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## Aerobic mitochondrial capacities in Antarctic and temperate eelpout (Zoarcidae) subjected to warm versus cold acclimation

Received: 3 September 2004 / Revised: 15 February 2005 / Accepted: 3 March 2005 / Published online: 15 April 2005  
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**Abstract** Capacities and effects of cold or warm acclimation were investigated in two zoarcid species from the North Sea (*Zoarces viviparus*) and the Antarctic (*Pachycara brachycephalum*) by investigating temperature dependent mitochondrial respiration and activities of citrate synthase (CS) and NADP<sup>+</sup>-dependent isocitrate dehydrogenase (IDH) in the liver. Antarctic eelpout were acclimated to 5°C and 0°C (controls) for at least 10 months, whereas boreal eelpout, *Z. viviparus* (North Sea) were acclimated to 5°C and to 10°C (controls). Liver sizes were found to be increased in both species in the cold, with a concomitant rise in liver mitochondrial protein content. As a result, total liver state III rates were elevated in both cold-versus and warm-exposed *P. brachycephalum* and *Z. viviparus*, with the highest rates in boreal eelpout acclimated to 5°C. CS and IDH activities in the total liver were similar in *Z. viviparus* acclimated to 5°C and 10°C, but decreased in those warm acclimated versus control *P. brachycephalum*. Enzyme capacities in the total liver were higher in eelpout from Antarctica than those from the North Sea. In conclusion, cold compensation of aerobic capacities in the liver seems to be linked to an increase in organ size with unchanged specific mitochondrial protein content. Despite its life in permanently cold climate, *P. brachycephalum* was able to reduce liver aerobic capacities in warm climate and thus, displayed a capacity for temperature acclimation.

### Introduction

The geographical distribution of ectothermic species is related to the ambient temperature regime, and tolerance to fluctuations of habitat temperature exists only within certain limits (for review see Pörtner 2001; Pörtner 2002a). Living in extreme Antarctic environment appears to be associated with reduced tolerance to higher temperatures. Low upper-lethal temperatures have been observed in the Antarctic brachiopod, *Liothyrella uva* between 3°C and 4.5°C (Peck 1989). Pörtner et al. (1999a) found a short-term upper lethal temperature of 4°C and a long-term upper limit of around 2°C in the bivalve *Limopsis marionensis*. An upper-lethal temperature of 6°C was found in three species of the Antarctic fish *Trematomus* by Somero and de Vries (1967). Antarctic eelpout, *Pachycara brachycephalum* and the Notothenioid, *Lepidonotothen nudifrons* survived temperatures of up to 9–10°C (van Dijk et al. 1999; Hardewig et al. 1999), and Mark et al. (2002) reported a lethal temperature of *P. brachycephalum* at around 13°C. Reduced heat tolerance in cold-stenothermal ectotherms is likely linked to cold induced functional adaptations, which compensate for the decelerating effect of low temperatures on metabolic processes.

Mitochondrial proliferation has been found in cold adapted Southern and Northern hemisphere species (Londrville and Sidell 1990; Johnston et al. 1998; Sommer and Pörtner 2002) as well as in cold acclimated eurythermal species (Campbell and Davies 1978; Egginton and Sidell 1989). Furthermore, enhanced enzyme activities are found in cold adapted as well as in cold acclimated animals (Crockett and Sidell 1990; Sokolova and Pörtner 2001; Kawall et al. 2002; Lannig et al. 2003). Besides enhanced mitochondrial densities, an increase in total organ size has also been reported to compensate for the effects of low temperatures (Kent et al. 1988 (liver and heart); Seddon and Prosser 1997 (liver); Lannig et al. 2003 (liver)).

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Despite compensatory increase in tissue aerobic capacities, Antarctic animals show uncompensated resting metabolic rates (Clarke 1991; van Dijk et al. 1999; for review see Pörtner et al. 2000; Peck 2002) and uncompensated specific maximal respiration rates of mitochondria (Johnston et al. 1994; Johnston et al. 1998; Pörtner et al. 1999b; Hardewig et al. 1999). Pörtner et al. (2000) suggested that mitochondrial and whole animal aerobic capacities remain uncompensated only in cold-stenothermal organisms, as opposed to cold adapted eurytherms. This may also be associated with the inability of cold-stenotherms to acclimate to warmer temperatures (e.g. Weinstein and Somero 1998).

