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Resistance to freshwater exposure in White Sea *Littorina* spp. II: Acid-base regulation

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Abstract Parameters of acid-base and energy status were studied by in vivo ³¹P-nuclear magnetic resonance spectroscopy in three White Sea Littorina spp. (L. littorea, L. saxatilis and L. obtusata) during prolonged anaerobiosis in freshwater. Intracellular pH decreased significantly, especially during the early period of anaerobiosis, but later the decrease in intracellular pH slowed down considerably, suggesting a capacity for intracellular pH regulation in all three species. There was a trend for intracellular pH to fall most rapidly in the least freshwater-resistant species, L. obtusata, as compared to the most resistant, L. littorea. Non-bicarbonate, non-phosphate buffer values estimated by the homogenate technique were similar in the three studied species (28–37 mmol $pH^{-1} kg^{-1}$ wet weight) and did not change during freshwater exposure. The CaCO₃ buffer value of the foot tissues was considerably higher (171-218 mmol pH⁻¹ kg⁻¹ wet weight) and decreased significantly during freshwater exposure. The contribution of the multiple tissue buffering systems to intracellular pH regulation in Littorina spp. shifts between different stages of freshwater exposure. Initially, the non-bicarbonate, non-phosphate tissue buffering system seems to be of major importance for metabolic proton buffering

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¹Alfred-Wegener-Institute for Polar and Marine Research, Ecophysiology and Ecotoxicology, Columbusstr., 30, D-27568, Bremerhaven, Germany e-mail: ISokolova@awi-bremerhaven.de Tel.: +49-471-48311-311; Fax: +49-471-48311-149 at intracellular pH between 7.5 and 7.0. During later stages of anaerobiosis and at lower intracellular pH, the $CaCO_3$ buffer is involved in proton buffering. Decrease in the $CaCO_3$ buffer value during freshwater exposure was in quantitative agreement with the amount of metabolic protons buffered, thus suggesting that $CaCO_3$ tissue stores may serve as a major buffering system during prolonged anaerobiosis in *Littorina* spp.

Key words Salinity stress \cdot Anaerobiosis \cdot Intracellular pH \cdot Tissue buffer values \cdot ³¹P-NMR

Abbreviations ANOVA analysis of variance $\cdot ATP$ adenosine-5'-triphosphate $\cdot F$ fraction of protonated phosphate $\cdot NMR$ nuclear magnetic resonance $\cdot NTA$ nitrilotriacetic acid $\cdot PCO_2$ partial pressure of CO₂ $\cdot PD$ phosphodiesthers $\cdot pH_i$ intracellular pH $\cdot P_i$ inorganic phosphorus $\cdot pKa$ apparent dissociation constant $\cdot PLA$ phospho-L-arginine $\cdot S/N$ ratio signal-to-noise ratio $\cdot W_{tiss}$ tissue wet weight $\cdot \beta_{CaCO_3}$ calcium carbonate tissue buffer value $\cdot \beta_H$ non-bicarbonate buffer value of tissue homogenate $\cdot \beta_{NB,NPi}$ nonbicarbonate, non-phosphate tissue buffer value $\cdot \beta_{Pi}$ inorganic phosphate tissue buffer value $\cdot \Delta H^+_{non-resp}$ non-respiratory proton change $\cdot \Delta H^+_{met}$ metabolic proton change $\cdot \Delta pH$ change of pH $\cdot \Delta pH_{obs}$ observed change of pH

Introduction

Many intertidal animals, including gastropods, possess an outstanding ability to withstand and successfully adapt to a wide range of salinity fluctuations. If salinity changes strongly and rapidly, a very typical adaptive behavioural response is evoked in intertidal gastropods. Snails withdraw into the shell and isolate themselves inside by closing the operculum. Isolation minimises the exchange of water and salts between the internal medium of the animal and the osmotically unfavourable environment (e.g. Berger 1986). However, many physiologically important functions like excretion, feeding, and reproduction are suspended during this period. In addition, animals remain practically anoxic during such isolation periods, explaining why this adaptive behavioural response results in only limited survival of the snails (Sokolova et al. 2000).

Prosobranch gastropods Littorina littorea, L. saxatilis and L. obtusata are common inhabitants of the White Sea intertidal zone. All three species are known to exhibit a fairly high resistance to extreme hypoosmotic stress and may survive from several days to more than a week in freshwater depending on environmental temperature (Berger 1986; Sokolova et al. 2000). This ability may be adaptive during the periods of spring ice-melting in the White Sea when the salinity of the surface water drops to < 2% in near-coastal areas for about 2 weeks (Babkov and Lukanin 1985). However, significant differences exist between the three Littorina spp. with respect to freshwater resistance. The low-shore periwinkle L. obtusata is most vulnerable to extreme hypoosmotic stress, whereas the resistance of the highshore L. saxatilis, and especially that of the subtidal species L. littorea is considerably higher. Our companion study (Sokolova et al. 2000) clearly demonstrated for the first time that exposure to an extremely hypoosmotic medium evokes anaerobic metabolism in the periwinkles evidenced by a large degree of succinate accumulation. Inability of anaerobic energy production to keep pace with energy demand leads to an impaired energy status of the tissue, reflected in a depletion of phosphagen stores and ATP levels, resulting in a reduction in the Gibbs free energy of ATP hydrolysis. Our data suggested that differential survival of the periwinkles under conditions of extreme hypoosmotic stress depends on their capacity to delay adverse changes in the energy status, which in turn is crucially related to the efficiency of metabolic rate depression and the ability to reduce ATP turnover during environmental anaerobiosis.

