Millennial-scale changes in abundance of brachiopods in bathyal environments detected by postmortem age distributions in death assemblage (Bari Canyon, Adriatic Sea)



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Abstract: Inferring the composition of pre-Anthropocene baseline communities on the basis of death assemblages (DAs) preserved in a surface mixed layer requires discriminating among recently-dead shells sourced by living populations and older shells from extirpated populations. Here, we assess the distribution of postmortem ages in the DA formed by the brachiopod *Gryphus vitreus* at 580 m depth in the Bari Canyon (Adriatic Sea), with no individuals collected alive. The *Gryphus* DA exhibits millennial time averaging (inter-quartile range = 1250 years) and two modes in abundance at 500 and 1750 years BP. As high abundance of species in time-averaged DAs can reflect passive accumulation of shells sourced by populations with low standing population density, we reconstruct changes in annual density on the basis of the abundance maxima detected in the distribution of postmortem ages and on the basis of estimates of per-specimen disintegration rate. We find that adults (>20 mm) achieved densities of at least 10–20 individuals/m² (assuming lifespan is 10 years), and the pulses in abundance were thus associated with a high population density in the past, followed by the decline over the last few centuries. We infer that bathyal populations were volatile during the Late Holocene, with brachiopods sensitive to siltation that was induced by temporal changes in sediment dispersal into the Bari Canyon due to deforestation and climatic changes.

Supplementary material: Age data and R language scripts that replicate analyses are available at https://doi. org/10.6084/m9.figshare.c.6228410

Bathyal environments are increasingly affected by anthropogenic impacts induced by trawling (Rogers 1999; Roberts et al. 2000; Edinger and Sherwood 2012), litter in submarine canyons incising narrow shelves (Fabri et al. 2014; Fernandez-Arcaya et al. acidification (Turley et al. 2017), 2007: Ramirez-Llodra et al. 2011; Maier et al. 2012, 2013; Georgian et al. 2016; Baco et al. 2017), and by changes in seawater temperature and in the strength of the biological carbon pump (Morato et al. 2020; Chaikin et al. 2022). Understanding the volatility and resilience of bathyal ecosystems to such disturbances requires the assessment of their long-term dynamics during the Pleistocene and Holocene (Yasuhara et al. 2008; Thiagarajan et al. 2013). Sediment cores can be used to extract

ecosystem history from fossil assemblages (e.g. Moffitt et al. 2015). However, bathyal environments are characterized by slow sedimentation or complex sedimentation affected by condensation, redeposition and episodic burial (Middelburg et al. 1997; Tyson 2001). In such settings, skeletal remains accumulating over multiple millennia are mixed within the upper decimetres of seabeds where they are actively affected by taphonomic processes. Such surface death assemblages (DAs) thus do not have sufficient stratigraphic resolution to discriminate ecological history at decadal or centennial resolution. However, they still (1) provide information on time-averaged composition of baseline communities (Kidwell 2007; Jonkers et al. 2019; Rillo et al. 2019) and (2) if coupled with dating of individual shells,

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they can be used to restore fine-scale temporal resolution and track fluctuations in abundance of shell producers in the past at decadal (Albano et al. 2016), centennial (Tomašových et al. 2017), millennial (Carroll et al. 2003; Krause et al. 2010; Perry and Smithers 2011; Dexter et al. 2014; Tomašových et al. 2016), or multi-millennial scales (Thiagarajan et al. 2013: Margolin et al. 2014). Age dating of species that occur at high abundance in DAs, especially when shell producers are missing in living assemblages (LAs) collected in the same area, can detect whether the lack of living individuals of such species reflects (1) short-term, annual variability in abundance (Albano et al. 2016), (2) an unprecedented decline in production during the recent decades or centuries (potential signature of anthropogenic impacts, Kowalewski et al. 2000; Simões et al. 2009; Tomašových et al. 2019), (3) long-term variability driven by millennial-scale climatic changes unrelated to anthropogenic impacts (Ritter et al. 2017; Pratt et al. 2019; Vertino et al. 2019), or (4) other sources of mismatch related to transport, reworking, and differences in species preservation rates (e.g. Dominici and Zuschin 2005; Kidwell 2008).

Frequency distributions of postmortem ages of shells in DAs collected in the uppermost centimetres of a seabed are expected to be exponential or heavytailed rather than flat under steady input of shells as they are subjected to burial and disintegration in the taphonomic active zone (TAZ, Davies et al. 1989; Flessa et al. 1993; Olszewski 1999, 2004; Kosnik et al. 2013; Albano et al. 2020; Agiadi et al. 2022). However, internal modes or significant gaps in abundance of cohorts within such distributions of postmortem ages are diagnostic of significant fluctuations in abundance of populations over the duration of time averaging (Krause et al. 2010; Tomašových et al. 2016; Vertino et al. 2019). Although high time averaging of shells within a single stratigraphic increment or in a seabed averages out high-frequency fluctuations in assemblage composition and thus complicates palaeoecological inferences about short-term dynamics, age distributions of time-averaged populations reverse the loss of high-resolution information. When burial and disintegration rates of shells and lifespans of their producers can be approximated, the species abundances in time-averaged DAs can be unmixed and scaled down to the yearly ecological density (Kowalewski et al. 2000; Tomašových and Kidwell 2017), thus overcoming the gap between the nature of ecological and palaeoecological abundance data. For example, the distributions of postmortem shell ages can be used to detect whether dead-only species had very low standing densities at yearly scales and just accrued their high abundance in DAs in the course of extended time averaging or whether they

achieved high annual standing densities (Tomašových et al. 2017).

When sedimentation rates are very low, age distributions of shells in a surface mixed layer can track fluctuations over relatively long (millennial) time-scales if shell disintegration rates in the mixed layer are slow (either within the TAZ or within protected microenvironments when the TAZ is patchy). because fast disintegration (1) prohibits preservation of old cohorts in the TAZ close to the sedimentwater interface, and (2) pulls the postmortem age of modal classes towards the Recent (Tomašových et al. 2016). With increasing disintegration rate in the mixed layer, age structure of surface DAs thus increasingly deviates from the original time series in abundance (Tomašových et al. 2016). Very slow disintegration can lead to almost uniform distributions of postmortem ages under steady input of shells, but such conditions are unusual in present-day shelf environments with intense recycling of organic matter inducing carbonate deterioration (Walter et al. 1993), leading to strongly right-skewed age distributions in surface DAs (Kidwell et al. 2005; Kosnik et al. 2007, 2009; Tomašových et al. 2014; Olszewski and Kaufman 2015).