The adjustment of aerobic scope by setting mitochondrial densities and capacities is seen as a crucial step in shifting thermal tolerance windows (cf. Pörtner 2002a). The present study was designed to investigate the ability to adjust aerobic metabolism to changing temperature in Antarctic compared to temperate eurythermal fish. We wanted to investigate whether a demersal Antarctic species like *P. brachycephalum* is able to undergo adjustments in aerobic metabolism to various temperatures despite living in a cold-stable environment. If so, the question arises whether the patterns of temperature compensation are similar for stenothermal and eurythermal ectotherms. Accordingly, the response to a 5°C temperature change was compared in closely related members of the zoarcid fish family. We acclimated cold adapted Antarctic eelpout, *P. brachycephalum*, from 0°C to 5°C, whereas temperate common North Sea eelpout, *Zoarces viviparus* were acclimated from 5°C to 10°C. We investigated mitochondrial respiration rates and the enzyme activities of citrate synthase (CS) and NADP<sup>+</sup>-dependent isocitrate dehydrogenase (IDH) in liver mitochondria at different temperatures. CS was chosen as an indicator of citric acid cycle activity and IDH as representing a key role in a mitochondrial substrate cycle, which may be relevant in the control of mitochondrial respiration, redox status and proton leakage (Sazanov and Jackson 1994; Pörtner et al. 1999b).

## Materials and methods

### Animals

Antarctic eelpout, *P. brachycephalum* were caught by baited traps close to King George Island (depth 397–455 m) during the RV Polarstern Antarctic expedition XV/3 1998. The fish were kept in aquaria onboard the RV Polarstern and later at the Alfred Wegener Institute (Bremerhaven) at ambient temperatures of 0 ± 0.5°C and a salinity of 34‰. North Sea eelpout, *Z. viviparus* were caught in the German Bight near Helgoland (in August 2001 and in March 2002) and kept in aquaria at the Alfred Wegener Institute at temperatures of 10 ± 1°C and a salinity of 34‰. Acclimation to 5°C started in spring 2002, with Antarctic and North Sea eelpouts

placed in the same aquarium at 5 ± 0.5°C and a salinity of 34‰. Other animals remained at control temperatures. All animals were kept under a 12:12-h light–dark cycle and fed live shrimp once per week. Experiments were carried out at the start of 2003, after an acclimation period at 5°C of at least 10 months.

### Preparation of mitochondria

The isolation of mitochondria was modified after Hardewig et al. (1999). After the fish were killed, the liver was removed immediately, chopped finely with scissors in an ice-cold glass dish, and extracted in 10–25 ml of homogenisation buffer, depending on liver wet mass, approximately 0.5 g per 10 ml (50 mM Hepes, 80 mM sucrose, 85 mM KCl, 5 mM EDTA, 5 mM EGTA, pH 7.1 (20°C) with freshly added 1% BSA and 1 µl ml<sup>-1</sup> aprotinin) with a Potter-Elvehjem homogeniser. After centrifugation (12 min at 400 g and 0°C) the pellet was re-homogenised in the same volume of homogenisation buffer and centrifuged again. The combined supernatants were centrifuged for 8 min at 10,000 g and 0°C. The mitochondrial pellet was re-suspended in 1–2 ml assay medium (50 mM Hepes, 85 mM KCl, 80 mM sucrose, 5 mM KH<sub>2</sub>PO<sub>4</sub>, pH 7.1 (20°C) with freshly added 1% BSA and 1 µl ml<sup>-1</sup> aprotinin). In some cases the livers of two animals were pooled to enhance the volume of mitochondrial suspension but counted as one in the statistical analyses and results.

### Mitochondrial respiration and enzyme assays

Mitochondrial oxygen consumption was measured at different temperatures (5, 8, 10, 15 and 20°C) using a Clark-type oxygen electrode in a thermostatted respiration chamber (Eschweiler, Kiel). A total of 100 µl of the mitochondrial suspension were combined with assay medium, without BSA and aprotinin, to a total volume of 1 ml, containing 0.5–1.2 mg mitochondrial protein per millilitre, 5 µM di-adenosine pentaphosphate (Ap<sub>5</sub>A, an inhibitor of myokinase) and 3.3 mM succinate as a substrate since respiration showed lower rates using pyruvate and malate. State III respiration was recorded after addition of 0.3 mM ADP and state VI respiration was determined after all ADP had been phosphorylated. Finally, the respiration induced by proton leakage (state IV<sub>ol</sub>) was recorded after adding 2.5 µg ml<sup>-1</sup> oligomycin, an inhibitor of mitochondrial F<sub>0</sub>F<sub>1</sub>-ATPase. P/O ratios were calculated by classical determination according to Chance and Williams (1956). Mitochondrial oxygen consumption was calculated as nanomole O min<sup>-1</sup> mg protein<sup>-1</sup> by adopting values of oxygen solubility in the assay medium from Johnston et al. (1994). Protein concentration was determined by the Biuret method after Gornall et al. (1949) using 5% deoxycholate to solubilize membrane proteins.

