Boron isotope systematics of cultured brachiopods: Response to acidification, vital effects and implications for palaeo-pH reconstruction

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Abstract

CO2-induced ocean acidification and associated decrease of seawater carbonate saturation state contributed to multiple environmental crises in Earth’s history, and currently poses a major threat for marine calcifying organisms. Owing to their high abundance and good preservation in the Phanerozoic geological record, brachiopods present an advantageous taxon for palaeo-proxy applications as well as studies on biological mechanism to cope with environmental change. To investigate the geochemical and physiological responses of brachiopods to prolonged low-pH conditions we cultured Magellania venosa, Terebratella dorsata and Pajaudina atlantica under controlled experimental settings over a period of more than two years. Our experiments demonstrate that brachiopods form their calcite shells under strong biological control, which enables them to survive and grow under low-pH conditions and even in seawater strongly undersaturated with respect to calcite (pH = 7.35, Ωcalc = 0.6). Using boron isotope (δ11B) systematics including MC-ICP-MS as well as SIMS analyses, validated against in vivo microelectrode measurements, we show that this resilience is achieved by strict regulation of the calcifying fluid pH between the epithelial mantle and the shell. We provide a culture-based δ11B–pH calibration, which as a result of the internal pH regulatory mechanisms deviates from the inorganic borate ion to pH relationship, but confirms a clear yet subtle pH dependency for brachiopods. At a micro-scale level, the incorporation of boron appears to be principally driven by a physiological gradient across the shell, where the δ11B values of the innermost calcite record the internal calcifying fluid pH while the composition of the outermost layers is also influenced by seawater pH. These findings are of consequence to studies on biomineralisation processes, physiological adaptations as well as past climate reconstructions.

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Keywords: Boron isotopic composition; Culturing experiment; Physiological response; Proxy calibration; Biomineralisation; Low-magnesium calcite; pH and pCO2 reconstruction

1. INTRODUCTION

Brachiopods have survived multiple Phanerozoic crises since their radiation during the early Cambrian and still

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today present widespread taxa (e.g. Carlson, 2016). They are abundant in the marine fossil record, and owing to their good preservation they have been increasingly used to assess past changes of ocean chemistry and climate state (e.g. Veizer et al., 1999; Farkas et al., 2007; Brand et al., 2012, Vollstaedt et al., 2014; Garbelli et al., 2017). Their shell composed of low-magnesium calcite makes them more resistant to post-depositional alteration of primary chemical signals and potentially an advantageous geochemical palaeo-archive (e.g. Brand and Veizer, 1980; 1981; Brand et al., 2003). Yet to date, brachiopod proxy relationships have been barely examined under controlled laboratory conditions, and in contrast to other marine calcifiers such as corals or foraminifera (e.g. Rollion-Bard et al., 2003; McCulloch et al., 2012a,b; Raddatz et al., 2014; Tanaka et al., 2015; Taubner et al., 2017; Jurikova et al., 2019) surprisingly little is known on the mechanisms that control the incorporation of various key elements and isotope systems into brachiopod calcite. Whereas the lack of culture-based studies poses an uncertainty for accurate interpretation of palaeo-records, it also limits our understanding of the biological processes that may have enabled brachiopods to endure strong environmental changes, which will be tested under rising anthropogenic pressures (such the present threat from CO2-induced ocean acidification).

The boron isotope ($\delta^{11}B$) composition of biogenic carbonates is considered to be one of the most reliable pH-proxies (e.g. Vengosh et al., 1991; Hemming and Hanson, 1992; Sanyal et al., 2000) as it reflects the pH of fluid from which it precipitated. Although this is primarily seawater, several organisms have been found to be able to modify internal chemistry at the site of calcification (Al-Horani et al., 2003; de Nooijer et al., 2009; Stumpp et al., 2012; Anagnostou et al., 2012; McCulloch et al., 2012a,b; Venn et al., 2013; Ramesh et al., 2017) as a result of active ion transport and precipitation under non-equilibrium conditions termed ‘vital effects’, complicating the interpretation and calibration of $\delta^{11}B$ data (Rollion-Bard and Erez, 2010). Calibrations based on $\delta^{11}B$ analyses of samples from known pH environments have enabled an evaluation of the extent of vital effects on $\delta^{11}B$. The latter has become relatively well established for foraminifera (e.g. Henehan et al., 2013; 2016), and to some degree for corals (e.g. Trotter et al., 2011), allowing new insights into perturbations of the marine carbon cycle (e.g. Gutjahr et al., 2017) and even species-specific pH-regulating capacities (e.g. McCulloch et al., 2012b; Ross et al., 2018). However, the use of these two groups is mainly limited to the Cenozoic; their insufficient preservation and availability limits their use for geological time scales predating the Cenozoic for which an alternative archive is needed. For this, brachiopods represent highly suitable candidates, provided that a preceding careful assessment of vital effects can be made. Moreover, detailed $\delta^{11}B$ examination may permit us to illuminate their pH-controlling mechanism, a key but not universally developed ability of calcifiers to withstand ocean acidification. So far, only few studies focused on the potential of brachiopods as $\delta^{11}B$–pH recorders (Lécuyer et al., 2002; Joachimski et al., 2005; Penman et al., 2013). Relationship between brachiopod $\delta^{11}B$ and ambient pH has been postulated by Lécuyer et al. (2002) and Penman et al. (2013) based on co-variations of $\delta^{11}B$ from natural samples and oceanic regimes with distinct seawater pH. These calibrations may, however, potentially involve large uncertainties as the role of a sole parameter on the calcite chemistry is likely to be obscured or hampered by additional unknowns such as environmental parameters, growth rates or internal vital effects.

In order to provide a systematic understanding of boron incorporation into brachiopod calcite, assess their potential as palaeo-archives, and decipher the role of vital effects, we investigated the $\delta^{11}B$ composition of three brachiopod species; Magellaniana venosa (Dixon, 1789), Pajaudina atlantica Logan, 1988 and Terebratella dorsata (Gmelin, 1791). Brachiopod specimens were cultured under controlled experimental settings comprising different pCO2 and thus pH treatments. Solution-based MC-ICP-MS (Multicollector-Inductively Coupled Plasma-Mass Spectrometer) analyses were used to establish a culture-based $\delta^{11}B$–pH calibration, discuss its implications for palaeo-pH and palaeo-CO2 reconstructions, and to approximate the extent of vital effects involved. Further, we carried out high spatial resolution SIMS (Secondary Ion Mass Spectrometry) measurements to study intrashell $\delta^{11}B$ variations and their link to biominalensation. Finally, we performed in vivo calcifying fluid pH measurements using microelectrodes, to our knowledge the first of its kind for brachiopods, to reconcile the interpretation of $\delta^{11}B$-related vital effects in the context of internal pH-regulating capacities, and to corroborate whether boric acid may in part be also incorporated into brachiopod calcite (Klochko et al., 2009; Noireaux et al., 2015; Movormatis et al., 2015; Balan et al., 2018).