DAs, however, with very old, early Holocene or late Pleistocene aragonitic skeletal remains, stained by ferromanganese coatings and carrying diverse early-diagenetic signatures, frequently occur in mixed-bottom sediments or on firm- or hardgrounds in bathyal environments (Noé et al. 2006; Freiwald et al. 2009; Angeletti and Taviani 2011; Angeletti et al. 2015), indicating that the disintegration rate of skeletal remains in the mixed layer is slow in these settings. For instance, cold-water corals in the Mediterranean Sea (with unusually warm and saturated conditions relative to other deep-sea environments), but also in the NE Atlantic Ocean, are frequently preserved in surface DAs not associated with living populations. Their radiocarbon ages indicate distributional shifts and significant fluctuations in their abundance on the transition to the Holocene (Delibrias and Taviani 1984; Schröder-Ritzrau et al. 2005; Vertino et al. 2019).

Here, we focus on the frequency distribution of postmortem ages of a surface DA formed by the epifaunal brachiopod *Gryphus vitreus* in the middle bathyal (580 m) mixed-bottom habitats in the Bari Canyon of the Southern Adriatic Sea (Figs 1 & 2). This species presently forms dense aggregations with more than 10–100 living individuals/m² in the Western Mediterranean Sea (Emig 1985, 1989*a*; Cartes *et al.* 2009). Such densities of living populations have not been observed in the Eastern Mediterranean Sea. We identify the timing of past fluctuations in abundance of the brachiopod *Gryphus vitreus* in the southern Adriatic Sea and discuss how the effect of disintegration modifies the shape of frequency



Fig. 1. (a) Geographical map of the southern Adriatic Sea with the location of the Bari Canyon, with light-grey colours corresponding to shelf segments, and the insets showing the positions of (b) and Figure 2. (b) Geographical distribution of stations in the southern Adriatic Sea with (dark grey squares) and without (light grey circles) *Gryphus vitreus* collected alive, and samples with dead-only collected *G. vitreus* (black triangles). The station POS514-14 in the Bari Canyon is investigated in this study. In total, 10% of all sites compiled on the basis of these published data (15 out of 130 sites in total, at water depths >100 m) contained individuals of this species collected alive in the Southern Adriatic Pit. These occurrences are based on samples collected during the Pola expedition in 1894 (Sturany 1896), grabs and trawls in 1969–70 documented by Gamulin-Brida (1983) off the Montenegrin coast, trawls documented by Marano *et al.* (1989), trawling survey PIPETA 1 in 1982 (Šimunović 1997), grab samples collected by the Poseidon cruise in 2017 (POS514), and samples documented in d'Onghia *et al.* (2015). The coordinates of death assemblages with *G. vitreus* are based on Trincardi *et al.* (2011) and on two stations collected by Poseidon in 2017. LA, living assemblage.

distributions of postmortem ages in DAs relative to the original chronological changes in abundances.

Gryphus vitreus

Gryphus vitreus is a suspension-feeding brachiopod with a smooth, punctate shell, attached to rocks, gravel or skeletal debris with a pedicle (Fig. 3). It can become secondarily free-lying when its substrate degrades or collapses (Emig 1987). It reaches 20–40 mm length in the adult stage (Boullier *et al.* 1986), and possesses a cyrtomatodont hinge (characterized by interlocked teeth and sockets) that prevents disarticulation to some degree. In contrast to most rhynchonelliformean brachiopods with two

shell layers represented by the outer, very thin cryptocrystalline primary layer and the inner, thicker secondary layer consisting of fibres, *G. vitreus* possesses also a third, innermost prismatic tertiary layer that is mainly developed in the posterior and central segments of the valves (Gaspard 1989, 2011; Emig 1990; Gaspard *et al.* 2007; von Allmen *et al.* 2010; Romanin *et al.* 2018).

Gryphus vitreus is abundant in level-bottom benthic communities inhabiting detritic, mixed-bottom, shelf-edge or bathyal habitats of the Mediterranean Sea. In the Western Mediterranean Sea, this species occurs between 110 and 250 m at Corsica and Provence (Emig 1985, 1987, 1989a), between 110 and 330 m in the Menorca Channel (Grinyó *et al.* 2018), and between 600 and 2200 m in the



Fig. 2. The location of the sampling station POS514-14 (black circle) on the northern edge of the northern channel of the Bari Canyon in the southern Adriatic Sea (modified according to Trincardi *et al.* 2007*a*, *b*; Angeletti *et al.* 2020). The grey-shaded areas mark the location of epibenthic communities with corals, sponges and serpulids as mapped by Angeletti *et al.* (2020).

Balearic Sea (Cartes *et al.* 2009). In the Eastern Mediterranean Sea, live *G. vitreus* was reported from 497 to 790 m in the Santa Maria di Leuca deep-water cold-coral mound province in the Ionian Sea (Mastrototaro *et al.* 2010; Rosso *et al.* 2010), and is also known to occur in the Aegean and Levantine seas (Brunton 1988; Logan *et al.* 2002; Gönülal and Güresen 2014). Former dredging and trawling surveys in the southern Adriatic Sea documented several occurrences of live-collected *G. vitreus* at bathyal depths between 300 and 1140 m in the nineteenth century (Sturany 1896) and in the twentieth century (Gamulin-Brida 1983; Marano *et al.* 1989;



Fig. 3. Life position of *Gryphus vitreus* attached with the pedicle to a hard substrate (shell, gravel) in the adult stage (c. 3 cm in length), with dorsal and ventral adjustor muscles that can rotate and tilt shells with respect to its substrate. Source: adapted from Cox (1934) and Delance and Emig (2004)

Šimunović 1997; Fig. 1). However, several more recent surveys did not detect this species alive (Angeletti *et al.* 2014, 2020; D'Onghia *et al.* 2015).

Study area

Sedimentation in the SW Adriatic Sea is affected by thermohaline bottom currents of the North Adriatic Dense Water (NADW) that advect sediment and organic detritus downslope and interact with a complex topography of the Bari Canyon, leading to a mosaic of non-depositional and depositional conditions (Trincardi et al. 2007a, b; Verdicchio et al. 2007; Ridente et al. 2007). The high fluxes of suspended food particles delivered by the cascading currents of the NADW in the Bari Canyon System also support epibenthic communities (Turchetto et al. 2007; Tesi et al. 2008; Bo et al. 2012; Foglini et al. 2016). The slopes and flanks of the Bari Canyon are inhabited by cold-water colonial and solitary corals that form three-dimensional structures (up to 80 cm high) at 305-650 m (Taviani et al. 2016), dominated by Madrepora oculata (280-550 m, d'Onghia et al. 2015), frequently in association with serpulids (Sanfilippo et al. 2013) and sponges (at 400–500 m; Bo et al. 2012; Angeletti et al. 2020).