Since the high lipid content in liver disturbed the photometric measurements, 3 M TCA was used to precipitate and separate the protein by centrifugation for 15 min at 16,000 g and 15°C. The pellet was re-suspended with 1 ml diethylether and ethanol (4:1) and, after re-centrifugation, the pellet was re-suspended with 0.2 ml H<sub>2</sub>O and 1 ml Biuret reagent. Since the extraction of mitochondria was still incomplete, mitochondrial protein content per gram of liver was calculated from comparisons of CS activities measured in the mitochondrial suspension and in liver tissue extracts (100% CS).

Citrate synthase activity was quantified in the mitochondrial suspension, specified in units per milligram of mitochondrial protein, and in liver tissue (units per g wet wt tissue) using a modified procedure after Sidell et al. (1987). Mitochondrial suspension was added to assay medium (2:1) and, after 15 min of ultrasonic treatment at 0°C (output control 8, duty cycle 50%, Branson Sonifier 450, Heinemann Ultraschalltechnik, Germany), was centrifuged for 5 min at 7,400 g and 0°C.

For the determination of CS activity in liver tissue (100% CS) approximately 0.02 g of tissue (taken after chopping the liver during the preparation of isolated mitochondria isolation) were added to 1 ml of assay medium without BSA and aprotinin, subjected to ultrasonic treatment (15 min, output control 8, duty cycle 50% at 0°C) and centrifuged for 5 min at 7,400 g and 0°C. The supernatants were kept on ice and enzyme activities remained stable for at least 24 h. CS activity was determined in 100 mM Tris-HCl, pH 8 (20°C), 5 mM DTNB, 20 mM Acetyl-CoA, 20 mM oxaloacetate. The increase in absorbance at 412 nm was monitored in a thermostatted spectrophotometer at 5, 10, 15 and 20°C, respectively. Enzyme activity was calculated using an extinction coefficient ( $\epsilon_{412}$ ) for DTNB of 13.61 l mmol<sup>-1</sup> cm<sup>-1</sup>.

For the determination of NADP<sup>+</sup>-dependent IDH activity, mitochondrial suspension was added to stabilisation buffer (2:1; containing 50 mM Hepes, 85 mM KCl, 80 mM sucrose, 5 mM KH<sub>2</sub>PO<sub>4</sub>, pH 7.1 (20°C), 2 mM DTT, 20 mM MgCl<sub>2</sub>, 20 mM isocitrate and 5 mM EDTA), treated for 5 min at 0°C in an ultrasonic bath (output control 2, duty cycle 50%) and centrifuged for 9 min at 7,400 g and 0°C. The enzyme remained stable for more than 24 h after this treatment. IDH activity was measured after Alp et al. (1976), by monitoring the appearance of NADPH at 339 nm in a temperature controlled spectrophotometer at 5, 10, 15 and 20°C, respectively. The assay contained 70 mM Tris-HCl pH 7.1 (20°C), 8 mM MgSO<sub>4</sub>, 2.5 mM NADP<sup>+</sup>, 1 mM MnCl<sub>2</sub>, 3 mM D/L-isocitrate and 20 mM citrate. Enzyme activity was calculated using an extinction coefficient ( $\epsilon_{339}$ ) for NAD(P)H of 6.3 l mmol<sup>-1</sup> cm<sup>-1</sup>.

#### Allometric scaling

Since liver size and lipid content influence liver aerobic capacity available to the animal, we calculated total

mitochondrial protein content, mitochondrial respiration rates and enzyme activities for total liver, normalized to a 40 g eelpout (see Pelletier et al. 1993; Lannig et al. 2003). Total mitochondrial protein content and total CS activities were calculated with the following formula:

$$N = T \cdot \left( \frac{40}{\text{body mass}} \right)^m,$$

where  $N$  indicates normalized mitochondrial protein or normalized CS activity and  $T$  indicates total mitochondrial protein or total CS activity. The body mass exponent 'm' was determined from the slope of the linear regression in a double logarithmic plot of mitochondrial protein or CS activity in total liver versus body mass in gram.

