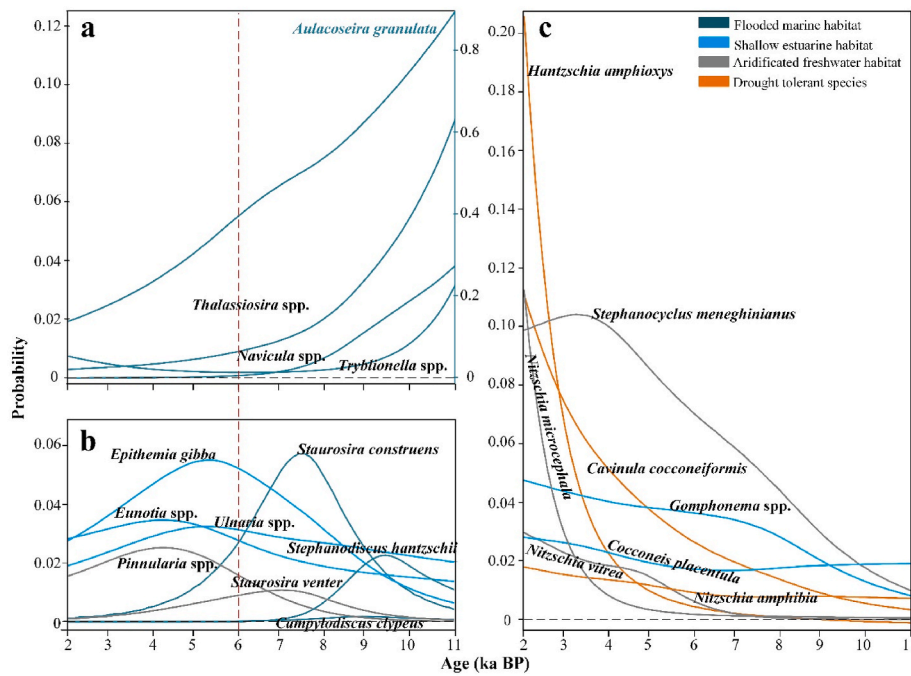


Fig. 8. GAM analysis of diatom taxa of B-1 and distribution patterns: a) descending; b) normal; c) ascending.

Lazrak et al., 2022). The representative species is *A. granulata*, predominant in B-1 assemblages (Figs. 4 and 5). Its relatively high dominance can be attributed to its competitive advantage in eutrophic to super-eutrophic waters. Therefore, *A. granulata* was widely used as a sensitive indicator of water areas of eutrophication and high productivity (Bicudo et al., 2016; Faustino et al., 2016). Particularly, *A. granulata* is identified as a characteristic Nile River diatom (Zalat, 2000), observed from upstream areas of both the Blue Nile and the White Nile, extending downstream to the estuary (Talling et al., 2009; Stager and Johnson, 2008; Hammad and Ibrahim, 2012). *A. granulata* frequently forms blooms in eutrophic waters (Kilham and Kilham, 1975; Campbell et al., 2006; Faustino et al., 2016).

3.6.3. Freshwater taxa

Throughout B-1, freshwater taxa from Groups 2–5 dominate (Fig. 4). The dominant freshwater species (Figs. 4 and 5) are characteristic of lakes, rivers and reservoirs, typically thriving in oligohalobous conditions, demonstrating tolerance to small amounts of salt. Amongst them, the genera *Pinnularia*, *Ulnaria*, *Eunotia* and *Gomphonema* exclusively comprise freshwater species (Foged, 1979; Van Dam et al., 1994). The dominant freshwater species, namely, *N. amphibia*, *H. amphioxys* and *C. cocconeiformis* are also recognised as drought indicators, since they dwell in either sand-soil or aerial and aeroterrestrial habitats (Van De Vijver et al., 2003; Zalat and Vildary, 2007; Cvetkoska et al., 2014; Maltsev et al., 2021). Meanwhile, primarily benthic and epiphytic, these freshwater taxa exhibit a preference for shallow environments owing to their phototropic response (Van Dam et al., 1994; Zalat and Vildary, 2007).