Sediments lining the Bari Canyon are formed by sandy muds and muddy sands that are interspersed with large-scale patches of bedrock, coral bioconstructions and patches of biogenic consolidated sediments (Prampolini *et al.* 2021). DAs on the NW slope of the Southern Adriatic shelf are formed by mixtures of autochthonous, relict and reworked Holocene and Pleistocene remains of corals, molluscs and foraminifers (Trincardi *et al.* 2007*a*; Freiwald *et al.* 2009; Tesi *et al.* 2017). Dead *Madrepora, Lophelia* and *Desmophyllum* corals occurring between *c.* 250– 585 m range in postmortem age between 230 and 5100 years BP (Taviani *et al.* 2019). The bottom

currents can generate sand waves and lead to reworking and downslope transport of inner-shelf foraminifers (Tesi et al. 2017). Molluscan DAs with high levels of bioeroded and encrusted skeletal remains that occur at 200-300 m are represented by Holocene species mixed with the glacial late-Pleistocene species (Panetta et al. 2013), including Pseudamussium septemradiatum, two specimens of which were dated to 19 000 + 370 and 15 350 + 250 cal. years BP (Colantoni 1975; Colantoni and Gallignani 1978; Taviani 1978). Dead shells of G. vitreus were also collected in the southern Adriatic Sea between 318 and 616 m (Trincardi et al. 2007a, 2011), and at two stations sampled during the Poseidon cruise in 2017. The station POS514-14 sampled in the Bari Canyon at 580 m depth is investigated here (Figs 1 & 2). Another station, POS514-26, sampled on the eastern margin of the Southern Adriatic Pit at 509 m also included dead shells of Gryphus (black triangles in Fig. 1).

Methods

The station POS514-14 sampled during Poseidon cruise No. 514 in 2017 (41.35002N, 17.19997E,

Fig. 2) is located at 580 m water depth on the northern slope of the northern branch of the Bari Canyon (Channel B in Trincardi et al. 2007a; Angeletti et al. 2014), close to a steep, almost vertical wall of the canyon channel. The seafloor at the site is bathed by high-density currents of the North Adriatic Dense Water and is topographically complex with many small depressions. At 580 m, the salinity is equal to 38.8 psu and bottom-water temperature to 13.6-13.8°C (Trincardi et al. 2007b). Three Van Veen grabs (numbered as 5, 6, and 7) were wet sieved with 1 mm mesh. They all contained dead specimens of G. vitreus (Fig. 4). No individuals of G. vitreus were collected alive. All Gryphus specimens larger than 20 mm (n = 59) were immediately picked from grab 6 (labelled as POS514-14-6) during the wet-sieving at the ship, and additional 33 specimens larger than 3 mm were exhaustively picked from the 1/8 of the grab to increase the size coverage of the G. vitreus distribution of postmortem ages.

In total, 92 specimens (31 articulated shells, 39 ventral valves, and 22 dorsal valves) were dated with ¹⁴C, using the direct carbonate accelerator mass spectrometry (Bush *et al.* 2013). This method is based on powdered carbonate targets and produces



Fig. 4. Skeletal-rich death assemblage residues sieved at 1 mm with tubes of *Ditrupa* and valves of *Gryphus vitreus* collected in three grabs 5, 6 and 7 at 580 m at the station POS514-14 in the Bari Canyon. (a) Grab 5; (b) grab 7; (c) grab 6. The residue of grab 6 lacks articulated shells (>20 mm) that were removed during initial sample sieving and processing at the ship (the photograph captures approx.1/8 of the grab, and thus 4–5 shells of *Gryphus* are missing from it).

postmortem ages that are consistent with the standard graphite ¹⁴C method (Bright et al. 2021). ¹⁴C ages of four specimens range from 7350 to 21 790 ¹⁴C years. All the other specimens are younger than 3000 ¹⁴C years (with two postbomb ages). Radiocarbon ages were calibrated to calendar ages with the R package rearbon (Bevan et al. 2017). We used the Marine13 database (Reimer et al. 2013), with the corresponding ΔR (Reimer and Reimer 2001), and considered the estimate from the southern Adriatic Sea ($\Delta R = 121$ years based on Mimachlamys varia, Taviani and Correggiari, in Reimer and McCormac 2002). The calibration results are similar if $\Delta R = 61$ years based on *Glycy*meris shells collected in the central Adriatic Sea (Siani et al. 2000). The reservoir age in the Mediterranean Sea was higher by a factor of two during the last deglaciation (Siani et al. 2001), but as we focus on shells younger than 3000 years in the quantitative analyses of postmortem age distributions, we used the same ΔR in all calibrations. The calibrated ages are reported in years before present (BP, i.e. before the year 1950, except in analyses of disintegration rates where postmortem ages are rescaled to the time of sampling in AD 2017). Age errors (represented by one standard deviation) of probability distributions of calibrated ages are c. 50 years for specimens <1000 years BP, c. 80-90 years for specimens that are c. 1000-3000 years old, and exceed 100 years for the oldest shells. To assess the effect of these errors on the shape of the distribution, on the location of modes, and on time averaging (measured with an inter-quartile age range, IQR), we estimated postmortem age of each specimen by sampling its age from the calibrated probability age distribution, computed the corresponding frequency of specimens in 50-year cohorts and repeated this sampling 10 000 times. We then estimated the mean frequency (with 95% confidence intervals) of specimens in 50-year cohorts. All radiocarbon and calibrated ages (Supplement A) and the script in R language (R Core Team 2021, Supplement B) are available as supplementary materials.

As postmortem age-frequency distributions of the two subsets of dated shells (59 larger specimens and 33 additional specimens) are similar (bimodal), we pool them in all analyses. In total, 285 specimens of *G. vitreus* (33 shells, 92 dorsal valves, and 160 ventral valves) were exhaustively picked from the grab 6 and 92 of them were dated with ¹⁴C. Boullier *et al.* (1986) documented that a shift in the slope of the relationship between shell length and loop length occurs at *c.* 20 mm shell length in *G. vitreus*, and we use this size threshold as an indication that all individuals >20 mm are adults, even when maturity can be achieved at smaller size in similarly-sized brachiopods (Thayer 1977). The resulting size distribution of all shells suggests that

most adult individuals >20 mm collected in grab 6 were dated. Therefore, the number of radiocarbon dated individuals exceeding 20 mm effectively corresponds to the total abundance of dead adult specimens in 0.1 m² sampled by one Van Veen grab. The preservation of all specimens (disarticulation and completeness of valves) and the presence of encrusters, borers and mineral precipitates on external and internal surfaces were scored under a light microscope at 10-20× magnification. Postmortem age distributions show number of specimens binned to 50-year, and 100-year cohorts (Fig. 5). The relative frequency of articulated shells per cohort (white bars in Fig. 5) is estimated relative to the total number of shells summed with the number of ventral valves in that cohort. The estimation of the location of modes and testing for the bimodality of postmortem age data (with the null hypothesis represented by one mode) follows Ameijeiras-Alonso et al. (2019), implemented in Ameijeiras-Alonso et al. (2021).