Besides the depression of metabolic and ATP turnover rates, other mechanisms, which can potentially contribute to anoxia resistance of the periwinkles, should not be neglected. Among them, the mechanisms of acid-base regulation preventing extreme acidification of the intracellular milieu during prolonged anaerobiosis are of primary importance. A decrease of intracellular pH (pH_i) may have adverse effects on enzyme functions and intracellular energy transductions (Pörtner 1987a, 1993). Although an extreme drop in pH_i would have deleterious effect on cellular functions, a moderate decrease may be advantageous during anoxia, providing a necessary background for slowing down enzymatic reactions and for improving protein stability, hence contributing to metabolic arrest (Juretschke and Kamp 1990; Kamp 1993; Pörtner 1993; Hand and Hardewig 1996; Schmidt and Kamp 1996).

To obtain a more complete picture of the mechanisms of freshwater and anoxia resistance, changes in the parameters of acid-base status were analysed by in vivo

³¹P-nuclear magnetic resonance (³¹P-NMR) and tissues homogenate techniques in White Sea gastropods L. littorea, L. saxatilis and L. obtusata. Changes in acidbase parameters were compared to anaerobic metabolic rates and mortality of the Littorina spp. studied in order to test whether the capacity for acid-base regulation contributes to differential survival of the snails under conditions of prolonged anoxia during extreme hypoosmotic stress. This approach also provides insight into the importance of acid-base parameters in metabolic regulation and vice versa. The buffer capacities of the multiple tissue buffering systems (non-bicarbonate, non-phosphate, inorganic phosphate and calcium carbonate) were compared in order to estimate the relative contribution of these systems to pH_i regulation at different stages of environmental anaerobiosis in Littorina spp.

Materials and methods

Sampling of Littorina spp.

Sampling for determination of tissue buffer capacities was performed in August 1997 in the Chupa Inlet of the Kandalaksha Bay of the White Sea (66°20' N, 33°40' E) at a water temperature of 10– 15 °C and 25–27‰ salinity. Details of the sampling procedure and the maintenance of the animals were as described elsewhere (Sokolova et al. 2000). Only adult snails with shell diameters of 7– 11 mm (*L. saxatilis* and *L. obtusata*) or 22–26 mm (*L. littorea*) were used in the experiments.

For ³¹P-NMR studies, adult *Littorina* were collected in October 1997 at the same sites (water temperature 5–7 °C, 26–27‰ salinity) and transported to the Alfred-Wegener Institute for Marine and Polar Research (Bremerhaven, Germany). Transportation took 2.5 days and all snails arrived alive. In the Alfred-Wegener Institute, snails were placed in a recirculating aquarium with natural North Sea water diluted with tap water to obtain a salinity of 26‰. Water temperature was maintained at 7.0–7.5 °C. Under these conditions snails were kept without feeding for 11 days prior to experimentation. Shell diameter of the periwinkles used in ³¹P-NMR studies was 8–12 mm for *L. saxatilis* and *L. obtusata* and 15– 27 mm for *L. littorea*.

Incubation procedure

Experimental incubation and tissue sampling for determination of tissue buffer capacities (August 1997) were performed at the White Sea Biological Station (Cape Kartesh, White Sea). Temperature of incubation was 7.7 \pm 0.6 °C. Control animals were kept in aquaria with aerated natural seawater of 26–27‰. Experimental animals were placed in plastic trays and covered with natural freshwater from the nearby Krivoye Lake. The water was changed daily. After specified exposure periods a portion of snails was taken out of the tray. Scoring for mortality and tissue sampling were performed as described elsewhere (Sokolova et al. 2000). Deep-frozen tissues were transported at liquid N₂ temperature to the Alfred-Wegener Institute where all analyses were performed.

For ³¹P-NMR studies the periwinkles were placed in separate 200-ml plastic bottles covered only with cotton mesh. Each sample consisted of 7 or 8 individuals (*L. littorea*) or 70–75 individuals (*L. saxatilis* and *L. obtusata*). Bottles were submerged into well-aerated aquaria with 26% seawater at 7.0–7.5 °C and left overnight in order to reduce the effect of handling. For the measurements the bottles were removed and filled with 80 ml of 26 ‰ seawater at the same temperature. The cotton mesh was gently replaced by a plastic grid to prevent movement of the snails during the NMR measurements.

After analysis of control values the bottles were drained, the plastic grid removed and the bottles refilled with fresh tapwater, covered with cotton mesh and fixed in the aquarium with recirculating aerated freshwater at 7.0–7.1 °C. After specified exposure periods, samples were taken out of the aquarium, drained and checked for the presence of dead snails. To reduce the effect of handling, poking by a needle was not applied (see Sokolova et al. 2000) and only those snails that failed to keep the operculum tightly closed were scored as dead. The final scoring after the experiment revealed that the error of this mortality estimate was below 2% for *L. saxatilis* and *L. obtusata* and was 0% for *L. littorea*. Dead periwinkles were discarded. The NMR measurements in freshwater-exposed animals were performed under 80 ml of freshwater. The whole procedure was repeated after every incubation period.

For each species, five samples were measured repeatedly as controls and at different exposure periods (3, 5 days of freshwater exposure for *L. obtusata* and 3 days, 5 days and 7 days for *L. saxatilis* and *L. littorea*). For *L. littorea*, ³¹P-NMR spectra were obtained for five independent additional samples after 7 days and 9 days of freshwater exposure.

³¹P-NMR measurements

For in vivo ³¹P-NMR measurements the bottles were fixed onto a 5.0-cm-diameter ¹H/³¹P/¹³ C surface coil and placed inside the magnet. The temperature was balanced via the water cooled gradient coil of the magnet (8 °C). All experiments were performed at 81 MHz using a 47/40 Bruker Biospec DBX system with actively shielded gradient coils. In vivo ³¹P-NMR spectra were recorded with a sweep width of 4000 Hz, 60°bp-pulses of 50 µs and a repetition time of 1 s for 600 scans resulting in an acquisition period of 10 min and a total measurement period of 20–30 min. All data were zero filled to 16 K and a line-broadening of 20 Hz was applied before Fourier transformation. Chemical shift values were expressed relative to phospho-L-arginine (PLA).