2. MATERIALS AND METHODS

2.1. Brachiopod sites and sampling

We cultured three different brachiopod species over the period of approximately two years under controlled experimental settings in a climatically controlled laboratory of the KIMOCC – Kiel Marine Organism Culture Center at GEOMAR in Kiel, Germany. Specimens of the cold-water species M. venosa (Fig. 1a), the largest living brachiopod, were harvested from three sites in the Comau Fjord in Chilean Patagonia (Jetty 42°22′47″S, 72°24′56″W, Cross–Huinay 42°23′28″S, 72°27′27″W, and Lliguay 42°9′43″S, 72°35′55″W) at 21 m depth. We also found few T. dorsata (Fig. 1b), a commonly associated brachiopod species with M. venosa, and co-cultured them together with M. venosa. The Comau Fjord encompasses a 41 km long, 4.5 km wide and 487 m deep basin, and presents an oceanographically and ecologically well-studied area (Sievers and Silva, 2008a,b,c; Häussermann and Försterra, 2009; Pantoja et al., 2011). The fjord is characterized by strong vertical stratification, with fairly homogenous temperatures below the surface layer of 8–12 °C, salinity about ~30 (Baumgarten et al., 2013) and pH of 7.8–8.0 at 20 m depth (Janitzen et al., 2013). M. venosa is an abundant species in the fjord and all throughout Southern Patagonia, and is
found attached to hard substrates of the shallow subtidal zone to 2000 m depth (McCammon, 1973).

Specimens of the warm-water species *P. atlantica* (Fig. 1c) were collected attached to rocks (app. 60 to 80 individuals per rock) at 18 m depth from La Palma, Canary Islands (Punta Malpique 28°27′17″N, 17°50′40″W). *P. atlantica* inhabit hard substrates of caves and shaded overhangs where they are found cemented to the substrate in great numbers up to 5000 individuals m⁻² (Alvarez et al., 2005; Logan, 2004). While the physical conditions of the surface ocean are driven by seasonal variations, with a temperature range between 17–26 °C throughout the year (Giumerans and Canavate, 1994), the crevices colonised by *P. atlantica* likely see only narrow oscillations. In January 2016 we measured temperature of ~21 °C, salinity of ~36, and pH ~8.1.

### 2.2. Culturing and experimental setup

The environmental conditions in our experimental culturing tanks simulated the natural conditions (as detailed in the previous Section 2.1) for both cold- and warm-water species. Prior to the start of the experimental treatments, brachiopods transported from the field remained in an acclimatisation phase at control conditions for five weeks for *M. venosa* and *T. dorsata*. A biofilter, protein skimmer and UV sterilizer were used for CM filtering. All aquariums were equipped with calibrated automated sensors for continuous measurements of salinity (Conductivity Electrode), temperature (Temperature Sensor), and pH (pH-Electrode) connected to a GHL ProfiLux aquarium computer 3.1T (Kaiserslautern, Germany) for data storing, which were validated against manual measurements several times per week. An optical membrane-inlet CO₂ sensor (CONTROS HydroC® underwater CO₂ sensor; Kongsberg Maritime Contros GmbH, Kiel, Germany) calibrated for a pCO₂ range of 200–6000 μatm (with accuracy ~1% of reading; Fietzek et al., 2014) was used for continuous monitoring of the pCO₂ in the aquariums as well as the CO₂ in the room air.

Additionally, CM from all brachiopod-culturing aquariums, reservoir tanks, and deionised water source was routinely (roughly every month) sampled and analysed for major and trace element content, isotopic composition, total alkalinity (TA) and total dissolved inorganic carbon (DIC) content. Total alkalinity was determined immediately after water sampling via open-cell titration of 0.5 ml samples with 0.01 M HCl in a titration vessel after Pavlova et al. (2008) using a Metrohm 876 Dosimat plus (Ω Metrohm, Florida, USA). During titration, the vessel
was continuously purged with N₂ to strip CO₂ released by acid addition. All TA measurements were calibrated with IAPSO seawater standard and had an accuracy of 5%. For DIC measurements, 1 ml of filtered water sample (28 mm diameter, SFCA membrane, 0.2 μm mesh size) was transferred into a CO₂-free 10 ml glass vial and sealed. Subsequently, 100 μl of 6 M HCl was added. After minimum of 6 h of equilibration time 100 μl of gas headspace was measured using a gas chromatograph (Shimadzu) equipped with a methane detector. DIC measurements were calibrated with bicarbonate solutions of concentrations ranging from 0 to 25 mM and an IAPSO seawater standard, and had an accuracy of 5%. The CM samples for elemental and isotopic analyses were collected in 4 ml vials using syringe filters (28 mm diameter, SFCA membrane, 0.2 μm mesh size), acidified (40 μl concentrated HNO₃ for the total volume of 4 ml), and stored in a refrigerator room at 4°C in a laminar flow hood. Targeted growth increments (approximately 250–500 μm) of each brachiopod species were achieved by continuous bubbling of CO₂-enriched air through an air stone, which we found to be more effective than via a skimmer. We started culturing enriched air through an air stone, which we found to be more effective than via a skimmer. We started culturing:

Table 1

|---------|-------------|---------|-------------|-------------|----|-------------|-------------|--------|
| CO₂ phase 1 (P1): August 2016–April 2017
| A1 | Control pH₂ | P. atlantica | 36 22 | 600 | 8.20 | 3.1 | 2.6 | 9.9 |
| A3 | pCO₂ pH₁ | P. atlantica | 36 22 | 2000 | 7.70 | 3.3 | 2.7 | 4.6 |
| B1 | pCO₂ pH₁ | P. atlantica | 36 22 | 1500 | 7.80 | 3.3 | 3.5 | 4.8 |
| C2 | Control pH₂ | M. venosa | 30 10 | 600 | 8.00 | 2.3 | 2.0 | 2.0 |
| C3 | pCO₂ pH₁ | M. venosa | 30 10 | 2000 | 7.60 | 2.8 | 2.7 | 1.1 |

CO₂ phase 2 (P2): April 2017–July 2017

| A1 | Control pH₂ | P. atlantica | 36 22 | 600 | 8.25 | 3.8 | 2.6 | 10.4 |
| A3 | pCO₂ pH₁ | P. atlantica | 36 22 | 2000 | 7.70 | 3.3 | 2.8 | 3.9 |
| B1 | pCO₂ pH₂ | P. atlantica | 36 22 | 4000 | 7.35 | 3.5 | 3.1 | 2.0 |
| C2 | Control pH₂ | M. venosa | 30 10 | 600 | 8.15 | 2.9 | 2.9 | 3.5 |
| C3 | pCO₂ pH₂ | M. venosa | 30 10 | 4000 | 7.35 | 3.1 | 3.5 | 0.6 |