4. Discussion

Throughout the Holocene, the coastline of the north-central Nile Delta underwent significant changes, closely tied to the evolution of palaeo-Nile branches and the dynamic interplay between land and sea interactions (Fig. 1). The environment of B-1, further inland than the Burullus lagoon, was relatively stable, protected by the formation of a lagoon-barrier system on the seaward side (see Br-1 in Giaime et al., 2022) (Fig. 1), evidenced by its relatively stable sedimentation rate

throughout the Holocene (Fig. 2). The diatom record of B-1 enabled us to reconstruct the Holocene hydroclimate changes, particularly in relation to marine invasion events (Figs. 3–5). Diatoms, being sensitive indicators of various environmental variables, offer valuable insights into these fluctuations (Hall and Smol, 1999; Bigler et al., 2003; Zalat and Vildary, 2007; Zalat, 2015).

4.1. Reconstructing Holocene hydroclimate-marine fluctuations

4.1.1. >10.0–8.5 ka BP: highest Nile flow and a diluted estuarine habitat

During the peak of the AHP, the highest abundance of *A. granulata* within assemblage I (prior to ca. 8.5 ka BP) was observed across B-1, alongside a notable presence of brackish and marine taxa from Group 1 (including *Thalassiosira* spp. and *S. hantzschii* etc.) (Figs. 4 and 5), consistent with the influx of significant volumes of Nile freshwater reaching the coastal areas. This period also corresponds to rapid rates of relative sea-level rise occurring during the Early Holocene (Fig. 3) (Stanley and Warne, 1993b), as recorded by the majority of the marine diatom communities (Figs. 4 and 5). Overall, it implies a brackish estuarine setting (Figs. 4 and 5), but high abundances of *A. Granulata* suggest that the estuarine waters were significantly diluted by the substantial influx of Nile floodwaters rich in terrestrial nutrients. These conditions facilitated the formation of algal blooms in eutrophic environments (Faustino et al., 2016; Bicudo et al., 2016).

4.1.2. ca. 8.5–7.5 ka BP: brief waning of Nile flow and oligotrophic habitat

The sharp decline of *A. granulata* (assemblage II; ca. 8.5–7.5 ka BP) indicates the waning of Nile flow (Fig. 5), which was closely associated with the retreat of the Intertropical Convergence Zone (ITCZ) during the later AHP peak (Tierney et al., 2011; Shanahan et al., 2015; Castañeda et al., 2016; Liu et al., 2017). Meanwhile, the flourishing freshwater taxa belonging to Groups 2–4 and assemblage II (Figs. 4 and 5), including species such as *Eunotia* spp., *E. gibba* and *Gomphonema* spp., suggest that the estuarine habitat was characterised by shallower water settings and lower nutrient concentrations, indicative of oligotrophic conditions (Patrick and Reimer, 1975; Floener and Bothe, 1980; Van Dam et al., 1994; Zalat and Vildary, 2007; Pavlov and Levkov, 2013). In comparison, reduced brackish and marine taxa in assemblage II attest to less

marine influence on the Nile coast during this period, although rapid sea-level rise did not slow until ca. 5 ka BP (Fig. 3; Giaime et al., 2022). Between, 8.5–7.5 ka BP RSL was around 10 m below present MSL and RSL rising rates were relatively modest at around 0.5 mm/yr. As fluvial input continued, the abundance and diversity of freshwater species increased during the later stage of the AHP peak (Figs. 5 and 6), reflecting a shift in the dominant influence from marine to freshwater (Arbouille and Stanley, 1991; Giaime et al., 2022).