Mean pH_i of the whole animal was determined by the chemical shift of the inorganic phosphorus (P_i) signal in the NMR spectra and calculated with a formula of a pH titration curve from muscle fibres of the horseshoe crab *Limulus polyphemus* (Doumen and Ellington 1992) and corrected for temperature according to Pörtner (1900). Intracellular free magnesium concentrations were also determined from ³¹P-NMR spectra as described by Doumen and Ellington (1992). We assume that both pH and free magnesium concentration reflect intracellular values of these parameters because the P_i concentration in haemolymph of marine gastropods is too low to become visible in NMR spectra (Kinne 1971; Berger 1986).

Integration of phosphorus compounds in the NMR spectra were obtained with the standard integration routine of the processing software (xwin-nmr, Bruker, Ettlingen). The typical line width of phosphorus signals was less than 40 Hz in control animals as well as in the spectra obtained under hypoosmotic stress conditions.

Determination of tissue buffer capacity

Tissue buffer capacity was determined in vitro by a method following the rationale of Pörtner (1990). Tissue samples were ground under liquid N₂. Frozen tissue powder (1–2 g) of were placed into a preweighed tonometer vessel precooled on ice. To obtain this amount of tissue powder, foot muscles from 3 (*L. littorea*) or 28–30 (*L. saxatilis* and *L. obtusata*) individuals were pooled. Four millilitres of ice cold medium consisting of 0.54 mol 1^{-1} KF and 0.01 mol 1^{-1} nitrilotriacetic acid (NTA) were added for each gram of tissue powder. The mixture was thoroughly stirred and allowed to thaw.

The tissue homogenate was placed into a tonometer (Instrumentation Laboratory, Padorno Dugano, Italy) and equilibrated for 20–30 min with normoxic humidified gas mixture at a partial

pressure of CO₂ (PCO₂) of 3.6 torr provided by a gas mixing pump (Wösthoff, Bochum, Germany) at 8 °C. Homogenate pH was continuously monitored using a fluoride-resistant pH electrode GAT IJ 42 (Gamma Analysentechnik, Bremerhaven, Germany) calibrated with WTW (Weilheim, Germany) precision buffers. The initial homogenate pH ranged between 8.4 and 8.6 and was considerably higher than the pHi measured in vivo by ³¹P-NMR probably due to the presence of calcium carbonate stores in the foot of the periwinkles (Mason et al. 1984; Brough and White 1990). Therefore, the homogenate was titrated back to approximately the respective average pH_i values determined by ³¹P-NMR (see below) using 3 mol 1^{-1} HCl. During titration, the homogenate was equilibrated with a normoxic gas mixture at $PCO_2 = 3.6$ torr. After HCl titration intracellular buffer value was determined using the CO₂ titration method as described by Pörtner (1990). Normoxic gas mixtures with three partial pressures of CO₂ were used: low (0.75 torr), medium (3.62 torr) and high (7.6 torr). At the end of experimentation, aliquots of the homogenate were withdrawn and subjected to ultrafiltration at 0-4 °C using Millipore (Bedford, USA) Ultrafree PF units (30 000 d) for 50 min. PCA extracts of the ultrafiltrates were prepared, and inorganic phosphate concentration was determined enzymatically (Pörtner 1990).

Calculations of non-bicarbonate buffer values in the tissue homogenate ($\beta_{\rm H}$) followed the procedure outlined by Heisler and Piiper (1971):

$$\beta_{\rm H} = \frac{0.5 \times (\text{Bic}_1 + \text{Bic}_3) - \text{Bic}_2}{0.5 \times (\text{pH}_1 + \text{pH}_3) - \text{pH}_2} (\text{mmol pH}^{-1} \, \text{l}^{-1} \, \text{homogenate water})$$
(1)

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where indices 1 and 3 refer to values at medium PCO_2 , and index 2 to values at high or low PCO_2 . In order to evaluate the non-bicarbonate, non-phosphate buffer value ($\beta_{\text{NB,NPi}}$)of the homogenate, the buffer value of inorganic phosphate (β_{Pi})was subtracted from β_{H} of the homogenate as proposed by Pörtner (1990). The buffer value of fluoride was not considered in the calculations because the apparent dissociation constant of hydrofluoric acid (2.8–3.2) in the given range of temperature and ionic strength is far enough from the pH of the tissue homogenate to have only negligible influence on its buffer value (Pörtner 1990). The resulting value was corrected by the dilution factor of the tissue in the homogenate water (including medium and tissue water, the amount of water added with HCl and derived from neutralisation of calcium carbonate) to obtain $\beta_{\text{NB,NPi}}$ of the tissue (mmol pH⁻¹ kg⁻¹ wet tissue weight).

Unfortunately, the exact determination of total CO₂ in samples by the gas chromatography technique was impossible due to the high mucus content of the tissue homogenate which interfered with the measurements. Therefore, the solubility coefficient of CO₂ and the bicarbonate concentration in the homogenate were calculated using temperature, total molarity and PCO₂ according to the formula proposed by Heisler (1986). Tissue concentrations of Na⁺ and K⁺ (126 mEq g^{-1} and 94 mEq g^{-1} wet tissue weight, respectively) for Littorina spp. were adopted from a study on the foot muscle of White Sea L. littorea (Mikhailova et al. 1979). Calcium and magnesium ions were assumed to be bound to NTA in the homogenate. Additionally, data on tissue molarity $(0.5607 \text{ mmol } l^{-1} \text{ tissue water})$ and protein concentration (180 g 1^{-1} tissue water) of *Sipunculus nudus* (Pörtner 1982) were used in the calculations. Due to the high degree of dilution of tissue by the KF+NTA medium, variation in the latter has only minor influence on the calculated intracellular buffer values. All concentrations were corrected for dilution by medium and tissue water, as well as water added with HCl and derived from neutralisation of calcium carbonate. Tissue water content was measured for the foot of L. littorea as the difference between tissue wet weight and dry weight obtained after drying at 80 °C for 24 h. Average tissue water content was 76.43 \pm 0.027% (n = 60) and did not significantly change after 5 days of freshwater exposure. This value was applied in subsequent calculations for all three species.