Brachiopod shells were gently rinsed with ultrapure water (Milli-Q), and dried over few days on a hotplate at 40°C in a laminar flow hood. Targeted growth increments precipitated under specific experimental conditions (Fig. 1) were sub-sampled for mini-bulk analyses under binoculars using a precision drill (Proxxon) with dental tips (~100 μm diameter). Approximately 1–3 mg of CaCO₃ powder was collected in 1.5 ml centrifuge vials, integrating a shell growth interval of about one to two months. Powders were rinsed with Milli-Q followed by an oxidative cleaning procedure (following Barker et al., 2003) during which organic matter was oxidised with 1% hydrogen peroxide buffered with 0.1 M ammonium hydroxide at 80°C (approximately 250–500 μl of oxidative mixture was used per sample). Cleaned samples were subsequently transferred into 1.5 ml centrifuge Teflon vials and treated with 0.5 mM HNO₃ (typically 250 μl per sample) for 30 seconds to remove any adsorbed contaminants (following Foster, 2008). Finally, we dissolved the samples in 0.5 M HNO₃ with the aid of ultrasonication (depending on sample size, approximately 100–200 μl of 0.5 M HNO₃ was required to achieve complete dissolution). An aliquot of 10% of each dissolved sample was analysed for major and trace elements (Li, Na, Mg, Ca, Sr, and Ba; for the present study only B and Ca are relevant, to fully explore all major and trace element data is beyond the scope of this manuscript and will be presented in a separate publication) using a Quadrupole ICP-MS (Agilent 7500x) at GEOMAR, Kiel. The long-term external reproducibility RSD (relative standard deviation,
2σ) was better than 3% for B/Ca assessed by repeated measurements of carbonate reference materials ICp-1, ICT-1 and our in-house brachiopod standard MVS-1. Culture medium samples were measured for major and trace elements (Li, B, Na, Mg, K, Ca, Sr, and Ba) with an ICP-OES (Inductively Coupled Plasma-Optical Emission Spectrometry) The long-term external reproducibility RSD (relative standard deviation, 2σ) was better than 2% for B/Ca determined by repeated measurements of IAPSO standard seawater.

Prior to boron isotope analysis, boron is separated from the sample’s carbonate matrix on micro-columns (25 μl volume) using boron specific anionic exchange resin Amberlite IRA 743 crushed and sieved to 63–120 μm (Kiss, 1988; Lemarchand et al., 2002a). Samples were buffered to a pH of 5 using 2 M sodium acetate–0.5 M acetic acid buffer (typically twice the amount of buffer than the dissolution of 0.5 M HNO3 is required) and carefully loaded on the chromatographic columns. Boron retention on the resin is pH dependent (see Lemarchand et al., 2002a); thus repeated rinses with Milli-Q water were necessary to elute the carbonate matrix and the buffer, while purified boron was collected with 720 μl of 0.5 M HNO3. All columns were individually calibrated with yields verified by a highly resolved elution scheme (consisting of minimum 30 steps), yielding >98% boron recovery. To ensure that no significant amount of the sample remained on the column, boron content of each elution tail was determined representing <1% of the total boron in the sample. Likewise, a total procedural blank (TPB) for each column batch was determined; all TPBs were below <100 pg (<0.1% of the sample size, n = 12). Purified boron samples were stored in Teflon vials and diluted to 35 ppb before analysis using 0.5 M HNO3, and based on previous B concentration assessments on the Quadrupole ICP-MS.

2.4. Boron isotope analyses on MC-ICP-MS

The boron isotopic composition of brachiopod CaCO3 and CM was determined at GEOMAR, Kiel on a Thermo Scientific Neptune Plus MC-ICP-MS following procedures outlined in Foster (2008) with modifications. Prior to each analysis the instrument was tuned for maximum 11B/10B stability. We used an ESI PFA 50 μl min⁻¹ nebuliser and a Teflon barrel spray chamber. We did not find any significant improvement in washout when running the instrument with ammonia add gas; on the contrary, the addition of ammonia led to precipitation of ammonium nitrate in the Teflon spray chamber and the injector, thereby requiring cleaning of the sample introduction unit after each ~24 h of running. Machine-induced mass fractionation was corrected by standard-sample bracketing with a 35 ppb standard NIST SRM 951 boric acid that consequently converts 11B/10B ratios to delta notations. We aimed for matched sample to standard concentration, although within the range of ~10 to 50 ppb boron per sample the required precision and accuracy was achieved. Each analysis consisted of 120 seconds lasting simultaneous collection of masses 11 and 10 with Faraday cups H4 and L4 (both with 1012 Ω resistors), respectively. The typical signal intensity was 10–20 mV per 1 ppb on mass 11. Each sample was analysed twice during each analytical session with the average value reported. Airborne boron contamination to the open sample vials in the autosampler caused significant blank accumulation. In order to obtain the best possible control over the boron memory in the spray chamber, an on-peak zero was run over 30 seconds immediately preceding the take-up of the following sample or standard containing the same volume of 0.5 M HNO3 (typically 1.5 ml). Following this protocol background blank contribution as high as 10% could be corrected without significant negative impact on either the standard or the sample composition as quantified with secondary standards run under identical conditions throughout sequences. This allowed for a maximum length of one analytical session of approximately 11 h during which a sequence of 10 samples, 6 standards and corresponding 16 blanks were measured twice (total of 32 positions in the autosampler). The external reproducibility for δ11B values was better than ±0.2‰ (2σ) assessed by repeated measurements of purified coral standard reference material JCP-1 (Okai et al. (2002); n = 15, δ11B = 24.41 ± 0.18‰, in agreement with the certified value and other laboratories e.g. Aggarwal and You (2016)) and in-house M. venosa brachiopod standard MVS-1 (n = 46, δ11B = 15.95 ± 0.19‰). The long-term Neptune Plus stability and performance was monitored by regular measurements of pure boron BAM reference materials (Vogl and Rosner, 2001), resulting in −20.31 ± 0.28‰ (n = 58); 19.70 ± 0.20‰ (n = 123); and 39.67 ± 0.22‰ (n = 104) for ERM-AE120, ERM-AE121 and ERM-AE122, respectively (between years 2016–2018).

2.5. Boron isotope analyses by SIMS

High spatial resolution boron isotopic analyses were performed at CRPG-CNRS, Nancy using Cameca IMS 1270 ion microprobe. The analytical settings are described in Blamart et al. (2007). In summary, a primary mass-filtered beam of 16O ions with an intensity of 60–70 nA was focused into an aperture-delimited spot of approximately 40 μm using Kohler illumination. The analyses were conducted in a mono-collection mode on an electron multiplier by peak jumping between mass 9.8 (background), 10B and 11B. The deadtime of the electron multiplier was set at 0.5 M HNO3 is required) and carefully loaded on the chromatographic columns. Boron retention on the resin is pH dependent (see Lemarchand et al., 2002a); thus repeated rinses with Milli-Q water were necessary to elute the carbonate matrix and the buffer, while purified boron was collected with 720 μl of 0.5 M HNO3. All columns were individually calibrated with yields verified by a highly resolved elution scheme (consisting of minimum 30 steps), yielding >98% boron recovery. To ensure that no significant amount of the sample remained on the column, boron content of each elution tail was determined representing <1% of the total boron in the sample. Likewise, a total procedural blank (TPB) for each column batch was determined; all TPBs were below <100 pg (<0.1% of the sample size, n = 12). Purified boron samples were stored in Teflon vials and diluted to 35 ppb before analysis using 0.5 M HNO3, and based on previous B concentration assessments on the Quadrupole ICP-MS.

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(δ$^{11}$B = 16.0‰; see Section 2.4 above, which is a well-established value as it is also our standard for solution-based analyses MVS-1) in-between measurements of the shell parts grown in the cultures.