4.1.3. ca. 7.5–6.0 ka BP: unstable Nile flow and fluctuating salinity levels in the estuarine habitat

This period saw two major pulses of *A. granulata* in diatom assemblage III (Fig. 5), with a low occurrence between ca. 7.0–7.4 ka BP., reflecting unstable hydroclimate conditions in the upper Nile watershed following a further decline of the AHP (Tierney et al., 2011; Wang et al., 2022). Studies by Tierney et al. (2011) and Shanahan et al. (2015) found that higher Nile freshwater levels during this period were associated with the transition from the AHP peak to the early stages of the AHP recession as the ITCZ migrated southwards. This caused marked fluctuations in precipitation over the Ethiopian Highlands due to strong interaction between the African and Indian monsoons (Castañeda et al., 2016; Liu et al., 2017). Also, an eastward shift in the Congo Air Boundary (CAB) during this period intensified the unstable hydroclimate conditions over the eastern African plateau (Castañeda et al., 2016; Liu et al., 2017). In addition, the decline of freshwater diatom species (*E. gibba*, *Gomphonena* spp., and *Eunotia* spp., etc.) in Group 3 and assemblage III suggests a shallowing estuarine environment, where benthic communities faced increased difficulty in adapting. Several brackish diatoms disappeared as the delta coastline prograded seawards due to high fluvial sediment supply and the creation of new accommodation driven by RSL rise (Stanley and Warne, 1993a; Giaime et al., 2022, Fig. 3).

4.1.4. ca. 6.0–3.2 ka BP: delta progradation and a drying coastal habitat

After 6.0 ka BP, RSL stabilised with rising rates between 0 and 1.5 mm/yr during the following period (Fig. 3). The diatom assemblage IV saw a ca. 50% reduction in *A. granulata* compared to previous assemblages, but also marked the reappearance of freshwater communities from Groups 2–4 (Fig. 4). This was related to the drying climate of the Nile basin (Wang et al., 2022; Zhang et al., 2022), confirmed by the contrasting trends of *A. granulata* (increase) and *E. gibba* (decrease) during the late period of the AHP recession after approximately 6 ka BP (Zalat and Vildary, 2007). Also, the appearance of sporadic fresh-brackish taxa from Group 5 is consistent saline coastal habitats due to high evaporation under post-AHP drying climate conditions (Figs. 4 and 5). The discontinuous occurrence of diatoms probably translates an “extreme aridity” event around 3.2 ka BP (Fig. 5). As the ITCZ moved further southward, Nile floods decreased significantly (Fig. 9b–g).

4.1.5. ca. 3.2–2.0 ka BP: intensifying aridification

This period witnessed a marked increase in drought-indicative species (15–20% of total), mainly *N. amphibia*, *H. amphioxys*, and *C. cocconeiformis* of Group 5 and the assemblage V (Figs. 4 and 5), consistent with an intensified arid climate of the Nile River basin. Unlike before, the Buto coast was seldom flooded by Nile flow (Giaime et al., 2022; Wang et al., 2022). *H. amphioxys* and *C. cocconeiformis*, both characterised as being aerial diatoms (Cvetkoska et al., 2014; Jahn et al., 2014), flourished in the extremely dry environment of the Buto coast, with blooms of up to 20%. These diatoms are even capable of living in soil and beach sand, and, in a context of RSL stability, they record the environmental change from a former lagoon coast to a coastal plain (Giaime et al., 2022).