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Additionally, tissue CaCO₃ buffer value (β_{CaCO_3}) was calculated as follows:

$$\beta_{\text{CaCO}_3} = \frac{3000 \times \text{V}_{\text{HCl}}}{\Delta \text{pH} \times \text{W}_{\text{tiss}}} (\text{mmol } \text{pH}^{-1} \text{ kg}^{-1})$$
(2)

where V_{HCl} is the volume (l) of 3 mol l⁻¹ HCl used for the back titration of the homogenate to physiological pH values, ΔpH is the difference between the initial homogenate pH and that reached after HCl titration, and W_{tiss} is the tissue wet weight (kg).

Chemicals

All chemicals were purchased from Sigma Chemical (St. Louis, Mo., USA) or Merck (Darmstadt, Germany).

Proton balance

The metabolic proton production ΔH^+_{met} was calculated according to Pörtner (1987a, b) using the data from our companion study (Sokolova et al. 2000). It was assumed that succinate accumulation is coupled to a production of 2 mol H^+ mol⁻¹, and ATP hydrolysis releases 0.43 mol H^+ mol⁻¹ at pH 6.9 (Pörtner 1987a). Influence of phosphagen depletion on proton balance was calculated considering the proton binding by the phosphate buffer at the final pH_i after 3 days or 5 days of freshwater exposure. The fraction of protonated phosphate (F) was calculated using Eq. 3 for pKa (8 °C) = 6.842 (Pörtner 1990):

$$F = \frac{1}{10^{\text{pH}-\text{pKa}} + 1} \tag{3}$$

Non-respiratory proton quantities $(\Delta H^+_{non-resp})$ were deduced from non-bicarbonate buffer lines on ΔH^+ versus pH_i plots. This procedure estimates a proton load to the tissue under the assumption that the observed pH_i changes are negligibly influenced by the bicarbonate system at low pH_i . Hence, any discrepancies between $\Delta H^+_{non-resp}$ and ΔH^+_{met} are assumed to represent the effects of other acid-base relevant processes like proton-equivalent ion transfer, calcium carbonate buffering, etc. (Pörtner 1987b).

Statistics

Statistical treatment was performed using standard Model I ANOVA procedures after testing the assumptions of normal distribution and homogeneity of variance of the data (Sokal and Rohlf 1995). We used Tukey's honestly significant difference test for unequal *N* as a method of post-hoc comparisons. For analyses of non-independent sequential measurements on the same samples obtained by ³¹P-NMR, distribution-free statistical procedures were used (Kruskal-Wallis test statistics and sign test; Sokal and Rohlf 1995). For constructing paired data for sign test, samples of each species were randomised within respective levels of the factor "exposure duration". All differences were considered significant if the probability level of type I error was less than 0.05. Results are expressed as mean \pm SE.

Results

In vivo NMR spectroscopy: Method validation and general results

Figure 1 shows typical *in vivo* ³¹P-NMR spectra of the three *Littorina* species under control conditions. All spectra exhibit the characteristic signals of high-energy phosphates: PLA and adenosine-5'-triphosphate (ATP), as well as that of Pi and a large signal arising from the



Fig. 1 In vivo ³¹P nuclear magnetic resonance (³¹P-NMR) spectra of *Littorina saxatilis*, *L. littorea* and *L. obtusata* sampled under control conditions. Each spectrum was accumulated over 10 min. Peaks are: *PLA* phospho-L-arginine, *ATP* adenosine-5'-triphosphate, *Pi* inorganic phosphate, *PD* phosphodiesthers

presence of phosphodiesters (PD). Although the phosphodiester signal seemed to be higher in the spectra of *L*. *obtusata*, no significant differences were found between the studied species (P > 0.05).

³¹P-NMR spectra (10-min acquisition period each) were collected repeatedly in *L. saxatilis* and *L. littorea* control samples demonstrating that signals remained unchanged and stable over a measurement period of almost 1 h and between repeated measurements during 24 h of control conditions.

It should be noted that the quality of spectra depends crucially on the conductivity and homogeneity inside the sample. The signal-to-noise (S/N) ratio varied between the spectra from control and freshwater-exposed animals as well as between the spectra obtained from different species. Considerably lower S/N ratios in spectra under control conditions were evidently due to the higher conductivity of seawater that decreased the performance of the NMR probe and furthermore, a higher inhomogeneity due to animal movements in control samples. This resulted in a broader line width of the signals in comparison to samples measured in freshwater. Impaired homogeneity in samples consisting of numerous small animals (as in L. saxatilis or L. obtusata) also led to a decrease in the S/N ratio as compared to samples containing few large L. littorea. Nevertheless, this does not influence estimation of relative peak areas and/or position of the peaks within one spectrum but would have a marked impact on calculations of absolute peak areas and their transformation into concentration units. Therefore, we only determined ratios of energyrelevant phosphorus compounds and intracellular pH_i values from the chemical shift of the inorganic phosphorus signal. We also calculated free Mg2+ concentrations for calculations of the Gibbs free-energy change of ATP hydrolysis (Sokolova et al. 2000).