2.6. Boron isotope systematics

Boron has two isotopes, $^{10}$B and $^{11}$B (approximately 20% and 80% of total B, respectively), and the $^{11}$B/$^{10}$B variations are expressed in the standard delta notations in permil (%ε) relative to the reference material NIST SRM 951 boric acid (Catanzaro et al., 1970) as follows:

$$\delta^{11}B(%) = \left( \frac{11B(\text{sample})}{11B(\text{standard})} \right) \times 1000 - 1000 \quad (1)$$

The principle of the $\delta^{11}$B proxy is based on the speciation of boron in seawater. Boron is present in seawater almost exclusively as trigonal boric acid [B(OH)$_3$] and tetrahedral borate ion [B(OH)$_4$]$^-$. The relative proportion of boron species is pH dependent as defined by the following equilibrium:

$$\text{B(OH)}_3 + \text{H}_2\text{O} \leftrightarrow \text{B(OH)}_4^- + \text{H}^+. \quad (2)$$

Because of the differences in coordination and the subsequent B-O bond vibrational frequencies, an isotopic fractionation exists between the two species. Thus, as the distribution of boric acid and borate ion change with pH, so does their isotopic composition. The associated isotopic fractionation may be described as:

$$^{10}\text{B(OH)}_3 + ^{11}\text{B(OH)}_3 \leftrightarrow ^{11}\text{B(OH)}_3 + ^{10}\text{B(OH)}_4^- , \quad (3)$$

where the equilibrium constant $^{11-10}K_B$ (also termed as $\alpha_{4-3}$ or $\beta_B$ in the literature) is defined as:

$$^{11-10}K_B = \left( \frac{[^{11}\text{B(OH)}_3]}{[^{10}\text{B(OH)}_3]} \right) \times \left( \frac{[^{10}\text{B(OH)}_4^-]}{[^{11}\text{B(OH)}_4^-]} \right) \cdot \quad (4)$$

Given a value for $^{11-10}K_B$, the isotopic composition of both boron species in seawater varies predictably with pH (Kakihana et al., 1977). The experimentally derived value 1.0272 ± 0.0006 in seawater at 25 °C is currently the preferred isotope fractionation factor (Klochko et al., 2006), which was previously also supported by theoretical studies (Zeebe, 2005). Providing $^{11-10}K_B$ and $\beta_B$, the dissociation constant for boric acid at in situ temperature, salinity and pressure (commonly calculated according to Dickson, 1990), pH can be calculated from the $\delta^{11}$B values of either boron species, e.g. borate ion:

$$\text{pH} = \text{pK}_B - \log \left( \frac{\delta^{11}B_{\text{sw}} - \delta^{11}B_{\text{borate}}}{\delta^{11}B_{\text{sw}} - \left( ^{11-10}K_B \times \delta^{11}B_{\text{borate}} \right) - 10^6 \left( ^{11-10}K_B - 1 \right)} \right) \quad (5)$$

where $\delta^{11}B_{\text{sw}}$ is the isotopic composition of seawater (39.61‰; Foster et al., 2010), consistent on timescales of 11 to 17 Ma due to the long residence time of boron in the ocean (Pagani et al., 2005; Paris et al., 2010; Lemarchand et al., 2012b).

Owing to the use of artificial culture medium (CM), the $\delta^{11}B_{\text{sw}}$ should be replaced by $\delta^{11}B_{\text{CM}}$, as its isotopic composition was different to that of natural seawater and was closely monitored during the full duration of the culturing.

Although we observed a steady drift towards more isotopically depleted values over the culturing period (from approximately −3 to −6‰), at each given time the $\delta^{11}B_{\text{CM}}$ remained highly consistent (within the current analytical errors) in all culturing treatments (control and low-pH). Hence an exact comparison of the $\delta^{11}$B of newly built shell material at the margin was possible between specimens cultured under different pH conditions (control pH$_0$ vs. and low-pH conditions pH$_1$ or pH$_2$). Following this approach we could precisely isolate the $\delta^{11}$B-pH response and establish a calibration for brachiopods. In order to make our culture calibration applicable to natural seawater conditions a simple normalisation was required. First, an adjustment of the $\delta^{11}$Bborate in the cultures to $\delta^{11}$Bborate of the natural seawater using the measured pH and $\delta^{11}$Bsw (39.61‰) was required. Subsequently, the linear regression slopes of the culture calibrations (average $y = 0.29x$, Fig. 2) were fitted to the data of natural $M. \text{venosa}$ (which are well constrained: $\delta^{11}$B = 15.95‰, see Section 2.4; pH = 7.9 ± 0.1; see Section 2.1). This resulted in a single general calibration applicable to natural conditions (Fig. 3) without affecting

![Image](https://via.placeholder.com/150)
was comparable to natural seawater; around 0.5 mmol/l in contrast to 0.4 mmol/l measured for IAPSO.

2.7. Electron microprobe (EMP) analyses

JEOL JXA 8200 electron microprobe (EMP) mapping was used to assess elemental distribution and variations associated with shell microstructures. Quantitative elemental maps were obtained by wavelength dispersive x-ray spectrometry, simultaneously measuring Mg, Ca, Sr, Cl and S following Liebetrau et al. (2014) with modifications. In summary, the electron beam was focused to a spot size of 5 μm, accelerating voltage set to 15 kV and beam current to 100 nA. Mapping was repeated to gather 10 accumulations of the selected area at a step size of 5 μm and accumulation time of 15 ms, resulting in a resolution of 5 × 5 μm per pixel. Standard materials (Calcite, Sphalerite, Modern Coral-A2, Dolomite, KANI, Strontianite, VG-2, Scapolite) were measured before and after sample analyses to calculate absolute elemental concentrations, with uncertainties typically better than 2% relative standard deviation, 1σ).

2.8. Calculifying fluid pH measurements

Using H⁺-selective microelectrodes we conducted in vitro pH measurements in the calcifying fluid. The H⁺ microsensor construction and measurements were essentially performed as previously described in de Beer et al. (2008); Stumpp et al. (2012, 2013, 2015) and Ramesh et al. (2017). By plotting the voltage output of the probe against log[H⁺] values, a linear regression yielded a Nernstian slope of >54 mV per 1 pH unit. With this method we were able to resolve a minimum difference of 0.1 pH units. For this, M. venosa from control (pH0, n = 7) and low-pH (pH2, n = 7) treatments were collected from the culturing tanks at GEOMAR and immediately transported to Kiel University in closed jars to minimise the gas exchange between culture medium and atmospheric air. Microelectrode measurements were performed within hours and once completed the brachiopods were returned to the cultures. The measurements were limited to juvenile individuals (<1 cm), which possess a transparent shell that enables manoeuvring of the microelectrode between the shell and the mantle epithelium. Measurements were performed under visual control of the sensor tip on an inverse microscope (Axiovert 135, Zeiss, Oberkochen, Germany) equipped with a temperature controlled water bath adjusted to 10 °C. Each sample was introduced into the measuring chamber containing 5 ml of seawater from the respective pH treatment. The ion-selective probe was mounted on a remote-controlled micromanipulator (Phytron) and introduced from the shell margin through the mantle into the calcifying fluid where the ionic activities were recorded.