Mitochondrial respiration rates and IDH activity in total liver were calculated from the normalized protein content. No differences were found in total CS activity between direct normalization, described above, and calculation from normalized protein content.

Calculation of the hepatosomatic index as:

$$I_H = \frac{\text{liver mass (g)}}{\text{body mass (g)}} \times 100,$$

was carried out with normalized liver mass for a 40 g eelpout by the following formula:

$$\text{Normalized liver mass} = \text{exact liver} \times \left( \frac{40}{\text{body mass}} \right)^m,$$

where the body mass exponent  $m$  was determined from the slope of the linear regression in a double logarithmic plot of liver mass versus body mass in grams. Data of animals and mass exponents for the different allometric scalings are summarized in Table 1.

#### Statistical analyses

All values are given as means ( $\pm$  standard deviation). Statistical significance was tested at the  $P=0.05$  level using analysis of variance (ANOVA, SigmaStat) and post hoc procedures (Tukey test). Before statistical testing, log transformations were used whenever necessary to meet statistical assumptions of the homogeneity of variances. However, non-transformed data are given in text and figures.

## Results

### Hepatosomatic index and mitochondrial liver protein content

Normalized hepatosomatic indices ( $I_H$ , Fig. 1) were significantly higher in cold-exposed than in warm-exposed North Sea and Antarctic eelpout.  $I_H$  was lowest in *Z. viviparus* acclimated to 10°C. However, no

**Table 1** Body and liver masses, condition factors (k-factor), gonadosomatic indices ( $I_G$ ) and mass exponents ( $m$ ) of regression equations (with  $P$  values) for the normalization to a 40-g specimen of the two eelpout species, *Z. viviparus* (North Sea) and *P. brachycephalum* (Antarctica) from the different acclimation groups

Species	<i>Zoarces viviparus</i> North Sea		<i>Pachycara brachycephalum</i> Antarctica	
Acclimation temperature (°C)	10°C (control)	5°C	5°C	0°C (control)
$n$	6–13	4–5	7–19	20–21
Body mass (g)	42.88 ± 22.29	63.34 ± 17.34	39.19 ± 8.11	35.25 ± 11.62
Liver mass (g)	0.85 ± 0.70	2.43 ± 1.01	1.18 ± 0.57	1.16 ± 0.58
Condition factor	0.37 ± 0.08	0.41 ± 0.02	0.36 ± 0.03	0.40 ± 0.04
Gonadosomatic index	0.54 ± 0.15	0.39 ± 0.19	0.42 ± 0.12	0.33 ± 0.24
$m$ for normalization of hepatosomatic index	1.36 (13); $P < 0.0001$	1.01 (5); $P = 0.01$	2.29 (19); $P < 0.0001$	1.37 (19); $P < 0.0001$
$m$ for normalization of mitochondrial protein content	1.91 (6); $P = 0.043$	1.52 (5); $P = 0.021$	1.38 (8); $P = 0.019$	0.90 (9); $P = 0.035$
$m$ for normalization of CS activity	1.11 (5); $P < 0.0001$	1.39 (5); $P = 0.0008$	1.20 (7); $P < 0.0001$	0.98 (8); $P < 0.0001$

Numbers of experimental animals are shown in parentheses. Data are means ± SD

differences were found between *Z. viviparus* acclimated to 5°C and *P. brachycephalum* acclimated to control conditions (0°C). *P. brachycephalum* acclimated to 5°C showed a significantly lower  $I_H$  than *Z. viviparus* at 5°C.