Dead specimens can be lost from the surface sediments of the mixed layer either by disintegration or by burial. This total loss rate controls how the original chronological variability in abundance translates to the distributions of postmortem ages observed in the DA. The internal modes in postmortem agefrequency distribution and the lack or rarity of recently-dead cohorts complicates the estimation of loss rates with taphonomic models that assume temporally-constant production of shells. Here, we account for the decline in abundance by truncating the distribution at the location of the first major pulse at 450 years BP (i.e. prior to the first main peak in the frequency of specimens at c. 500 years BP) and at 5000 years BP (i.e. benthic environments were later exposed to the onset and offset of the sapropel phase). We then approximate the loss rate by fitting the postmortem age-frequency distribution (setting postmortem ages relative to the time of sampling, i.e. AD 2017) to a model with a single, temporally-constant rate of loss from the TAZ (i.e. loss rate in the mixed layer does not change with shell postmortem age) and to a model with a temporally-declining rate of loss from the TAZ (i.e. loss rate declines as a dead shells age in the mixed layer), both accounting for abrupt onset and offset in shell production (abruptly-truncated models, Tomašových et al. 2016). The first model assumes that disintegration rate in the TAZ and burial rate below the TAZ remain constant in time. The second model postulates that loss rate λ_1 declines to λ_2 at sequestration rate τ . The burial rate of skeletal remains below the surface increment sampled by the Van Veen grab is probably limited at the sampling station because abundant iron-stained skeletal remains and abraded and encrusted intraclasts indicate condensation and very slow net sedimentation rate. Therefore, we neglect the contribution of burial



Millennial-scale changes in brachiopod abundance

Fig. 5. (a) and (b) The complete frequency distribution of postmortem ages of *Gryphus vitreus* at POS514-14-6 in 200-year cohorts (a) and the subset (b) showing specimens younger than 3000 years BP in 50-year cohorts, with internal modes at *c*. 500 years and 1750 years BP. Each specimen age corresponds to the median of the probability distribution of calibrated ages. Shading highlights the contribution of articulated shells (white) and disarticulated ventral (black) and dorsal valves (grey). The distributions are based on all dated specimens. The two modes are captured by both articulated and disarticulated specimens, although the contribution of articulated shells diminishes in older cohorts. Both small and large specimens contribute to both modes. The inset in (a) shows the fit of the two taphonomic models to age data that were used to approximate loss rates of specimens from the taphonomic active zone (disintegration rates), truncating the distributions at 450 and 5000 years. (c) Size distribution of all *G. vitreus* specimens (n = 285) collected in grab 6, with age-dated specimens of *G. vitreus* in grey. The age-dated subset closely captures the mode formed by adult specimens exceeding 20 mm in length.

to loss rate and refer to this loss-rate estimate as disintegration rate (λ) below.

We unmix time-averaged abundance of *G. vitreus* to abundance of specimens produced during 100 years by dividing the counts in 100-year age cohorts by the survival function of the model with temporally-constant rate (the fit of the model with temporally-declining rate to postmortem ages is poor), using the estimate of disintegration half-life derived from the model. The survival function corresponds to $e^{-\lambda t}$ (λ refers to disintegration rate and *t* refers to time). We scale then these counts to yearly population density within each of the 100-year time bins. The lifespan of *G. vitreus* is unknown, but as the valves at the sampling station possess about ten or more regular growth increments, we assume that the lifespan of adult individuals is *c*. 10 years (similar in magnitude when compared to lifespan of other temperate brachiopods of similar size, Doherty 1979; Baird *et al.* 2013; Baumgarten *et al.* 2014) and discuss the sensitivity of the estimated density if the lifespan would be just 5 years. We use both disintegration models (with temporally-constant and temporally-declining disintegration rate) to (1) visualize the transition between the original time series in abundance and the postmortem age-frequency distribution observed in the surface DAs and to (2) illustrate the procedure of unmixing and scaling of

time-averaged abundance observed in a DA to yearly standing population density expected in a living community.

Models that assume that the original increase and decrease in abundance can be approximated with a symmetric normal-shaped trajectory can be used to evaluate the time-scale of the decline in abundance of Gryphus, by estimating the standard deviation of the postmortem age-frequency distribution with the so-called truncated-normal models (Tomašových et al. 2016). These models estimate the mean and the standard deviation of the age distribution sourced by a population that exhibited a temporally symmetric trajectory in abundance and was subjected to disintegration in the mixed layer. The mean is a function of (i) the original timing of maximum abundance but also of (ii) disintegration rate and (iii) the rate of change in abundance. However, the standard deviation is just a function of rate of change and is independent of disintegration rate. We fit the so-called truncated-normal model to postmortem ages, using both models of disintegration. We limit this estimation to the last pulse as detected in the death assemblage (i.e. at 500 years BP), and thus exclude specimens older than 1500 years.

Results

DAs in all three grabs are rich in skeletal and intraclastic sand and gravel (Fig. 4). They consist of abundant polychaetes (Ditrupa), benthic molluscs (e.g. Pseudamussium clavatum, Abra longicallus, Kelliella miliaris, Bathyarca sp., Asperarca sp., anomiid bivalves), pteropods, bryozoans, decapod fragments (Ebalia nux), cidaroid spines, benthic foraminifers (e.g. Ammolagena clavata), otoliths, and G. vitreus. In addition to Gryphus, the assemblage also contains disarticulated valves of the micromorphic brachiopod Platidia anomioides. Only dead individuals of G. vitreus were collected. The time averaging (IQR) of G. vitreus specimens is 1250 years, the 90th percentile range is 2170 years (Fig. 5a, b). The median postmortem age is 640 years BP and the mean postmortem age is 1750 years BP. The two youngest specimens represented by articulated shells show postbomb ages. The calibrated ages of the three oldest specimens fall between 18 210-25 620 years BP, and the age of one specimen is 7360 years BP. All other 88 specimens are younger than 3000 years BP (Fig. 5a). The fitting of the model abruptly truncated at 450 and 5000 years BP to postmortem age data (rescaled to AD 2017 so that residence time reflects time since death of individual brachiopods) generates the estimate of disintegration rate λ equal to 0.00055 (i.e. half-life is c. 1250 years, Fig. 5a). The fitting of the abruptly-truncated model with temporally-declining

disintegration to postmortem age data significantly underestimates the older pulse at 1750 years, and we thus use the model with temporally-constant disintegration in converting time-averaged abundance to annual population density.