4.2. 6.0 ka BP: timing of saline to arid habitat transformation

Our PCA analysis of the diatom assemblages revealed that salinity

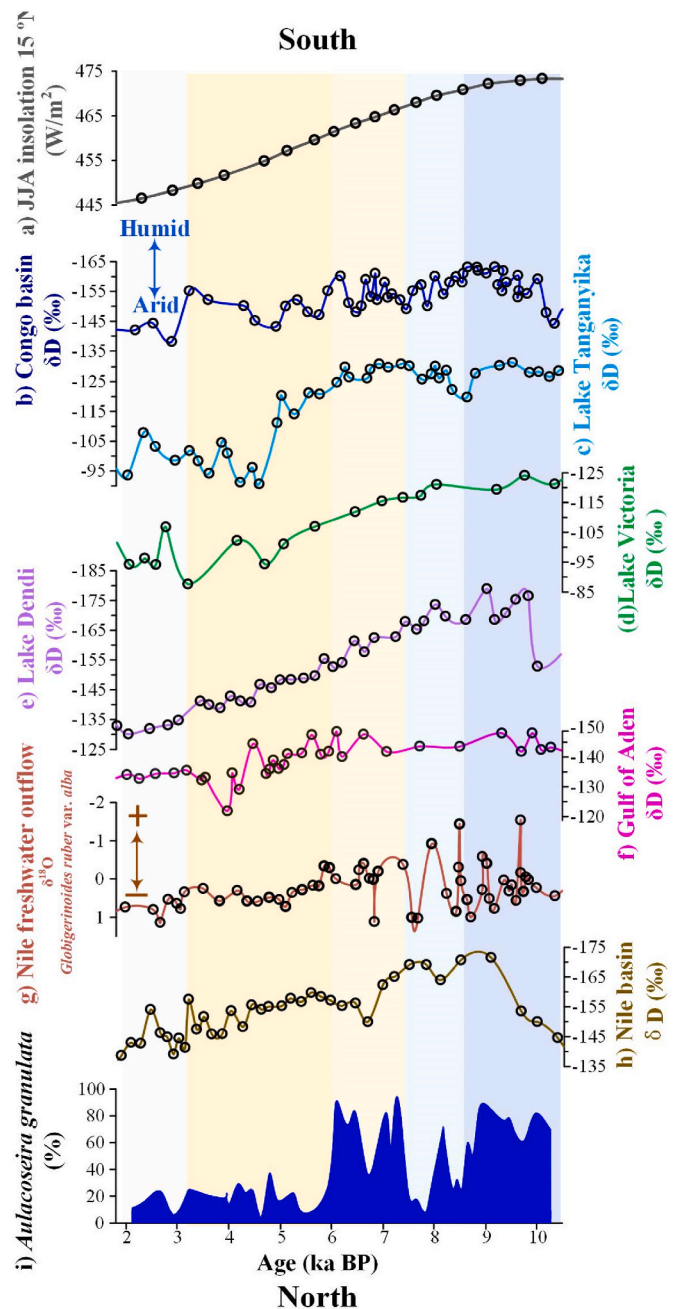


Fig. 9. Hydroclimate fluctuations of the Nile River basin. Comparison of the Nile flow indicator *Aulacoseira granulata* of B-1 with other basin-wide hydroclimate proxies: a) Summer insolation (June, July and August) at 25° N (Laskar et al., 2004); b) Congo Basin δD_{wax} (Schefuß et al., 2005); c) Lake Tanganyika δD_{wax} (Tierney et al., 2008); d) Lake Victoria δD_{wax} (Berke et al., 2012); e) δD values of the n-C29 from Lake Dendi sediment core (Jaeschke et al., 2020); f) Gulf of Aden δD_{wax} (Tierney and de Menocal, 2013); g) $\delta^{18}O$ data of the core MS27PT (Revel et al., 2015); h) Nile Basin δD_{wax} (Castañeda et al., 2016); i) Relative abundances of *Aulacoseira granulata* of B-1. δD values from the Nile and Congo River Basins were determined using C31 n-alkane (Schefuß et al., 2005; Castañeda et al., 2016), whereas δD measurements for the Gulf of Aden, Lake Tanganyika and Lake Victoria were conducted using C28 (Tierney et al., 2008; Berke et al., 2012; Tierney and de Menocal, 2013).

variations affected by Nile flow played a key role in controlling the eco-environmental evolution of the Buto coast throughout the Holocene. As shown in the coordinate graph (Figs. 4, 6 and 7), all “brackish and marine” taxa exhibited positive loading on PC1, while freshwater taxa were negatively loaded. The proportion of drought-tolerant species not

only consistently varied with PC2 (Figs. 4, 6 and 7), but also exhibited a positive loading on PC2 (Figs. 4, 6 and 7). Apparently, “freshwater-brackish” and “brackish and marine” taxa varied proportionally with freshwater taxa (Fig. 6). The transition from brackish-marine to freshwater dominance occurred ca. 6.0 ka BP (Figs. 5 and 6). There was an apparent increase in both the quantity and diversity of freshwater taxa after 6.0 ka BP (Figs. 5 and 6), indicating fluvial inputs into a dominant estuarine eco-setting (Arbouille and Stanley, 1991; Giaime et al., 2022). The oscillation between planktonic and benthic proportions (Fig. 6), further corroborates the dominance of estuarine habitats influenced by high Nile floods before 6.0 ka BP. Subsequently, a notable shift occurred whereby land surface processes began to exert a prevailing influence. In the meantime, increases in drought-tolerant species (*N. amphibia*, *H. amphioxys*, and *C. cocconeiformis*) demonstrates that the post-AHP drying hydroclimate prevailed throughout the river basin (Figs. 5–7) (Castañeda et al., 2016).