3. RESULTS

3.1. Brachiopod growth in cultures

Experimental conditions of the different treatments are summarised in Table 1; the cultured species are shown in

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Fig. 3. Mini-bulk δ¹ⁱB results normalised to δ¹¹Bsw = 39.61‰ (as described in Section 2.6); a) new calibration for brachiopods plotted in a δ¹¹Bmeasured − δ¹¹Bborate space. The black dashed line is the 1:1 relationship (δ¹¹Bmeasured = δ¹¹Bborate). The black circle shows the value for natural M. venosa. (b) Resulting δ¹¹B−pH relationship for brachiopods, with the measured data points indicated. (c) Comparison between our δ¹¹B−pH relationship and other curves from literature. The empty symbols indicate the typical spread as a result of varying degree of vital effects.
Fig. 1. For *M. venosa* (Fig. 1a), we did not observe a difference in the growth of new shell material between the control (pH0 = 8.0 and 8.15) and the low-pH treatments (pH1 = 7.6 and pH2 = 7.35) during either of the culturing phases. The typical growth rate of cultured *M. venosa* was about 1 mm per month, in line with observations made in nature (Baumgarten et al., 2013). Similarly, we noted that the growth rate was lower in larger individuals (>5 cm), possibly due to being close to reaching sexual maturity. The growth rate refers to simple measurements of the length of the shell valves (i.e. extension rate).

Information on *P. atlantica* regarding natural growth conditions is generally very limited and their growth rates are yet to be fully constrained (Álvarez et al., 2005; Logan, 2004). For the cultured organisms (Fig. 1c) it was not possible to distinguish any apparent new growth increments built during the culturing in either of the treatments by microscopic observations and calcein labelling. The only clue to their growth comes from SIMS δ11B analyses, which enabled us to identify shell parts precipitated from natural seawater and from the culture medium because of the pronounced differences in the respective δ11B values down to minimum ~20‰, implying that the area covered by the ion spot size corresponded to calcite almost entirely precipitated within the culture medium (Fig. 4a). Spots with more positive values, but lower than the δ11B of natural *P. atlantica* (δ11B = 19‰), implied that the measured shell area contained material precipitated partially from natural seawater and partially from culture medium. In comparison, intrashell variations were rather small, potentially resulting from the different microstructures of the shell (Fig. 4b). This suggests a growth rate of less than 40 μm (the SIMS spot size) in 1.5 years. Whether this extremely slow growth rate may be an artefact of the artificial environment or is also true for natural organisms remains unclear, but if real could suggest surprisingly long lifespans of these organisms considering the average adult size of approximately 1 cm. As a consequence of the minimal growth in cultures, this species was not used for further δ11B investigations.

3.2. Boron isotope calibration

Mini-bulk δ11B data from *M. venosa* shell increments precipitated under specific pH conditions (as described in Section 2.3 and shown in Fig. 1) were used to establish a δ11B-pH calibration equation. The measured δ11B values of our cultured brachiopods present a unique signature in contrast to natural brachiopods, because of the differences in our artificial δ11Bsw and the natural δ11Bsw (see Section 2.6). The initial δ11Bsw at the start of experimental phase 1 (P1) was ~3‰, with a progressive drift towards more depleted values throughout, resulting in ~6‰ at the end of the experimental phase 2 (P2). The reason behind this progressive depletion of δ11Bsw remains unclear, however we can exclude the successive refilling of evaporated volume by deionised water (δ11B > 80‰), as the boron background is expected to be heavy rather than light. Moreover, the B content of the deionised water source was regularly monitored (along with CM) on the ICP-OES and negligible (~0.005 mmol/l in deionised water against ~0.5 mmol/l in CM). This trend was highly consistent (within the current analytical errors) in all culturing treatments (control and acidification). By comparing the low-pH treatments to the respective controls we can isolate the δ11B-pH sensitivity for *M. venosa*. In Fig. 2 each pair of data points connected with a line represents the δ11B composition of a control and a low-pH sample at a given time of culturing. The offset between these lines illustrates the slowly decreasing δ11Bsw trend. All four calibration lines have highly consistent slopes (average y = 0.29x, Fig. 2), being controlled by the δ11B-pH sensitivity in *M. venosa*. By normalising the culture data to natural seawater and borate (δ11Bsw = 39.61‰, δ11B of natural *M. venosa* = 16.0‰; see Section 2.6, for more details), we generate a universal calibration for conversion of natural brachiopod δ11Bcal to δ11Bborate for pH-calculations (Fig. 3):

\[
\delta^{11}B_{\text{borate}} = \frac{\delta^{11}B_{\text{cal}}}{11(1.572 \pm 0.14)}
\]

The scatter in the data is minimal resulting in a close-fitting regression slope (R² = 0.98), and a clear response of δ11Bcal to the different δ11Bborate (Fig. 3a). Eq. (6) is shown with standard errors for the slope and intercept, determined using the MS Excel function LINEST. In contrast to the aqueous borate ion [B(OH)₄]⁻ to pH relationship
(dashed line), our brachiopod δ11B data show a much shallower slope, plotting above the 1:1 line at the lower pH-end, and slightly below at the higher pH-end (Figs. 2a and 3a). While this calibration is predominantly based on *M. venosa* specimens, the sparse data available on *T. dorsata* from one treatment overlap within errors with the former. The resulting sensitivity to pH is shown in Fig. 3b. Although our curve confirms a clear δ11B–pH dependency for brachiopods, in comparison to the other relationships from the literature it records a rather diminished response to seawater pH changes. At present analytical uncertainties this suggests a limited proxy range to minimum pH of approximately 7.5, reconstructions below this value could potentially lead to large uncertainties due to the flattening of the δ11B–pH curve. For instance, a reverse pH reconstruction from a sample from the lowest pH treatment (pH2 = 7.35) results in an apparent pH range within 7.1 and 7.5, entirely caused by the minimal analytical scatter on the order of below 0.2‰ (Fig. 3c).

### 3.3. Intrashell δ11B variations

Prior to discussing our results we briefly review the terminology and structure of brachiopod shell for guidance (schematic diagram is available in Fig. 5). Brachiopod shell is secreted by epithelial cells of the mantle (*Williams, 1966*) in an accretionary growth mode (*Ackerly, 1989*). New increments are added to the shell margin as well as to the growth surface lining the mantle, resulting in a progressive elongation and thickening of the shell (Fig. 5). Accordingly, a multi-layered skeleton is developed composed of calcite increments secreted at different stages of life. Hence, the posterior part of the shell is dominated by an ontogenetic trend, while the anterior part of the shell consists of the youngest increment. At a microstructural level, the shell of terebratelid brachiopods including *M. venosa* is differentiated into two layers (although some species with an additional tertiary layer also exist): a thin (only few tenths of μm thick) outer primary layer of granular structure and a fibrous secondary layer (Fig. 6; see also e.g. *Ye et al., 2018* for further details).