For specimens younger than 3000 years BP, the distribution is bimodal ($p [H_0 = \text{one mode only}]$ <0.001; p [H₀ = 2 modes] = 0.44), with the location of the two modes estimated at 510 and 1710 years BP (Fig. 5b). Accounting for the calibration errors, the frequency distribution of specimen postmortem ages binned to 50-year cohorts is also bimodal (Fig. 6), with the mean age of the first mode at 500 years BP (95% c.i. = 460-530 years BP) and of the second mode at 1740 years BP (95% c.i. = 1680-1800 years BP). The 95% confidence intervals on time averaging (IQR) fall between 1190 and 1305 years (Fig. 6). The sizefrequency distribution is also bimodal, with abundant juvenile specimens smaller than 10 mm, and with a second mode formed by shells 25-30 mm in length (Fig. 5c).

The truncated-normal model with a temporallydeclining disintegration rate (grey lines) fitted to postmortem ages of specimens younger than 1500 years indicates that the standard deviation is 110 years (all specimens) or 120 years (specimens >20 mm). Rate of change in the increase (prior to the younger pulse at *c*. 500 years BP) and in the decline in abundance as fitted by this model thus indicates that most of the decline occurred at the scale of *c*. 100 years, and thus was not protracted over several centuries (grey lines in Fig. 7). This model does not overestimate the abundance of young cohorts, in contrast to the model with a temporally-constant disintegration (black lines in Fig. 7).

Preservation of G. vitreus exceeding 15-20 mm is variable, and includes complete shells with loops (brachidia) preserved (Fig. 8a-f), disarticulated but still complete valves with broken loops (Fig. 8g-i), and umbonal fragments that lack peripheral margins and are worn and discoloured, with few worn relicts from the Last Glacial Maximum (Fig. 8j, k). External surfaces of articulated shells are frequently encrusted by foraminifers, serpulids and bryozoans and bored on external surfaces with microendolithic organisms, clionid sponges (Entobia), whereas disarticulated valves are encrusted and bioeroded on both sides. A large proportion of old shells (c. 500–1800 years BP) is still articulated (white bars in Fig. 5a, b), contain complete loops (Fig. 8a-f) and is frequently internally coated by Mn and Fe oxides, especially along edges of the internal valve surfaces. The relative frequency (percentage) of articulated shells older than 500 years exceeds 50% in several 100-year cohorts (Fig. 5a, b). Small-sized specimens (<20 mm) are rarely articulated. Their preservation ranges from transparent or semi-transparent



Fig. 6. (a) Accounting for the uncertainty in age calibration shows the mean frequency of specimens, with specimen ages binned to 50-year cohorts, predicted by repeated (10 000) sampling of ages from the probability distributions of specimen calibrated ages, with 95% confidence intervals (dashed lines). (b)–(d) The mean age of the two modes (b–c) and the mean inter-quartile age range (d), with 95% confidence intervals.

complete valves to microbored, encrusted and discoloured valve fragments. The mean postmortem age of articulated shells is 650 years, the mean postmortem age of disarticulated valves with complete outlines is 1236 years, and the mean postmortem age of fragments is 2950 years. The oldest articulated shell is 1771 years old (Fig. 8j, k). The four oldest specimens (>5000 years) are represented by strongly worn and discoloured relicts of the umbonal part of the shell.

Transition from chronological trajectory in abundance to age distribution

Assuming that burial of shells below the TAZ and out-of-habitat transport is negligible, we can model the transition from the original, biphasic chronological series in species abundance to the abundance of postmortem-age cohorts observed in the surface sediment. We specify that two pulses occur at 500 and 1750 years BP, and standard deviation of rate of



Fig. 7. The truncated-normal model with a temporally-declining disintegration rate (grey lines) fitted to postmortem ages of specimens younger than 1500 years estimates the standard deviation at 110 years (\mathbf{a} , all specimens) or 120 years (\mathbf{b} , specimens >20 mm), indicating that the decline in abundance of *Gryphus* occurred at the scale of one century and was thus not protracted over several centuries. (\mathbf{a}) The truncated-normal models with a temporally-constant disintegration (black lines) overestimate the abundance of the youngest cohort (standard deviation is years on the basis of all specimens and years on the basis of specimens >20 mm).



Fig. 8. Preservation states of *Gryphus vitreus*, with (**a**–**h**) articulated shells with (d–f) complete and (g and h) incomplete loops, (**i** and **j**) disarticulated valves and (**k**) worn umbonal relicts. Specimen numbers and their postmortem ages (BP) follow the letters. (a) 39 (742 years), (b) 4 (406 years), (c) 2 (503 years), (d) 18 (1157 years), (e) 4 (406 years), (f) 2 (503 years), (g) 8 (399 years), (h) 49 (514 years), (i) 49 (514 years), (j) 43-4 (1931 years), (k) 54 (2113 years). Scale bars: (a)–(c), (i), (j) 10 mm; (d)–(h), (k) 5 mm.

change in abundance is equal to 120 (as estimated from empirical data, Fig. 9a, b) and 300 years (to approximate a slower rate of change taking place over several centuries, Fig. 9c, d). These two scenarios are subjected to two preservation regimes (1) with a temporally-constant disintegration rate (Fig. 9a, c) and (2) with a temporal decline in disintegration rate (Fig. 9b, d). As shells located in the TAZ are exposed to disintegration, the original pattern of change in abundance through time is transformed into the DA postmortem age-frequency distribution, in which the number of individuals naturally declines in each age cohort owing to disintegration. The decline is not equal across the cohorts when burial rate is neglected or is smaller than disintegration rate. The abundance of older cohorts is thus penalized by disintegration rate more strongly relative to the abundance of younger cohorts, leading to the smaller abundance of older cohorts in the DA (Fig. 9a-d).