Fig. 2 Typical stack plot of in vivo ³¹P-NMR spectra from *L. littorea* over all incubation periods. *a* control conditions; *b*, *c*, *d* 3 days, 5 days and 7 days of freshwater exposure, respectively. Note that the peak of inorganic phosphate (P_i) increased and shifted to low field indicating a more acidic intracellular pH value. PLA signal decreased during freshwater exposure and practically vanished at days 5 and 7

A typical stack plot of in vivo ³¹P-NMR spectra from *L. littorea* over all incubation periods is presented in Fig. 2. Spectrum a was accumulated under control conditions. After 3 days of hypoosmotic stress (spectrum b) the peak of P_i increased and shifted indicating a more acidic intracellular pH value. The PLA peak decreased, whereas the ATP signal remained almost constant throughout all exposure periods. After 5 days and 7 days of incubation the P_i signal rose even further (spectra c, d), and the PLA signal decreased. These changes are reflected in the dynamics of P_i/PLA and P_i/β ATP ratios which showed large and statistically significant increases in freshwater exposed animals as compared to controls in all studied species (Fig. 3).

Intracellular pH

Average levels of pH_i as estimated by in vivo ³¹P-NMR spectroscopy were similar in the studied species under control conditions (K-W test, P > 0.65) and varied between 7.45 and 7.56. During freshwater exposure a drastic drop in pH_i was observed (Fig. 4A). After 3 days of freshwater exposure pH_i had declined by 0.4– 0.5 units. Further pH_i changes were small and remained insignificant between conspecifics at later exposure periods (K–W, P > 0.10). In a between-species comparison, there is a trend for pH_i to decrease faster and reach lower values in the least resistant species *L. obtusata*. The rate of reduction and pH_i levels reached were intermediate in *L. saxatilis*. The slowest rate and highest



Fig. 3 Changes of NMR indices in White Sea *Littorina* spp. during freshwater exposure: A P_i/PLA ; B $P_i/\beta ATP$. Data are means of five samples measured for each exposure period, except for 7 days in *L. littorea* (n = 10). Increases in the NMR indices in freshwater-exposed animals are statistically significant compared to controls. Vertical bars represent standard errors

 pH_i values were observed in the most resistant *L. littorea* (Fig. 4A). However, a sign test showed that differences in pH_i were only statistically significant between the least resistant *L. obtusata* and the most resistant *L. littorea*.

Calculation of intracellular free Mg²⁺ concentration from in vivo ³¹P-NMR spectra resulted in average means of about 0.7 mmol l⁻¹ (range 0.55–0.84 mmol l⁻¹) in the three *Littorina* spp. studied (Fig. 4B). Variability was large in control samples of *L. obtusata* and *L. saxatilis* due to difficulties determining the exact position of the peak in rather broad β -ATP signals (data not shown). However, changes in Mg²⁺ concentrations during long-term incubation were minor and insignificant between species, and between different exposure periods.

Tissue buffer capacity

Non-bicarbonate, non-phosphate buffer values measured in homogenates of foot tissues varied between





Fig. 4 Intracellular pH (**A**) and free magnesium concentrations (**B**) calculated from ³¹P-NMR spectra in White Sea *Littorina* spp. during freshwater exposure. Data are means of five samples measured per exposure period, except for 7 days in *L. littorea* (n = 10). Vertical bars represent standard errors. Intracellular pH (pH_i) was significantly below control values for all exposure periods. Intracellular concentration of free Mg²⁺ did not change significantly during freshwater exposure in either species

28.5 mmol pH⁻¹ kg⁻¹and 42.8 mmol pH⁻¹ kg⁻¹ wet weight. However, two statistically significant outliers were detected: $\beta_{\text{NB, NPi}}$ of 67.8 mmol pH⁻¹ kg⁻¹ wet weight (a sample of *L. saxatilis* foot tissues after 5 days of exposure) and 73.6 mmol pH⁻¹ kg⁻¹ wet weight (a sample of *L. obtusata* under control conditions). No significant differences in $\beta_{\text{NB,NPi}}$ values were found between studied species nor between control and exposed animals (ANOVA, P > 0.05) regardless of whether outliers were rejected or not. Mean $\beta_{\text{NB,NPi}}$ (outliers excluded) are given in Fig. 5A.

Fig. 5 Foot muscle non-bicarbonate, non-phosphate $(\beta_{\text{NB,NPi}})$ (A), $P_i(\beta_{\text{Pi}})$ (B) and calcium carbonate (β_{CaCO3}) (C) buffer values of White Sea periwinkles during freshwater exposure. n = 6-10 (control) or 2–4 (5 days). Vertical bars represent standard errors or range (for n = 2)

Buffer values of P_i increased significantly during freshwater exposure in all three species studied from 0.26 mmol pH⁻¹ kg⁻¹ to 0.35 mmol pH⁻¹ kg⁻¹ wet weight under control conditions to 0.72– 0.81 mmol pH⁻¹ kg⁻¹ wet weight after 5 days of freshwater exposure (ANOVA, factorial effect of "freshwater exposure": P < 0.0001). There were also species-specific differences in the value of β_{Pi} in controls, the level of β_{Pi} being the highest in *L. littorea* (P < 0.001). No differences in β_{Pi} were found between studied species after 5 days of freshwater exposure (Fig. 5B).