High spatial resolution data on δ11B distribution within selected growth increments and shell layers are presented in Fig. 7. SIMS spots were arranged in vertical profiles that traverse the different calcification increments, which can be distinguished from Mg concentration EMP maps. High Mg content (in mostly green and yellow colours) is apparent on the edges of each successive increment and traces the growth lines produced under different ambient conditions: nature and the two experimental culturing phases (P1 and P2). The observed Mg ‘banding’ is present in shell parts grown under natural as well as culture conditions, indicating that this feature is not caused by seasonal variations. Since similar Mg patterns are obvious for both the control and the low-pH specimen the influence of pCO2/pH changes on Mg distribution can also be excluded. Potentially, these Mg trends could be caused by growth related processes, which may lead to non-equilibrium effects during incorporation of Mg into the crystal lattice (*Cusack et al., 2008*). Increased Mg concentration is particularly apparent at the transition zones between the increments produced under different experimental conditions (indicated by arrows in Fig. 7), possibly as a result of short-term alteration of the precipitations rate due to stress from transport or calcine labelling.

The SIMS profiles trace the δ11B composition of the different increments and calcite layers from the outermost primary layer to the innermost secondary layer. An example of a typical SIMS profile within the shell microstructures is shown in the scanning electron microscope (SEM) image in Fig. 6. Based on an ontogenetic trend, the δ11B profiles can be separated into anterior (Fig. 7: profiles a1–a5 and b1–b5) and posterior profiles (Fig. 7: profiles a1–a2 and b1–b2). While the anterior profiles are located in a recent shell increment formed entirely in the cultures, the posterior profiles cross the multiple shell increments precipitated at different stages of growth, including nature and culture conditions.

In the anterior part of the shell of the control specimen (Fig. 7a) the δ11B decreases from isotopically heavier values...
in the outermost primary layer to lighter values in the innermost secondary layer (Fig. 7a: profiles a1–a5). In the low-pH specimen (Fig. 7b), the trend is opposite; the primary layer is relatively light while the secondary layer is more enriched in the heavy isotopes (Fig. 7b: profiles b1–b5). In contrast to the anterior shell margin, the posterior area is composed of multiple layers in both specimens (control and low-pH) as indicated by the EMP Mg maps, which results in more complex $\delta^{11}B$ profiles (Fig. 7: profiles a1–a5 and b1–b5). The $\delta^{11}B$ of the upper increments (older) precipitated under natural conditions (before culturing) is typically positive (about 16‰), whereas underneath the inner increments bear a negative $\delta^{11}B$ signal indicating that they were formed within the culture medium.

Fig. 7. Secondary ion mass spectrometry (SIMS) $\delta^{11}B$ profiles in cultured M. venosa: (a) cultured under control conditions (pH2), and (b) low-pH conditions (pH1 and pH2). Position of the SIMS $\delta^{11}B$ profiles in the shells is shown in black boxes on the electron microprobe (EMP) Mg maps. Mg distribution maps highlights growth features, with elevated Mg content (in green) tracing specific shell increments. Outer protuberances (indicated by arrows) mark transition zones between increments formed in nature and the different experimental culturing phases P1 and P2. In the panels with $\delta^{11}B$ data (a1–a5 and b1–b5) black circles show measurements located in the shell parts grown purely under culturing conditions (negative $\delta^{11}B$ values, top x-axis); blue circles represent measurements located in shell parts grown in nature prior to culturing (positive $\delta^{11}B$ values, bottom x-axis). The pronounced difference between culture and nature $\delta^{11}B$ of the shell is due to precipitation from solution of different $\delta^{11}B$: $\delta^{11}B_{CM}$ vs. $\delta^{11}B_{sw}$ (see Section 2.6 for more details). Filled symbols illustrate measurements located entirely in the secondary layer, empty symbols in the primary layer, and half-filled show the spots with significant contribution of both layers.
3.4. Calcifying fluid pH

Microelectrode measurements of pH in the calcifying fluid between the shell and the mantle tissue (Fig. 8a and b) were carried out to evaluate the internal pH-regulatory capacity of *M. venosa*. Our results are summarised in Fig. 8c, and indicate a relatively narrow pH range in the calcifying fluid, with a median around ~7.8 (Fig. 8c) for individuals cultured under control (pH₀ = 8.15) as well as low-pH (pH₂ = 7.35) conditions. The box plots are based on measurements in seven brachiopods from each treatment and thus reflect the variance within individuals. We note that these measurements were performed outside of the culturing tanks, and thus a slight offset (~0.1 pH units) in the pH of the CM measured in culturing facilities (as given in Table 1) and under the microscope setup using microelectrodes (particularly at low-pH) was observed due to CO₂ degassing and different ambient conditions.

4. DISCUSSION

4.1. Boron isotope composition of brachiopods

Brachiopods have been considered faithful δ¹¹B recorders, as their composition apparently reflects the seawater borate ion. Measurements of natural samples have shown that δ¹¹B values of modern samples are within the range of foraminifers and thus could be used for palaeo-reconstructions (Lécuyer et al., 2002). Leaving the culturing experiments aside for now, our measured δ¹¹B of natural *M. venosa* and *P. atlantica* samples already indicate the feasibility of the δ¹¹B – pH proxy in brachiopods (Table 2). For instance, at pH of 8 and δ¹¹Bsw ~39.6‰, the δ¹¹Bborate is expected to be ~15.7‰. Providing that only borate ion is incorporated into brachiopods, as largely expected for biogenic carbonates (e.g. Hemming and Hanson, 1992), the δ¹¹B of brachiopod shells should mimic the seawater borate. Our measurements of natural *M. venosa* give a δ¹¹B ~16.0‰, which agrees very well with the expected borate ion values (~15.7‰). A similar exercise can be applied to *P. atlantica*, using the pH and ambient conditions as provided in Table 2, the δ¹¹Bborate of local seawater is expected to be ~19.1‰ while the *P. atlantica* give a δ¹¹B ~18.9‰, highly consistent within the uncertainties. Moreover, both species are morphologically very different, possess distinct biomineralisation mechanisms and have contrasting growth rates. This supports the notion that only borate ion appears to be incorporated into brachiopod calcite. Potentially, it could also indicate that inter-specific differences are not too large and do not obscure the seawater δ¹¹Bborate, but this needs further confirmation with other modern species. We note, however, that for a definite eval-

![Fig. 8. Microelectrode pH measurements in the calcifying fluid. Left panels: photographs taken from below of the *M. venosa* specimen (anterior part of the valve) through an inverted microscope; (a) the electrode is pressed into the mantle and is in measurement position between the shell and the epithelium. SH–shell (outside), EM–epithelial mantle, ME–microelectrode (V-shaped contour). Right panel: (c) Measured pH of seawater (SW) and calcifying fluid (CF) in pH₀ (control) and pH₂ (low-pH) treatments. The minimum and maximum values are indicated by the whiskers, lower and upper boxes show the third and first quartiles, respectively with a thicker median line between them (n = 7 for each box plot). (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)](image)

<table>
<thead>
<tr>
<th>Species</th>
<th>Location</th>
<th>Sal.</th>
<th>T. [°C]</th>
<th>pH</th>
<th>δ¹¹B [%‰]</th>
</tr>
</thead>
<tbody>
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<td><em>Magellania venosa</em></td>
<td>Comau Fjord, Chile</td>
<td>30</td>
<td>10</td>
<td>7.9</td>
<td>15.95</td>
</tr>
<tr>
<td><em>Pajaudina atlantica</em></td>
<td>La Palma, Canary Islands</td>
<td>36</td>
<td>21</td>
<td>8.1</td>
<td>18.86</td>
</tr>
</tbody>
</table>
ulation a good knowledge on seawater pH is critical, which is not straightforward in nature. Ultimately, it will be difficult to convincingly rule out presence of boric acid in the calcite until these measurements are attempted, but at present all points towards borate ion being the only species incorporated into brachiopod shells.