When increasing disintegration half-lives (i.e. the natural logarithm of 2 divided by disintegration rate) from 12 (very fast disintegration) to 1200 years (i.e. as approximated from empirical postmortem age data of all specimens) in the model with temporally-constant disintegration in the mixed layer, an original time series with two abundance pulses occurring at c. 500 and 1750 years BP (dashed lines in Fig. 9) is

increasingly closer in shape to the distribution expected to be observed in a surface DA (black lines in Fig. 9). The timing of maximum abundance is taphonomically pulled towards the younger cohorts with decreasing disintegration half-life (black arrows on Fig. 9a, c), especially when the temporal decline in abundance is not abrupt (Fig. 9a) but gradual (Fig. 9c). However, when disintegration halflife of shells increases as they reside in the TAZ, the mode shifts towards the younger cohorts only slightly when disintegration half-life exceeds median time to sequestration (Fig. 9b, d, half-time to sequestration is assumed to be 200 years). Only when disintegration half-life is very short (12 years), the resulting distribution is right-skewed, dominated by very young shells, and the original modes disappear (Fig. 9a, c). Long disintegration half-lives in the TAZ thus minimize the bias in the shape of surface postmortem age-frequency distributions relative to the original chronological changes in abundance.

Effects of disintegration rate on temporal variability in abundance of Gryphus

Dividing the counts of shells binned to 100-year cohorts in the observed *Gryphus* distribution of



Fig. 9. The effect of shell disintegration rate and rate of change in abundance of living individuals on the shape of age-frequency distributions in the surface death assemblages in two preservation models (a model with the temporally-constant disintegration rate in (a) and (c) and a model with the declining disintegration rate in (b) and (d)). The original chronological trajectory in abundance is defined by two pulses at *c*. 500 years and 1750 years ago (dashed lines), and by a fast (a and b) or slow increase and decline in abundance (c and d). (a) and (b) Scenario with fast increase and decline in abundance (standard deviation = 120 years, as approximated from empirical age distribution of *Gryphus*). (c) and (d) Scenario with slower increase and decline in abundance (standard deviation = 200 years). Disintegration half-lives range in four scenarios from 12 to 1200 years in the model with temporally-constant disintegration (a, c). The disintegration half-lives of 12 and 120 years increase to a half-life of 1200 years in all scenarios in the model with age-declining disintegration (b, d). In (a) and especially in (c) where the rate of decline in abundance is slow, the mode of the distribution is visibly pulled towards the youngest cohorts with increasing disintegration rate, i.e. with shorter disintegration half-life (black arrows). In (b) and (d), the mode shifts only slightly unless disintegration half-life = 12 years).

postmortem ages (dashed lines in Fig. 10a, b, based on all dated specimens) by survival function of the two taphonomic models, the reverse procedure occurs, and abundance of individuals in cohorts increases towards the original chronological pattern in abundance. If shells degrade at a temporallyconstant rate, with long half-life equal to 1200 years (black lines), the two pulses at 500 and 1750 years BP have similar magnitudes. If half-life declines to 700 years in the model with constant disintegration half-life (grey lines in Fig. 10a), the older pulse is more pronounced than the younger pulse (Fig. 10a). If disintegration half-life increases with increasing postmortem age of shells in the TAZ (Fig. 10b), with the initial decadal-scale half-life of 90 years increasing to 1200 years in all scenarios,

the slow sequestration in the model with temporallydeclining disintegration rate is equivalent to longer duration of the taphonomic regime with fast disintegration in the model with temporally-constant disintegration half-life. The scenario with slow sequestration thus increases the magnitude of the younger mode relative to that of the older mode, and the original trajectory in abundance thus can be characterized by differences in the magnitude of abundance pulses (black lines, Fig. 10b). When median time to sequestration is similar to disintegration halflife, the difference between the initial trajectory in abundance and the final postmortem age-frequency distribution is smaller (grey lines, Fig. 10b).

In contrast to the model with temporally-constant disintegration rate, very old shells disintegrate at



Fig. 10. (a) and (b) The empirical distribution of *G. vitreus* postmortem ages binned to 100-year cohort (observed in a surface death assemblage) is shown in dashed lines. The modelled distributions (prior to disintegration) in solid and grey lines represent the original trajectories in abundance (computed per 100 years at the scale of grabs, i.e. 0.1 m^2) that generated, under a given combination of disintegration and sequestration rates, the empirical age distributions. In (a) (shells degrade at temporally-constant rate), the pulses at 500 and 1750 years have similar magnitudes in the reconstructed trajectory of abundances when the half-life is 1200 years (black lines). If half-life rate declines to 700 years (grey lines), the older pulse is more pronounced than the younger pulse. In (b) (disintegration declines with shell age, the initial half-life of 90 years increases to 1200 years in all scenarios), the slow sequestration is equivalent to longer duration of the initial phase of disintegration. When time to sequestration is short, the difference between the age distribution and the chronological trajectory in abundance is small. (c) and (d) Reconstruction of annual population density (per 1 m²) of *G. vitreus* individuals larger than 20 mm (lifespan = 10 years), in models with temporally-constant disintegration (c) and temporally-declining disintegration (half-life declines from 90 years to 1200 years in all scenarios, (d)). The empirical distribution of postmortem ages is based on the unique number of individuals (shells and ventral valves), with the ventral valve exceeding 20 mm (dashed lines in (c) and (d)).

negligible rate in models with temporally-declining disintegration rate, and the magnitude of older modes (i.e. abundance maxima) thus does not increase in the reconstructions of the original chronological trajectory in abundance of Gryphus. These scenarios show that, although the magnitude of modes can shift, depending on the model and the magnitude of disintegration rate, the disintegration alone or its temporal decline does not generate internal modes. Notably, the disintegration rate does not shift the timing of modes in abundance of cohorts markedly (Fig. 10a, b) because the modes of the empirical postmortem age-frequency distribution are relatively sharply separated from the lows. The decline in abundance of *Gryphus* had to take place at the scale of a century (standard deviation of the

rate of change in abundance as estimated for the younger pulse is 110–120 years) rather than over several centuries (as in Fig. 9c, d, where abundance was modelled with the standard deviation of 300 years).

Scaling time-averaged abundance to population density

Using the minimum number of individuals (the sum of shells and ventral valves), assuming zero net sedimentation rate, lifespan not shorter than 10 years, and the range of half-lives centred around 1200 years, reconstructing standing densities indicates that populations of adults of *G. vitreus* (exceeding 20 mm) achieved at least *c.* 20 individuals/m² at

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500 years BP in the Bari Canyon (Fig. 10c). A yearly density of 40–60 individuals/m² is predicted when the disintegration half-life of 90 years increases to 1200 years in the course of 100 years (light grey line in Fig. 10d) or 200 years (dark grey line in Fig. 10d). If the half-life of 1200 years also subsumes the loss of specimens below the TAZ by burial, the yearly density is also expected to be higher relative to the initial estimate of 20 individuals/m².