Tissue calcium carbonate buffer values (β_{CaCO_3}) were considerably higher than $\beta_{NB,NPi}$ (Fig. 5C) and varied



Fig. 6 Relationship between metabolic proton accumulation and pH_i (A) and contribution of different metabolic processes to the metabolic proton load (B) during freshwater exposure in Littorina spp. A The relationship between pH_i and cumulative metabolic proton change (ΔH^+_{met}) reflects the effective buffer value of the tissue for metabolic protons and quantifies the extent to which protons are buffered in vivo. Filled circles: L. littorea; filled triangles: L. saxatilis; open squares: L. obtusata. Non-bicarbonate buffer lines $(\beta_{\rm H})$ determined in vitro for L. littorea (L.l.), L. saxatilis (L.s.) and L. obtusata (L.o.) (dotted lines) reflect physicochemical buffering starting from the respective control pH_i values. Note that during the initial phase of freshwater exposure (days 0-3) the drop in pH_i is close to or even somewhat larger than expected from non-bicarbonate tissue buffering alone, whereas later on the pH_i appears to be efficiently positively regulated probably due to calcium carbonate buffering. **B** Proton changes due to the production of succinate and cleavage of ATP and phosphagen are shown. All calculations are based on the mean changes of metabolite concentrations for the respective periods

between 148.2 and 277.3 mmol pH⁻¹ kg⁻¹ wet weight in control snails and between 94.7 mmol pH⁻¹ kg⁻¹ and 148.7 mmol pH⁻¹ kg⁻¹ wet weight in exposed snails. As shown by two-way ANOVA, the factor "freshwater exposure" had a significant influence on β_{CaCO_3} values of the foot tissues of *Littorina* spp. (P < 0.001). In general, calcium carbonate buffer values of *Littorina* spp. foot tissues decreased after 5 days of freshwater exposure as compared to the respective control levels (Fig. 5C). Between-species differences in β_{CaCO_3} were not significant neither in the control, nor after 5 days of freshwater exposure (P > 0.07).

Proton balance

 ΔH^+_{met} in the studied *Littorina* spp. varied between 11 µmol H⁺ g⁻¹ and 14 µmol H⁺ g⁻¹ wet weight after 3 days of freshwater exposure and up to ca. 30 µmol H⁺ g⁻¹ wet weight at the end of the experiment (Fig. 6A). Substantial increase in the metabolic proton load over time during freshwater exposure was predominantly a result of succinate accumulation (Fig. 6B). Contribution of phosphagen depletion to the total metabolic proton balance was relatively small, and that of ATP breakdown practically negligible, in particular during sustained freshwater exposure (Fig. 6B).

Between-species comparisons of the rates of metabolic proton production suggest that in the two more resistant species (*L. littorea* and *L. saxatilis*) the increase in metabolic proton load was lower during sustained anaerobiosis as compared to the early period of freshwater exposure (Fig. 6B). In contrast, the rate of metabolic proton production in the most susceptible *L. obtusata* increased considerably during the later stage of anaerobic exposure.

In general, intracellular pH_i was negatively related to the metabolic proton load. However, the rate of pH_i decrease changed considerably over the freshwater exposure period. During the first 3 days of freshwater exposure a fast and drastic drop in pH_i was observed with a rate close to or even slightly higher than expected from the exclusive buffering by the non-bicarbonate tissue buffers (Fig. 6A). Later on during freshwater exposure the rate of acidification of the intracellular milieu slowed down significantly despite continuing metabolic proton production. In fact, intracellular pH of *Littorina* spp. remained fairly constant over a wide range of cumulative metabolic acidification between 11 μ mol H⁺ g⁻¹ and 30 μ mol H⁺ g⁻¹ wet weight. At the highest proton loads (>20 μ mol H⁺ g⁻¹ wet weight) the drop in pH_i was less than expected from non-bicarbonate tissue buffering alone by approximately 0.2–0.4 pH units (Fig. 6A).

Discussion

In vivo NMR spectra

Changes of energy-relevant NMR ratios during freshwater exposure of White Sea *Littorina* spp. were in qualitative agreement with the biochemically determined changes of high-energy compounds in the tissues of the studied species (Sokolova et al. 2000). Ratios of P_i /PLA and $P_i /\beta ATP$ increased significantly in freshwater exposed animals as compared to controls, thus reflecting impaired energy status of the experimental snails.

Analysis of ³¹P-NMR spectra showed that P_i concentrations were rather high under control conditions in *Littorina* spp. Presence of high concentrations of P_i in the tissue is generally considered to indicate stressful conditions (e.g. van den Thillart and van Waarde 1996) The experimental design used in our study may have imposed a certain degree of stress due to crowding and restrained movements of the snails during acquisition of the NMR spectra. However, repeated measurements of ³¹P-NMR spectra showed no changes over time during control conditions in the magnet. This observation suggests that stress had only minimal effects on the animals. Additionally, rather high P_i concentrations were found in tissue extracts of control animals which experienced no crowding (Sokolova et al. 2000). Therefore, relatively high P_i levels in *Littorina* spp. under control conditions may reflect a higher content of NMR-visible phosphorus compounds including P_i in these species, which is a well-known feature in some invertebrates (van den Thillart and van Waarde 1996). Comparison of ³¹P-NMR spectra obtained from

Littorina spp. after different periods of freshwater exposure clearly demonstrates an increase in intracellular P_i concentrations in freshwater-exposed animals. This partially contrasts our previous invasive study (Sokolova et al. 2000) which showed relatively small (and only statistically significant in some species) increases in P_i contents in the foot tissues of the periwinkles under conditions of hypoosmotic stress. Reasons for this were discussed elsewhere (Sokolova et al. 2000). However, it should be noted that the increase in P_i concentrations observed in ³¹P-NMR spectra was higher than the decrease in PLA contents and suggests that there must have been other sources of P_i mobilisation during freshwater exposure in the studied Littorina spp. This change was not visible in tissue extracts (Sokolova et al. 2000) and indicates a change in free P_i concentration. Possibly the release of free P_i ions occurred from an insoluble complex of calcium phosphate at low intracellular pH (Weich et al. 1989), as phosphates are known to be components of the calcium carbonate granules found in connective tissue cells of molluscs (Mason et al. 1984). This suggestion is indirectly supported by the observed decrease of calcium carbonate buffer values in the foot in *Littorina* spp., which probably indicates mobilisation of material from tissue calcium carbonate granules during hypoosmotic stress. Nevertheless, further experiments must be performed to verify this hypothesis.