4.2. Effect of acidification on brachiopods

Reduced skeletal growth due to CO₂-induced ocean acidification and associated changes in calcium carbonate saturation state of seawater \((Ω_{\text{calc}})\) has been well documented in a broad range of marine organisms including corals, molluscs, foraminifera, coccolithophores, or coralline algae (e.g. Gazeau et al., 2007; Fabry et al., 2008; Silverman et al., 2009; Ries et al., 2009; Kroeker et al., 2013), yet only little is known on the effect on brachiopods. During our experiments, all cultured brachiopods survived without obviously affected growth prolonged acidification such as that predicted for the end-century surface ocean \((pH_1 = 7.6; \text{Table 1; e.g. Brewer, 1997})\), or even much more extreme conditions involving further reduction of seawater pH \((pH_2 = 7.35)\) and calcite saturation state \((Ω_{\text{calc}} = 0.6)\). This points to a strong biological control on the shell formation and the internal pH of *M. venosa*. Potentially, similar controlling mechanism may also be present in *T. dorsata*, and *P. atlantica*, although this could not be fully evaluated by the present study due to the lack of individuals of the first species, and elusive growth of the second. Unaffected survival and shell building under acidification has also been reported for the Antarctic brachiopod *Liothyrella uva* (Cross et al., 2015) and the New Zealand *Calloria inconspicua* (Cross et al., 2016). It appears that several brachiopod species, and especially terebratellids, may not be negatively affected by regimes with reduced pH, or at least the effect of low-pH alone. Experiments combining acidification with additional stressors such as temperature increase or nutrient limitation indicate that it is the synergic effect of stressors that is particularly detrimental to physiological processes of calcifying organisms (e.g. Reynaud et al., 2003; Holcomb et al., 2010). These experiments present a more representative scenario for natural systems under increasing anthropogenic pressures, and thus it remains highly questionable whether brachiopods will be able to accommodate to these changes despite the apparent robust pH-buffering capacities.

4.3. Vital effects

The survival and calcification of *M. venosa* under acidified conditions and at thermodynamically unfavourable conditions with respect to calcite precipitation \((Ω_{\text{calc}} < 1)\), point towards a strong biological control over shell growth. This is also evident from the shallow slope of our brachiopod boron isotope fractionation line in comparison to the expected aqueous \(δ^{11}B_{\text{borate}}\) composition (Fig. 2a and a), and the resulting moderate response of \(δ^{11}B_{\text{calcite}}\) to pH changes (Fig. 3b and c). The calibration crosses the 1:1 line \((\delta^{11}B_{\text{measured}} = \text{aqueous } δ^{11}B_{\text{borate}})\) recording similar \(δ^{11}B_{\text{calcite}}\) than that of aqueous \(δ^{11}B_{\text{borate}}\) at normal culture medium pH, but higher \(δ^{11}B_{\text{calcite}}\) at acidified conditions (Fig. 3a). This suggests a reduced pH range at precipitation, which is likely caused by regulation of their calcifying fluid. To determine whether internal physiological pH-buffering may be responsible for the observed trend we performed *in vivo* calcifying fluid measurements at the interface of the epithelial mantle and the shell as shown in Fig. 8 on several *M. venosa* individuals from control \((pH_0)\) and low-pH \((pH_2)\) treatments. The calcite formation is anticipated to occur at the mantle-shell interface via ion-transporting channels in a fluid-filled medium – which is here referred to as the calcifying fluid – of μm to sub-μm dimensions. Our results indicate a rather narrow pH range in the calcifying fluid \((\sim 7.8, \text{Fig. 8e})\), below normal seawater conditions, which does not appear to be affected by the pH of the culture medium, or at least could not be resolved considering the variance in the date. Although these measurements were performed on juvenile organisms \((<1 \text{ cm})\), we do not expect the pH to differ significantly in the typical adults \((2-5 \text{ cm})\), since the growth rates of *M. venosa* are relatively constant over the different life stages (with the exception of the oldest organisms >5 cm that have lower growth rates; Baumgarten et al., 2013). This demonstrates that *M. venosa* are able to strictly control their calcifying fluid pH, leading to stable pH conditions at the calcification front even in an environment with low-pH conditions. Similar phenomenon has also been reported for corals (Georgiou et al., 2015; Wall et al., 2015), and our study now confirms it for brachiopods.

What remains to be answered is why would brachiopods calcify at such low-pH, when compared to most of the other calcifiers. For instance, the cold-water coral *Desmophyllum diathus* from the same habitat as *M. venosa* in Chile has a calcifying fluid pH of \(\sim 8.7\) (McCulloch et al., 2012b), much higher than seawater. Since brachiopods are an evolutionary old taxon (earliest known brachiopods occur in the lowermost Cambrian, Holmer and Popov, 1996) it could be that their calcification mechanisms differ from many other calcifiers and potentially bear a link to ancient seawater chemistry. Seawater pH estimates for the early Phanerozoic predict values as low as \(\sim 7.4\) (Halevy and Bachan, 2017). Early ancestors would have to accommodate to these environmental conditions, and calcification pathways requiring substantial up-regulation of calcifying fluid pH with respect to seawater would not pose a physiologically efficient mechanism. The problem of low seawater pH can be overcome if the preferred substrate for calcification is bicarbonate, and not carbonate, which is also the dominant DIC species at the respective pH. Relevant enzymatic systems such as carbonic anhydrases catalyse the reversible hydration of carbon dioxide into bicarbonate and free protons: \(\text{CO}_2 + \text{H}_2\text{O} \rightleftharpoons \text{HCO}_3^- + \text{H}^+\) and could present a highly effective mechanism for calcification under low-pH/high-DIC conditions (Moya et al., 2008; Bertucci et al. 2009; Khalifah, 1971). This could explain the ability of *M. venosa* and as well as other terebratellids to thrive in low-pH environments without hampered shell growth.

In summary, we identify vital effects on \(δ^{11}B\) in brachiopods as internal pH-regulation. Taking into account the reduced \(δ^{11}B\)-pH sensitivity (Fig. 3), it is apparent that the mini-bulk \(δ^{11}B\) composition of the shell, integrates to a
certain (and potentially large) extent the calcifying fluid pH. The distortion of the ambient pH signal (deviation from the aqueous $^{11}\text{B}_{\text{aq}}$) thus seems to be primarily caused by homeostasis. Further work and especially refining the uncertainties of the calcifying fluid measurements is required for a more conclusive evidence, but if true it would mean that $^{11}\text{B}$ in brachiopods fully reflects the pH of solution from which it precipitated. This would imply that incorporation of boron into brachiopod calcite follows the inorganic fractionation relationship (Klochko et al., 2006), and that no boric acid is co-precipitated, which is rather unexpected for a calcite matrix (e.g. Noireaux et al., 2015).