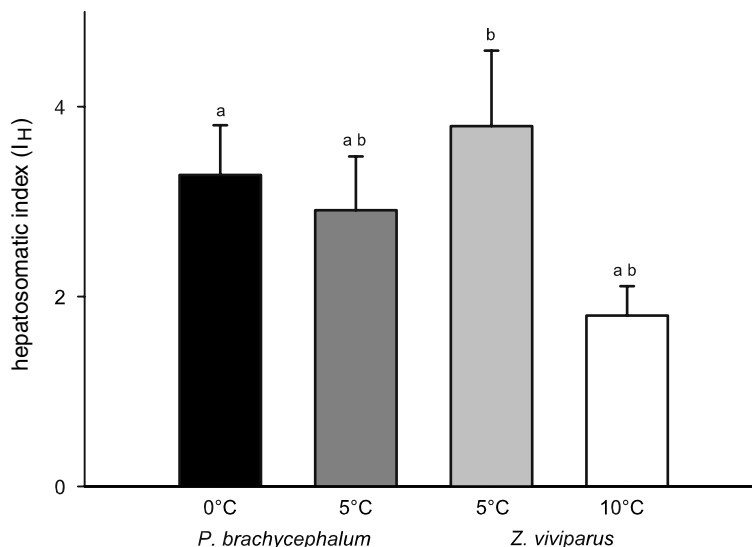
Specific mitochondrial protein contents (mg g liver<sup>-1</sup>) were more or less similar in control and acclimation groups of one species as well as the two species. Values were 35.41 ± 5.31 (North Sea, 10°C), 43.15 ± 10.97 (North Sea, 5°C), 30.16 ± 10.37 (Antarctica, 5°C) and 37.10 ± 7.48 (Antarctica, 0°C) (means ± standard deviation). Normalized protein content in total liver showed acclimation effects in both species, with higher values in cold compared to warm acclimated boreal and polar eelpout (Fig. 2). Species comparison revealed significantly higher mitochondrial liver protein in *P. brachycephalum* (0°C and 5°C) compared to control (10°C) *Z. viviparus*, but similar total protein contents compared to *Z. viviparus* acclimated to 5°C.

### Mitochondrial respiration

Mitochondrial coupling of all groups was significantly higher when assayed at 5°C than at 20°C. At 5°C respiratory control ratios (RCR<sub>ol</sub>, ratio of state III and state IV under oligomycin) were 4.85 ± 1.29 (North Sea, 10°C), 8.37 ± 1.13 (North Sea, 5°C), 4.39 ± 0.93 (Antarctica, 5°C) and 4.87 ± 1.08 (Antarctica, 0°C) (Fig. 3). The traditional RCR (ratio of state III and state IV without oligomycin, data not shown) was found to be only slightly lower than RCR<sub>ol</sub>. Comparison of RCR<sub>ol</sub> between groups of *Z. viviparus* revealed significantly higher values in animals acclimated to 5°C than in controls (10°C), whereas no acclimation effect was found in RCR<sub>ol</sub> of Antarctic eelpout. Coupling was significantly higher at all temperatures in North Sea eelpout acclimated to 5°C than in control and in 5°C acclimated Antarctic eelpout.

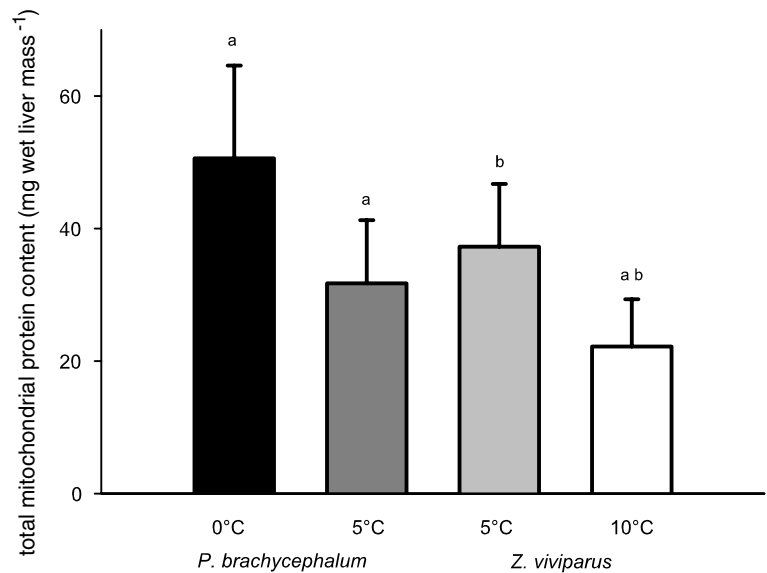
Determination of classical P/O ratios yielded numbers between 1.5 and 2.5 for all acclimation groups assayed in

**Fig. 1** Hepatosomatic indices, normalized to a 40 g animal, in Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) ( $n = 19$ ) and acclimated to 5°C ( $n = 19$ ), in *Z. viviparus* (North Sea) acclimated to 5°C ( $n = 5$ ) and 10°C (control conditions,  $n = 13$ ). Data marked with the same letters are significantly different