Interestingly, intracellular Mg^{2+} concentrations as determined by the non-invasive ³¹P-NMR technique, appear to be efficiently regulated in White Sea periwinkles. Free Mg^{2+} content is maintained at fairly constant

levels of ca. $0.6-0.8 \text{ mmol } \text{l}^{-1}$ even during prolonged anaerobiosis when many other crucial parameters of the intracellular milieu (including pH_i, energy status, etc.) undergo significant changes. Moreover, intracellular Mg²⁺ levels appear to be highly conserved between species of the genus *Littorina* studied. This finding is in accordance with the important role that this ion plays in many cellular processes (Simkiss and Mason 1983; Frederich et al. 2000).

Acid-base status and anaerobic metabolic rate

Even though NMR spectroscopy was able to provide only a qualitative picture of the changes in energy status, it was the only method allowing the determination of intracellular pH, which is crucial for investigating the acid-base status during prolonged anaerobiosis caused by freshwater exposure in *Littorina* spp. The homogenate technique of pH_i analysis (Pörtner et al. 1990) was not applicable in the calcium carbonate-rich tissues of White Sea *Littorina* spp.

Onset of anaerobiosis during freshwater exposure resulted in a substantial intracellular acidification in the three *Littorina* spp. studied. Intracellular pH dropped considerably at the beginning of freshwater exposure, with high rates of acidification during the first 3 days of exposure. Later on pH_i levelled off in *L. saxatilis* and *L. littorea* but not in *L. obtusata*, although differences in pH_i values between different exposure periods were not statistically significant in either species. This finding suggests that the periwinkles are able to regulate intracellular pH during anaerobiosis. In general, anaerobic pH regulation was most efficient in the most resistant species, *L. littorea*, as compared to *L. saxatilis* and especially *L. obtusata*.

Changes in intracellular pH values can theoretically be attributed to several processes, including (1) net proton production or consumption in metabolic reactions, (2) tissue proton buffering, (3) proton-equivalent transmembrane and transpithelial ion transfer, and (4) respiratory processes. The impact of respiratory proton production on acid-base balance of the snails could not be investigated since application of the homogenate technique was not possible (see above). However, the analysis of proton balance suggests a close correlation between pH_i changes observed and those expected from metabolic acidification and tissue buffering capacity (Fig. 6). Moreover, measurements of the oxygen consumption rates of *Littorina* spp. in freshwater show that there is no respiration detectable under these conditions (I. Sokolova, unpublished data). Therefore, the respiratory influence on acid-base balance of the snails can be considered small.

Proton buffering in the foot tissues of White Sea *Littorina* spp. was provided by multiple buffer systems including P_i, imidazole groups of proteins and probably oligopeptides (which are generally believed to contribute to non-bicarbonate, non-phosphate tissue buffering;

Eberlee and Storey 1984) and, most importantly, $CaCO_3$. Our data showed that the proton-buffering systems differ and shift in their contribution to tissue buffering during freshwater/anoxia exposure. This complex pattern reflects the potential for an efficient pH_i regulation during prolonged anaerobiosis in the periwinkles. Average values of $\beta_{NB,NPi}$ varied between 28.5 mmol pH^{-1} kg⁻¹ and 36.7 mmol pH^{-1} kg⁻¹ wet weight in the Littorina spp. studied and were in the low range reported for other invertebrate and lower vertebrate species, possibly for methodological reasons (Eberlee and Storey 1984; Heisler 1986; Wiseman and Ellington 1989; Zange et al. 1990; Ellington 1993; Pörtner 1990; Pörtner at al. 1996; van Dijk et al. 1997). Exposure to an extremely hypoosmotic environment for 5 days had no effect on $\beta_{\text{NB,NPi}}$ values of the periwinkles. In general, no correlation was found between freshwater resistance and $\beta_{\text{NB,NPi}}$ of foot tissues in the species studied, suggesting that differential resistance to freshwater and the discrepancy in the time-course of pH_i changes between White Sea Littorina spp. could not be accounted for by the variations in $\beta_{\text{NB,NPi}}$.

The buffer values for P_i were low in the three studied species (< 1 mmol pH⁻¹ kg⁻¹ wet weight) under control conditions as well as after freshwater exposure. The increase in β_{P_i} after 5 days of freshwater exposure was partially a result of a moderate elevation in P_i concentration (by ca. 27%) found in ultrafiltrated homogenate samples from freshwater exposed animals as compared to controls. This increase was evidently due to the cleavage of phosphagen and adenylate compounds during freshwater exposure (Sokolova et al. 2000). In addition, the twofold elevation of β_{P_i} in freshwater-exposed animals is related to tissue acidification which brings pH_i close to the apparent dissociation constant of H₂PO₃.

High initial pH values of tissue homogenates of Littorina spp. far exceeded in vivo pH_i determined by ³¹P-NMR and were in the range of 8.4–8.6. These high pH values can most likely be explained by the presence of crystalline CaCO₃ stored in connective tissue calcium cells of the foot (Mason et al. 1984; Brough and White 1990) and possibly, also by CaCO₃ inclusions in the mucus produced by the foot epithelium of these species (Davies and Hutchinson 1995). Buffer values of the CaCO₃ stores were five to six times higher than $\beta_{\text{NB,NPi}}$ values of the tissues. Again, no species-specific differences were found neither under control conditions, nor after 5 days of freshwater exposure. The β_{CaCO3} value decreased upon prolonged freshwater exposure in all species studied, thus indicating that part of this buffer was probably consumed during anaerobiosis. It is worth noting that an important role of shell CaCO₃ mobilisation in the neutralisation of acidic anaerobic end products in molluscs, has repeatedly been stressed since early in this century (Collip 1921; Dugal 1939; Akberali et al. 1977; Byrne and McMahon 1991). This role has also been suggested for CaCO₃ granules in the connective tissue calcium cells of gastropods (Mason et. al. 1984).