The importance of the temporal inflection point at ca. 6.0 ka BP was also expressed in the GAM and diversity analyses (Fig. 8a, b, c). The brackish and marine taxa of Group 1 (Fig. 4) exhibited a descending trend but eventually stabilised at low relative abundances after 6.0 ka BP (Fig. 8a), whereas the drought-indicative species displayed a notable increase after 6.0 ka BP. (Fig. 8c). In the type of normal distribution, all freshwater taxa within Groups 2–3 reached their peak abundance after 6.0 ka BP (Fig. 8b). Both the variation of D_{α} and β_{sim} also identified the significant transformation after 6.0 ka BP. From >10.0–6.0 ka BP, the diversity variation pattern remained consistent alongside changes in the proportion of taxa with varying salinity tolerance (Fig. 6). Since 6.0 ka BP, the persistent high values of D_{α} coupled with low β_{sim} values (Fig. 6) suggests that the freshwater environment had sufficiently stabilised, allowing the diatom community to maintain a consistently high and steady diversity.

It is worth mentioning that the timing of the habitat transformation ca. 6.0 ka BP in B-1 coincides with both RSL stability and with the establishment of the settlement of Buto, precisely after 6.0 ka BP. This chronological correlation underscores the significance of Buto settlement in the context of the predynastic Buto Culture of Egypt, serving as a crucial link between Upper and the Lower Egypt societies (Mączczyńska, 2014).

4.3. Diagnostic diatom taxa: implications for hydroclimate events

Diagnostic diatom communities were associated with specific hydroclimate events. Higher abundances of *S. meneghinianus* ca. 8.5 ka BP, 7.8 ka BP and 4.2 ka BP (Fig. 5) may also reflect an estuarine setting richer in nutrients, given its competitive edge in eutrophic waters (B-Béres et al., 2014; Fontana et al., 2014; Faustino et al., 2016). The remarkably high abundances of *E. gibba* and *Gomphonema* spp. ca. 7.8 ka BP and 6.4 ka BP (Fig. 5) point to more oligotrophic conditions (Zalat and El-Sheekh, 1999; Adler et al., 2014; Cvetkoska et al., 2014; Glushchenko and Kulikovskiyb, 2017).

S. construens and *S. venter* are small sized (3–20 μm) benthic/tycho-planktonic species (Morales, 2001), most common in coastal settings (De Wolf, 1982; Vos and De Wolf, 1993). The higher abundances of *S. construens* >50% ca. 7.6 ka BP (Fig. 5) probably indicates increasing marine processes in a context of high RSL rising rates (Vos and De Wolf, 1993; Hassan et al., 2003), when the Nile flow was lower (seeing lower *A. granulata* of Fig. 5). The high abundance of *S. construens* is likely a result of taphonomic processes rather than ecological attributes (Hassan et al., 2005). Its frustules are easily accumulated in high concentrations through tidal transport processes.

4.4. *Aulacoseira granulata*: Nile flow indicator

A. granulata is a widespread centric and planktonic diatom, blooming in freshwater lakes, reservoirs and rivers (Kilham and Kilham, 1975; Costa et al., 2009). It can also follow the outflow of freshwater into