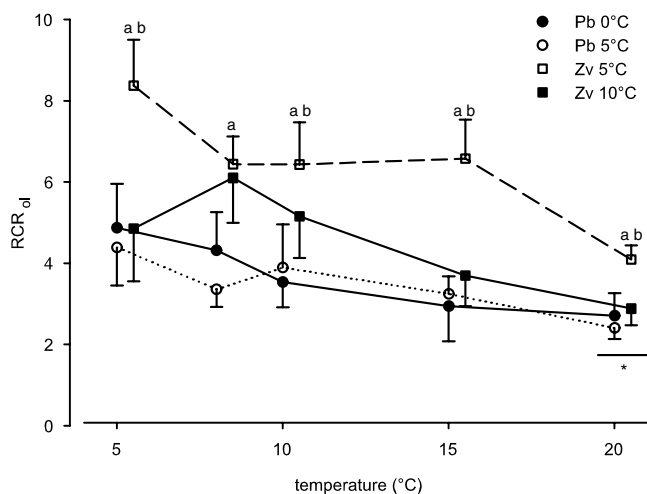
**Fig. 2** Total mitochondrial protein content (normalized to a 40 g animal), in liver of Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) ( $n=8$ ) and acclimated to 5°C ( $n=8$ ), in *Z. viviparus* (North Sea) acclimated to 5°C ( $n=4$ ) and 10°C (control conditions,  $n=6$ ). Data marked with the same letters are significantly different



the temperature range of 5°C and 10°C. Above 10°C P/O values of mitochondria from warm acclimated *P. brachycephalum* and *Z. viviparus* (at 20°C only) decreased significantly, resulting in significantly lower values compared to the more or less unchanged and temperature independent P/O ratios of cold acclimated Antarctic and North Sea eelpout (data not shown).

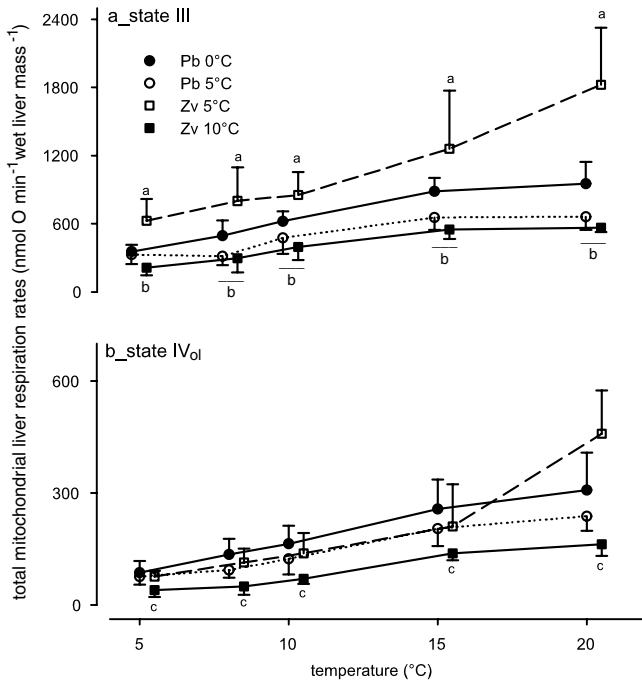
Mitochondrial respiration rates (nmol O min<sup>-1</sup> mg protein<sup>-1</sup>; state III, state IV and state IV<sub>ol</sub>) increased significantly with rising temperatures in all groups (data