However, to the best of our knowledge, this is the first experimental indication that tissue CaCO₃ stores of the foot support proton buffering during sustained anoxia.

Analysis of the contribution of different metabolic processes to the proton balance of the tissue (Fig. 6B) suggests that succinate formation predominantly caused the acidification. An incomplete assay of metabolites can be excluded since succinate was the only end product among all tested metabolites (including alanopine, strombine, lactate, acetate and propionate) which was found to accumulate in the foot tissues of the Littorina spp. studied (Sokolova et al. 2000). The priority of endproduct accumulation to cause metabolic acidification of intracellular milieu was previously also reported for other marine organisms including the squids Illex illecebrosus and Lolliguncula brevis (Pörtner et al. 1991a, 1996), the lugworm Arenicola marina (Sommer and Pörtner 2000) and the peanut worm Sipunculus nudus (Pörtner 1987b, Hardewig et al. 1991).

During the first 3 days, metabolic proton production was similar to or even lower than non-respiratory proton changes deduced from the actual changes in pH_i and the non-bicarbonate tissue buffer values (Fig. 6A). The latter is an unexpected finding which suggests net proton uptake by the cells of *Littorina* spp. during the early stage of freshwater exposure, and may argue for a downregulation of intracellular pH during this period (Pörtner 1993). Considering a possible overestimation of the metabolic proton load due to the lack of data on ammonia concentrations, the discrepancy between ΔH^+_{met} and $\Delta H^+_{non-resp}$ may be even larger. For example, ammonia may bind between 5% and 20% of metabolic protons in temperature stressed lugworms Arenicola marina (Sommer and Pörtner 2000) and about 15% in anoxic peanut worms Sipunculus nudus (Pörtner 1987b). Later on during freshwater exposure, pH_i appears to be strongly positively regulated and is maintained at fairly constant levels despite the increase in the cumulative metabolic proton load. The difference between expected non-respiratory and observed metabolic proton loads changed to positive values, such that expected $\Delta H^+_{non-resp}$ was markedly higher than the observed acidification. This strongly indicates involvement of an additional mechanism in pH buffering, possibly ion transfer mechanisms and/or increased buffering by the $CaCO_3$ system. It is worth noting that the role of transmembrane and transepithelial ion exchange mechanism is expected to be relatively small due to the depressed overall metabolic rate during environmental anaerobiosis (Pörtner et al. 1991b; Sokolova et al. 2000). It is highly probable that the reduction in accumulated proton quantities is mostly caused by the CaCO₃ buffer system. This system is evidently not involved in pH_i regulation during the early stages of freshwater exposure and becomes mobilised only when pH_i drops to sufficiently low levels probably due to the increased solubility of calcium carbonate in an acidic medium. Comparison of the observed decrease in pH_i (e.g. $\Delta p H_{obs}$ of 0.1–0.15 between ΔH^+_{met} of 10 µmol g⁻¹ and 30 µmol g⁻¹ wet tissue weight) with those expected from non-bicarbonate tissue buffering alone (ca. 0.5–0.7 pH units) suggests that additional proton buffering of between 40 µmol H⁺ pH⁻¹ g⁻¹ and 50 µmol H⁺ pH⁻¹ g⁻¹ wet weight is involved. This value is roughly equivalent to what would be expected from the observed decrease in β_{CaCO_3} during freshwater exposure of the periwinkles. This quantitative agreement between the observed decrease in the CaCO₃ buffer value and the amount of the buffered metabolic protons evidences in favour of our hypothesis that a fraction of CaCO₃ stores is consumed for pH_i regulation during sustained anaerobiosis in White Sea *Littorina* spp.

In general, the observed dynamics of pH_i changes versus metabolic and non-respiratory proton production may suggest that pH_i regulation diverts from downregulation (or nearly no regulation) during early stages of environmental anaerobiosis to considerable positive regulation during sustained stress. Maintenance of lower set points of pH could help, on the one hand, to enhance the onset of metabolic arrest state during early anaerobiosis (Pörtner 1993), and on the other hand, to restrain fast acidification of the intracellular milieu caused by high metabolic proton production when anoxic conditions persist.

In conclusion, investigation of anaerobic metabolism during prolonged freshwater exposure in White Sea *Littorina* spp. reveals that various acid-base, metabolic and energy parameters are correlated and interconnected to provide the physiological basis for the remarkable freshwater resistance of periwinkles. During prolonged freshwater exposure, the change in pH_i follows a complex pattern and suggests a high capacity for pH_i regulation in the periwinkles, especially during sustained environmental anaerobiosis when high rates of succinate accumulation would otherwise lead to a deleterious extreme acidification of the intracellular space. The pH_iregulatory ability is probably in part provided by the multiple buffering systems present in the tissues of Littorina spp., especially by the CaCO buffer. Relative contribution of the multiple buffer systems to pH_i regulation evidently changes during freshwater exposure. At the initial stage of freshwater exposure the nonbicarbonate, non-phosphate tissue buffering system seems to be of a major importance for the metabolic proton buffering at pH_i between 7.5 and 7.0. During later stages of anaerobiosis and at lower pH_i, the CaCO₃ buffer is involved in proton buffering. A quantitative agreement between the decrease in the CaCO₃ buffer value and the quantity of buffered metabolic protons suggests that CaCO₃ tissue stores may serve as a major buffering system during prolonged anaerobiosis in Littorina spp. Species-specific differential resistance to extreme hypoosmotic stress can be attributed to the improved abilities for metabolic arrest (Sokolova et al. 2000), as well as to the increased ability to regulate pH_i and to defend relatively higher levels of energy potential of ATP in the more resistant species. However, interspecific variation in values of $\beta_{\text{NB,NPi}}$ and CaCO₃ tissue buffers does not apparently contribute to between-species differences in freshwater resistance.

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