Discussion

The effect of disintegration rates on the detection of past abundance fluctuations

The highest abundance of the youngest cohorts and the declining abundance or the lack of older cohorts is a null expectation under steady-state production and any non-negative disintegration or burial rates: it is expected to produce right-skewed, exponential or heavy-tailed frequency distributions of postmortem ages (Olszewski 1999, 2004; Carroll et al. 2003). The modelling scenarios in Figures 9 and 10 visualize that high disintegration rates in the TAZ can maximize the bias in the shape of surface postmortem age-frequency distributions relative to the original chronological changes in abundance (with fast sequestration minimizing the bias). The multi-centennial estimate of disintegration half-life generated by fitting the abruptly-truncated taphonomic model to age data (half-life = 1200 years) can be biased upwards because young cohorts are rare, and the initial phase of their residence time in the TAZ potentially characterized by high disintegration cannot be observed. However, as the proportion of articulated shells is very high even in very old cohorts and exceeds 50% in younger age cohorts, we assume that the disintegration half-life of several centuries is not markedly biased up by the rarity of recent Gryphus individuals, and that brachiopod disintegration rates at this sampling location are probably very low. All modelling scenarios show that internal modes in abundance of cohorts within surface postmortem age frequency distributions indicate past maxima in abundance of G. vitreus.

Subsurface assemblages can be also characterized by internal modes that reflect their time to burial into subsurface layers even when temporal production of shells at the sediment–water interface is constant in time. The exhumation of such subsurface assemblage back to the TAZ (with the original surface sedimentary particles that were overlying the DA with abundant *Gryphus* being fully removed by transport elsewhere, thereby exhuming the *Gryphus* assemblage to the seabed surface) can thus generate a surface assemblage with an internal mode in cohort abundance. However, this scenario with subsurface

exhumation of formerly-buried assemblage predicts the lack or rarity of young cohorts of any species. Abundant, well-preserved and delicate bivalve shells of Kelliella miliaris and Abra longicallus do not show any signs of coatings formed by Fe or Mn oxides in the death assemblage collected at the station POS514-14 (in contrast to well-preserved Gryphus shells). Their high abundance in present-day living assemblages in the southern Adriatic Sea (Nasto et al. 2018) and their low alteration indicate that these shells are young, and the scenario with full removal of the surface assemblage and the exhumation of older shells from subsurface zones is unlikely. Therefore, we suggest that temporal variability in abundance of G. vitreus, driven by changes in the population dynamics is responsible for the presence of two distinct pulses at 500 and 1750 years BP in postmortem age-frequency distributions of G. vitreus.

Reconstructing population density

Gryphus presently has not been observed to form any densely-populated beds, patches or aggregations on the seafloor in the Southern Adriatic Sea, in contrast to occurrences in the Western Mediterranean Sea where *Gryphus* forms shell-rich aggregations, with densities between tens and hundreds of living individuals/m² (Emig 1989a; Grinyó et al. 2018). The population density of Gryphus in the Mediterranean Sea falls into three categories: (1) 1–5 individuals/ m² in sparsely-populated habitats (at 110–330 m in the Menorca Channel, Grinyó et al. 2018), (2) several tens of individuals/ m^2 (in the upper bathyal at 180– 250 m off Corsica and Provence), or (3) dense populations exceeding 100s of individuals (at 130-180 m off Corsica and Provence, and at Banc de Magaud near the Hyères Islands, Emig 1987). Multiple camera surveys performed in the Bari Canyon in the twenty-first century did not detect any sites with dense populations of Gryphus (D'Onghia et al. 2015; Angeletti et al. 2019). However, although distributions of postmortem ages indicate that the present-day abundance is smaller than in the past, it is unclear whether the internal modes observed in the distribution of postmortem ages correspond to ecologically-significant population densities equivalent to those observed in the Western Mediterranean Sea today. When time averaging is extensive and disintegration slow, even a species with low standing abundance can become abundant in DAs.

If disintegration were to be neglected, when computing the standing (yearly) density per 1 m² on the basis of 100-year cohorts, the division of abundance in each cohort by ten generations (assuming lifespan of 10 years) and the multiplication of the same count by ten (from the scale of Van Veen grab capturing 0.1 m^2 to 1 m^2) would cancel each other out. A

shorter lifespan of 5 years would divide the estimate by half. The resulting estimates of standing densities that do account for disintegration do not markedly differ from those predicted from the raw distribution because the inferred disintegration rates of Gryphus are very low (Fig. 10c, d). These estimates of population density are minimum estimates and higher loss by disintegration or other sources of loss by burial or transport to adjacent habitats would imply even higher standing density. To conclude, the estimates of several tens of individuals of Gryphus per 1 m² indicate that this species was truly abundant at times corresponding to the two modes in the postmortem age-frequency distribution. These estimates are comparable to intermediate levels of its densities presently observed in the Western Mediterranean Sea.

Ecological causes of volatility

Bathyal benthic communities with epifaunal suspension-feeders in the Southern Adriatic Pit were affected by major multi-millennial variability in deep-water circulation during the Holocene. The bottom currents decreased in velocity at about c. 7-10 kyr BP, leading to the deposition of sapropel in the Southern Adriatic Pit (Narciso et al. 2012; Tesi et al. 2017). Corals re-colonized the Bari Canyon after the sapropel phase (Taviani et al. 2019). The distribution of living cold-water corals is asymmetric in the Southern Adriatic Pit during this postsapropel phase, with higher abundance on the western side (Taviani et al. 2019). In the eastern side of this basin, the growth of cold-water corals is sporadic and localized, and most records are based on surface DAs (Angeletti et al. 2014, 2015; Nasto et al. 2018). However, this turnover related to the onset and offset of the sapropel phase occurred prior to the volatility detected in abundance of Gryphus as observed here.

The bathyal communities with G. vitreus belong to the EUNIS category (European Nature Information System) 'Communities of bathyal detritic sands with Gryphus vitreus' and the LPHME category (Lista Patrón de los Hábitats Marinos Presentes en España) 'Bathyal grounds in the continental shelf edge with Gryphus vitreus'. They are considered as vulnerable community types in the Mediterranean Sea negatively affected by deep-water trawl fishing (d'Onghia et al. 2003; Maynou and Cartes 2012; Aguilar et al. 2017; Carpentieri et al. 2021). The Bari Canyon, however, is not subjected to deepwater trawling operations owing to its steep topography (Sion et al. 2019), and trawlers primarily operate on the shelf up to 200 m, only occasionally affecting the borders of the canyon (Angeletti et al. 2021). The re-suspension of fine-grained sediments produced by trawling on the shelf may trigger increasing siltation of the canyon ecosystems (e.g. Martín et al. 2008).

However, the bottom trawling activities became industrialized during the second half of the twentieth century in the Adriatic Sea (Fortibuoni *et al.* 2010, 2017; Carpi *et al.* 2017) whereas the decline in abundance of *Gryphus* occurred earlier, as suggested by a relatively fast, centennial-scale decline in abundance after the maximum abundance was achieved at *c.* 500 years BP (Fig. 7).