not shown), whereas acclimation had no effect on specific mitochondrial respiration within both species. When measured at 5°C specific state III respiration rates of North Sea eelpout were 11.5±3.7 (10°C) and 14.2±3.5 (5°C) and those of Antarctic eelpout were 6.8±2.1 (0°C) and 9.7±1.4 (5°C). Respiration rates were significantly lower in polar eelpout acclimated to 0°C than in boreal eelpout acclimated to 5°C and 10°C, for state III at all temperatures and for state IV<sub>ol</sub> at 20°C only. Warm acclimated *P. brachycephalum* showed significant differences only in state III respiration, which was lower than in *Z. viviparus* controls (10°C) at assay temperatures between 8°C and 15°C. At investigated temperatures, no significant discontinuity in Arrhenius plots was found for all groups (data not shown).  $Q_{10}$  values and Arrhenius activation energies ( $E_A$ ) were similar in all groups: *P. brachycephalum*: state III:  $Q_{10}=2.0\pm0.4$ ,  $E_A=49.2\pm11.9$  kJ mol<sup>-1</sup>; state IV:  $Q_{10}=2.9\pm0.7$ ,  $E_A=67.1\pm12.9$  kJ mol<sup>-1</sup> and state IV<sub>ol</sub>:  $Q_{10}=3.2\pm0.8$ ,  $E_A=74.3\pm14.7$  kJ mol<sup>-1</sup> and *Z. viviparus*: state III:  $Q_{10}=2.2\pm0.4$ ,  $E_A=53.4\pm10.5$  kJ mol<sup>-1</sup>; state IV:  $Q_{10}=3.2\pm0.8$ ,  $E_A=76.9\pm17.5$  kJ mol<sup>-1</sup> and state IV<sub>ol</sub>:  $Q_{10}=3.2\pm0.5$ ,  $E_A=79.8\pm8.6$  kJ mol<sup>-1</sup> (means±standard deviation for control and acclimation groups of each species).



**Fig. 3** Temperature dependence of respiratory coupling ratios under oligomycin ( $RCR_{ol}$ ) of mitochondria isolated from liver of Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) (filled circles  $n=4-8$ ) and acclimated to 5°C (open circles  $n=4-8$ ) and in *Z. viviparus* (North Sea) acclimated to 5°C (open squares  $n=4-5$ ) and 10°C (control conditions, filled squares  $n=4-6$ ). For better viewing squares were shifted to the right. \* indicates values significantly different from those at 5°C. **a** indicates  $RCR_{ol}$  of *Z. viviparus* at 5°C significantly different to data from *P. brachycephalum* at 0°C and 5°C, when compared at the same temperature. **b** indicates  $RCR_{ol}$  of *Z. viviparus* at 5°C significantly different to data from *Z. viviparus* at 10°C, when compared at the same temperature

Mitochondrial state III respiration normalized for total liver of a 40 g animal showed significant acclimation effects in both species (Fig. 4a). Respiration was higher in total liver of cold-exposed versus warm-exposed *Z. viviparus* and *P. brachycephalum*. Liver respiration capacities (state III) in 5°C acclimated boreal eelpout were significantly higher than in control (0°C) and 5°C acclimated polar eelpout at all temperatures. However, total liver mitochondria of control *P. brachycephalum* (0°C) respired at higher rates than total liver mitochondria of control *Z. viviparus* (10°C). State IV respiration under oligomycin, which is assumed to quantify oxygen consumption due to proton leakage through the inner mitochondrial membrane (IV<sub>ol</sub>,



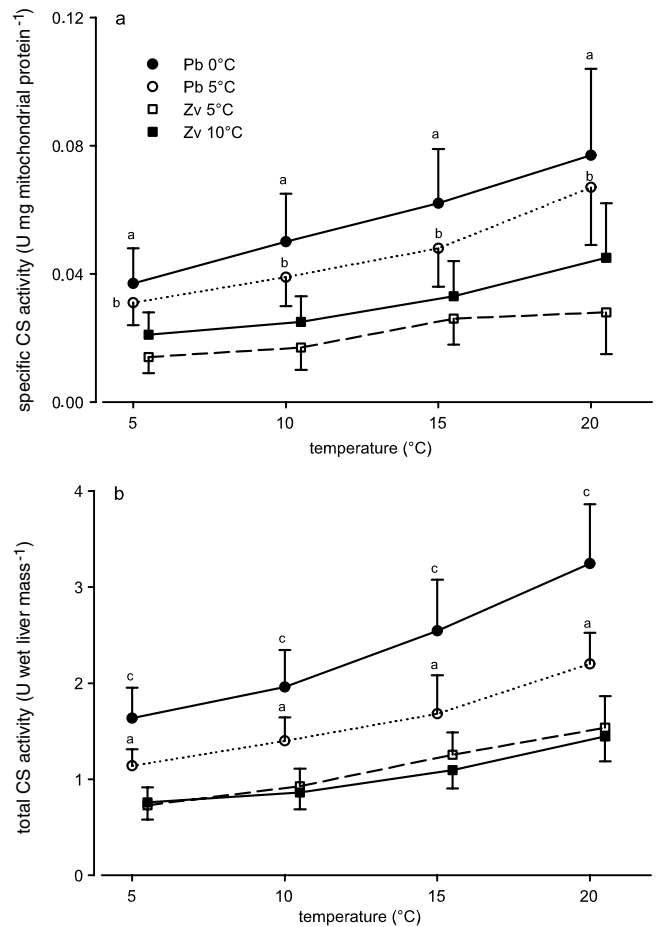
**Fig. 4** Temperature dependence of total mitochondrial respiration rates, state III (a) and state IV<sub>ol</sub> (b, state IV under oligomycin) in liver (normalized to a 4 g animal) of Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) (filled circles  $n=4-6$ ) and acclimated to 5°C (open circles  $n=4-8$ ) and in *Z. viviparus* (North Sea) acclimated to 5°C (open squares  $n=4-5$ ) and 10°C (control conditions, filled squares  $n=3-5$ ). For better viewing squares were shifted to the right. **a** indicates data from *Z. viviparus* at 5°C significantly different to data from the other acclimation groups, when compared at the same temperature. **b** indicates data from *P. brachycephalum* at 5°C and from *Z. viviparus* at 10°C significantly different to data from *P. brachycephalum* at 0°C, when compared at the same temperature. **c** indicates data from *Z. viviparus* 10°C significantly different to data from the other acclimation groups, when compared at the same temperature