4.4. Boron isotope composition of shell microstructures

Above we directly and indirectly demonstrate that brachiopods regulate their calcifying fluid pH, which appears to be to a certain extent probed by the (mini-bulk) $^{11}\text{B}$. Still, despite the confined calcification pH range, a $^{11}\text{B}$–pH dependency exists. This raises a question on the underlying basis of the $^{11}\text{B}$–pH dependency, which could be caused by two potential mechanisms. The first possibility would be that the brachiopod calcifying fluid is systematically impacted by ocean acidification, as apparently true for corals (Hönisch et al., 2004; Reynaud et al., 2004; Krief et al., 2010; Venn et al., 2013; Gagnon, 2013; Wall et al., 2016; Stewart et al., 2016), but the effect is relatively minor and could not be resolved by our calcifying fluid pH measurements. An alternative or additional option would be that the ambient seawater pH signal is, at least partially, attained in specific parts of the calcite (in this case shell layers), and cannot be detected via mini-bulk or bulk approaches. To reconcile the interpretation of the (mini-bulk) $^{11}\text{B}$ data we examine the $^{11}\text{B}$ composition of M. venosa shell microstructures from SIMS measurements (Fig. 7). Description of the brachiopod shell structure is provided in Section 3.3. Particularly relevant are also Figs. 5 and 6, which illustrate the shell features discussed in this section.

The shell of M. venosa is composed of two calcite layers with different microstructures: an outer primary layer of granular structure and a fibrous secondary layer (Fig. 7). Differences in boron and other isotope system compositions of the two layers have been documented, attributed to distinct biomineralisation processes during the formation of the layers. In brachiopods from natural settings, a shared trend was observed where the primary layer was isotopically lighter in contrast to the secondary layer, which was closer to equilibrium with seawater (Penman et al., 2013; Cusack et al., 2012; Bajnai et al., 2018; Romanin et al., 2018). Strikingly, this is not apparent in the cultured brachiopods and our results indicate that the respective direction of the trend principally depends on the ambient seawater pH. Progressive depletion in $^{11}\text{B}$ from the outermost primary layer to the innermost secondary layer in the control specimen (Fig. 7a: profiles a1–a2) and enrichment in the low-pH specimen (Fig. 7b: profiles b1–b2) indicate that biomineralisation differences between the primary and the secondary layer do not seem to significantly affect the $^{11}\text{B}$. Instead, the intrashell variations may be explained by a physiological gradient produced by an interplay of two end members of different pH, the calcifying fluid and the seawater, the first driving the $^{11}\text{B}$ composition of the innermost and the latter of the outermost layer. This would imply that kinetic effects resulting from the different biomineralisation modes between primary and secondary layer do not significantly affect $^{11}\text{B}$ values, or at least cannot be resolved with the present analytical error of SIMS. This is also in line with what is expected for the boron isotope system, considering the relative slow growth of brachiopods and the fast equilibration of boric acid with borate ion ($\sim$125 µs; Zeebe et al., 2001). More recently, however, precipitation experiments using synthetic calcite found growth rate effects on boron mineral-fluid partitioning and fractionation (Gabitov et al., 2014; Mavromatis et al., 2015; Uchikawa et al., 2015; Kaczmarek et al., 2016).

In the posterior region, where the shell structure is dominated by an ontogenetic trend the $^{11}\text{B}$ profiles enable us to trace the calcite sheets precipitated from solutions of different $^{11}\text{B}$ composition – $^{11}\text{B}_{\text{CM}}$ vs. $^{11}\text{B}_{\text{sw}}$ (Fig. 7: profiles a1–a2, b1–b2). Together with the Mg distribution maps, this permits us to identify the origin of each of the shell increments and illustrates the accretionary growth mechanism. The oldest shell part precipitated from natural seawater forms the outermost layer and has a composition of $\sim$16‰ (Fig. 7: profiles a1–a2 and b1, shown in blue symbols). Below, the intermediate shell layer was formed at the onset of the culturing experiments and was built from initial $^{11}\text{B}_{\text{CM}}$ ($\sim$3‰). The final increment is the youngest and has the lightest $^{11}\text{B}$ values, implying that it was built towards the end of the culturing experiments from most depleted $^{11}\text{B}_{\text{CM}}$ ($\sim$6‰). A particularly interesting feature of these profiles are the $^{11}\text{B}$ values of about $-13\%$ measured in the acidification specimen (Fig. 7b: profiles b1–b2). This composition may only be achieved by mixing of both, nature and culture grown components, yet the $-13\%$ values were measured in an increment built entirely under culture conditions. Terebratelids have been found to possess robust repair processes (Cross et al., 2015, 2016), which could explain the observed value, providing that mobilisation of older material was also involved.

5. CONCLUSIONS

This study provides novel direct and indirect evidences on the internal pH-controlling mechanisms of the brachiopod M. venosa, which potentially may also be relevant for other brachiopod species, in particular T. dorsata or other terebratelids. M. venosa regulate their calcifying fluid pH, which is maintained at a narrow pH range below normal seawater pH, suggesting the use of bicarbonate ion as the preferred calcification substrate. This physiological
adaptation makes these calcifiers resilient to CO₂-driven ocean acidification and explains their ability to survive in low-pH conditions, even at Qcrit < 1 without hampered shell growth. Presumably, this mechanism developed early during metazoan evolution and played an important role in determining brachiopod survival through the multiple Phanerozoic environmental crises. We establish a δ¹¹B-pH relationship for brachiopods, which supports its application as an archive for palaeo-pH and palaeo-CO₂ reconstructions. Our culture-based calibration, however, shows a more attenuated δ¹¹B-pH response than previously thought due to the strong role of biological processes over shell formation. This results in a reduced dynamic proxy range for brachiopods, demanding caution as small δ¹¹B variations in the calcite matrix would lead to potentially large differences in the calculated pH values, particularly at very low-pH conditions. In contrast to the clear pH-dependency confirmed by mini-bulk data, at a micro-scale level we observe marked δ¹¹B intra-shell heterogeneities. While an ontogenetic trend certainly plays an important role, at present it appears that δ¹¹B variations between the outermost and the innermost layers may be explained in context of a physiological gradient across the shell. Accordingly, boron incorporation into the innermost shell layers is primarily driven by calcifying fluid pH, whereas the outermost calcite layers are to some extent influenced by ambient seawater pH. Increased analytical precision is needed before a conclusive statement can be made, but if true, this may suggest rather unexpected δ¹¹B fractionation for a biogenic calcite. In addition, this could offer an explanation for the basis behind the mini-bulk δ¹¹B-PH dependency, although it is also possible that calcifying fluid pH is systematically impacted by ocean acidification, which could contribute to the δ¹¹B-PH trend. Finally, we would like to encourage further research on brachiopods involving controlled culturing experiments under both, laboratory and/or mesocosm conditions, which clearly proves invaluable for advancing our understanding on the geochemistry, biology as well as physiological adaptations of these marine calcifiers to cope with environmental change.

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REFERENCES


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