Siltation and smothering have significant effects on the viability of epifaunal, immobile, suspensionfeeding brachiopod populations (Simões et al. 2009; Tomašových and Kidwell 2017), even when long-term sedimentation rates are negligible and smothering of living individuals is transient. The Bari Canyon represents a conduit through which suspended fine sediment generated by the Po and other rivers is redistributed southwards along the Italian coast by the predominant count-clockwise circulation pattern (Cattaneo et al. 2003), being conveyed into the deeper parts of the Southern Adriatic Pit. The river discharge and sediment load at the prograding deltas sourced by the multitude of rivers that drain the Apennines, and enter into the central Adriatic Sea, increased over the past centuries driven by climatic (with the Last Ice Age phase leading to higher runoff) and anthropogenic changes of the watershed (deforestation and levee construction, Cattaneo et al. 2003, 2004, 2007; Correggiari et al. 2005a, b; Stefani and Vincenzi 2005; Mercuri et al. 2019). Cattaneo et al. (2003, 2007) documented an overall increase in net sedimentation rate over the past 500 years near the Garganto promontory as recorded in sediments of the Garganto subaqueous delta. We hypothesize that these changes led to higher extent of siltation that negatively affected suspension-feeding brachiopods in the Bari Canyon. This hypothesis is supported by the orientation of downward-facing growth forms of corals in the Bari Canyon system that indicates that they avoid the smothering effect (Freiwald et al. 2009). Furthermore, sponges in the Bari Canyon are affected by siltation as well (Bo et al. 2012), indicating that the currents are not strong enough to remove sediment and to buffer the deleterious effects of siltation. Pedunculate brachiopods such as Gryphus with limited repositioning abilities and smaller body size are probably more vulnerable to siltation than epibenthic corals or sponges (Emig 1989b).

Changes in deep-water circulation in the Southern Adriatic Sea over the past millennia (Piva *et al.* 2008; Siani *et al.* 2013) may also have led to changes in sediment accumulation and contribute to the volatility of bathyal communities. Piva *et al.* (2008) documented millennial-scale turnovers in the deepwater circulation in the Southern Adriatic Sea owing to changes in the rate of formation and depth of flow of the North Adriatic Dense Water, in response to changes in precipitation intensity

and in the importance of river floods. The two reconstructed peaks in Gryphus abundance at c. 500 and 1750 years coincide with the former pulses of stronger contribution of the North Adriatic Dense Water to deep-water circulation (at 0.6-0.8 kyr BP and at 1.8-2 kyr BP, Siani et al. 2013). Although the dynamics of sediment dispersal and deposition in the Bari Canvon can be decoupled from sediment accumulation induced by river deltas, with sediment sourced from the outer shelf (Tesi et al. 2008), the marked expansion in sediment accumulation observed in the mudbelt can be expected to propagate, even with some delay, to bathyal environments. However, as shifts in deep-water circulation coincided with climatic and anthropogenic changes affecting the watersheds and the amount of riverborne sediment discharge along the western Adriatic coast, disentangling the effects of these potential drivers may be difficult. The estimates of postmortem age structure of Gryphus assemblages will need to be expanded to other locations and to broader geographical extents to uncover whether the ecological roots of pulses in its abundance in the Bari Canyon is local or regional in scale.

Postmortem age-frequency distributions as the source of information on baseline communities

The conditions with slow disintegration in the TAZ can lead to the formation of relatively complete and unbiased DAs. DAs can efficiently sample patchy or rare species and thus are used in systematic or monitoring surveys to capture long-term abundance structure of regional species pools and spatial distribution patterns (Nebelsick 1996; Bouchet et al. 2002; Warwick and Turk 2002; Zuschin and Oliver 2005; Smith 2008; Weber and Zuschin 2013; Powell et al. 2017; Casebolt and Kowalewski 2018; Bhattacherjee et al. 2021). For example, more than 70% of molluscan species were identified based only on dead shells in monitoring surveys off New Caledonia in habitats deeper than 100 m (Bouchet et al. 2009), and the identification of new molluscan species, especially in tropical environments, is frequently based on dead shells (Dijkstra and Maestrati 2010; Dijkstra and Janssen 2013).

The slow shell disintegration rates typical of many marine habitas, however, can represent a challenge for marine biologists and taxonomists that use dead shells as indicators of active inhabitants of the present-day seafloor. In such conditions well-preserved shells can be hundreds to thousands of years old, as articulated shells of *Gryphus* in the death assemblage observed here. Taphonomic clock can be poorly manifested in Holocene brachio-pod shells (Carroll *et al.* 2003) when relatively young dead shells are poorly-preserved as they are

permanently exposed in the TAZ, whereas some old dead shells can be well-preserved if they have been temporally sequestered below the TAZ and were exhumed to the surface mixed layer only recently. Old Gryphus shells tend to be extremely well-preserved and are still articulated, and their good preservation thus does not ensure that they died recently. On the one hand, estimation of postmortem ages performed in our study shows that wellpreserved articulated shells can be misidentified as members of a living local population or a regional species pool in monitoring and taxonomic surveys, thus misrepresenting the current distribution of a species or over-estimating local and regional abundance (Bouchet and Taviani 1992). On the other hand, when taking into account the long-term average of Holocene level-bottom benthic communities inhabiting the Bari Canyon, age dating identifies the temporary dominance of G. vitreus at this location during the Late Holocene. Therefore, although the apparently pristine preservation of millennia-old articulated shells of brachiopods can bias systematic or monitoring studies that describe species on the basis of dead shells and attempt to characterize the composition of communities at high temporal resolution, their postmortem age structure expands the utility of surface DAs in tracing the composition of pre-impact baseline communities and identifying their long-term volatility.

Conclusions

Highly time-averaged DAs represent a paradox for conservation palaeobiologists or historical ecologists: although they integrate the total community composition over increasingly longer time-scales, they lose their ability to track ecologically significant volatility. However, the postmortem age structure based on dating of multiple (tens to hundreds) individual shells can effectively detect historical changes over longer, millennial scales, without the need to collect sediment cores, and thus address both the mean composition and its volatility. The DA that consists of largely autochthonous shells uniquely records two pulses of very high abundance of the brachiopod G. vitreus in the Bari Canyon during the Late Holocene, in contrast to its present-day rarity. With rapid advances in radiocarbon dating, examination and age dating of dead shells from sample residues collected during surveys of living assemblages, should become a part of the toolkit used in biomonitoring and in identification of baseline communities.

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