brackish estuarine, lagoonal and coastal settings (Shen et al., 1995; Odebrecht et al., 2005; Hassan, 2010; Zong et al., 2010). Some studies have classified it as a brackish species (e.g. Huang et al., 2004; Zong et al., 2010). However, as a typical freshwater species only tolerant to low salinity, its abundance and distribution obviously displayed a negative correlation with salinity (Huang et al., 2004; Zong et al., 2010), but a positive correlation with runoff discharge (Costa et al., 2009). Previous studies have recognised *A. granulata* as a Nile flow indicator (Zalat, 2000; Williams et al., 2010), with a wide latitudinal range, from the Blue Nile and White Nile to downstream and estuarine areas (Stager and Johnson, 2008; Talling et al., 2009; Hammad and Ibrahim, 2012). Our analysis demonstrates that the Holocene abundances of *A. granulata* in B-1 are significantly correlated to the hydroclimate dynamics of the Nile basin (Fig. 9a–i). In light of these observations, we suggest that *A. granulata* serve as a palaeo-Nile flow proxy. Its proportion of total taxa ranging from 60 to 80% during the Early Holocene to ca. 20% during the Late Holocene, supporting the estimation that the palaeo-Nile flow ca. 8000 years ago was approximately three times greater than that of the recent past (Figs. 5 and 9i). A similar estimate of palaeo-Nile flow during the AHP can be also inferred from sedimentation patterns off the Nile coast (Vadsaria et al., 2019; Zaki et al., 2021; Blanchet et al., 2024), and the unique pollen species (Nile headwater indicators) derived from the upper Nile source (seeing Figs. 5 and 6 of Zhao et al., 2020).

5. Conclusions

This study has demonstrated that.

- (1) *Aulacoseira granulata* is a sensitive freshwater-brackish diatom species, which fluctuated in proportion to Nile flow throughout the Holocene, making it a reliable proxy for reconstructing past Nile flow.
- (2) The ^{14}C -dated diatom spectra show that 6.0 ka BP was a key change point, marking the transition from a Nile-influenced estuary to a freshwater-dominated setting, corresponding to the shift from the peak AHP to the subsequent aridification phase and a stabilisation of relative sea level.
- (3) 5 diatom assemblages were elucidated in this study. Assemblage I (ca. >10.0–8.5 ka BP) exhibited the highest abundance of *A. granulata* (80–60%) alongside marine taxa, indicating a diluted estuarine environment during the Holocene marine transgression and the peak of the AHP. The subsequent lowering of *A. granulata* (to 20–40%) in assemblage II (ca. 8.5–7.5 ka BP) saw an incipient decrease in Nile flow during the latter phases of the AHP. Assemblage III (ca. 7.5–6.0 ka BP) witnessed fluctuations in *A. granulata* (50–80%) and no marine taxa, reflecting unstable Nile flow caused by monsoon hydro-dynamics in eastern Africa in response to the southerly migration of the ITCZ. After 6.0 ka BP, Nile flow decreased further (*A. granulata* to 20–30%), leading to a more oligohalobous coastal habitat under the persistent drying climate (assemblage IV). The drought-tolerant diatom communities of assemblage V (3.2 ka BP onward) are consistent with basin-wide aridification.
- (4) Higher abundances of the drought-tolerant taxa *Nitzschia amphibia*, *Hantzschia amphioxys* and *Cavinula cocconeiformis* are consistent with climate aridity during both the late period of 4.0 ka BP and 3.0 ka BP. The sudden increase in abundance of small species such as *Staurosira construens* and *Staurosira venter* suggests intense tidal dynamics along the estuarine coast ca. 7.6 ka BP, a period coinciding with diminished Nile flow intensity. High abundances of *E. gibba* ca. 7.3–6.0 ka BP and 5.3 ka BP suggests that the estuarine habitat exhibited markedly oligohaline conditions during these periods.

Authors individual contributions

Y. Wang, X. Zhao, Z. Chen, A. Salem designed the project. Y. Wang, X. Zhao, Z. Chen, A. Salem., and J. Chen collected sediment sample; Y. Wang and J. Zhou identified diatoms. X. Zhao and Z. Chen worked for AMS-dating materials; Y. Wang, X. Zhao, N. Marriner, D. Kaniewski, and Z. Chen wrote the manuscript, and all individuals participated in discussion and agreed this submission.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.quascirev.2024.109070>.

Data availability

Data will be made available on request.

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