Fig. 4b), showed higher rates after cold acclimation in the total liver of *Z. viviparus* only, whereas state IV<sub>ol</sub> capacity remained similar in total liver of warm acclimated (5°C) and control (0°C) *P. brachycephalum*. State IV<sub>ol</sub> rates were again higher in the liver of 0°C and 5°C exposed Antarctic eelpout than of North Sea eelpout at 10°C.

### Enzyme activities

Citrate synthase activities in the mitochondrial suspensions (U mg mitochondrial protein<sup>-1</sup>) were similar in acclimated and control groups of both species (Fig. 5a). Control groups (0°C) of Antarctic eelpout displayed significantly higher specific CS activities than both control and acclimation groups of North Sea eelpout. However, this difference decreased with warm acclimation and specific CS activity of liver mitochondria from 5°C acclimated *P. brachycephalum* was significantly higher only compared to those from *Z. viviparus* accli-

mated to 5°C. No discontinuities of Arrhenius plots were found in the investigated temperature range (data not shown) and  $Q_{10}$  values and Arrhenius activation energies ( $E_A$ ) were similar in all groups, with  $Q_{10}=1.6\pm 0.1$  and  $E_A=32.5\pm 5.3$  kJ mol<sup>-1</sup> for *P. brachycephalum* and with  $Q_{10}=1.6\pm 0.1$  and  $E_A=33.5\pm 4.2$  kJ mol<sup>-1</sup> for *Z. viviparus* (means  $\pm$  standard deviation for control and acclimation groups of each species). CS activities in the total liver of a 40 g animal again showed again no differences between groups of *Z. viviparus*, whereas higher total CS activities were found in control (0°C) compared to 5°C acclimated *P. brachycephalum* (Fig. 5b). The normalized liver of Antarctic eelpout (0°C and 5°C) displayed significantly



**Fig. 5** Temperature dependence of citrate synthase activity (CS), specific (a) and total (b normalized to a 40 g animal), in liver of Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) (filled circles  $n=6-8$ ) and acclimated to 5°C (open circles  $n=7$ ) and in *Z. viviparus* (North Sea) acclimated to 5°C (open squares  $n=4-5$ ) and 10°C (control conditions, filled squares  $n=5$ ). For better viewing squares were shifted to the right. **a** indicates data significantly different to data from *Z. viviparus* at 5°C and 10°C, when compared at the same temperature. **b** indicates data from *P. brachycephalum* at 5°C significantly different to data from *Z. viviparus* at 5°C, when compared at the same temperature. **c** indicates data from *P. brachycephalum* at 0°C significantly different to data from the other acclimation groups, when compared at the same temperature

higher total CS capacity compared to North Sea eelpout (5°C and 10°C).

For NADP<sup>+</sup>-dependent IDH activities in the mitochondrial suspension (U mg mitochondrial protein<sup>-1</sup>), activities were higher in control (0°C) Antarctic eelpout than in control (10°C) and cold acclimated (5°C) North Sea eelpout at all temperatures (Fig. 6a). Evaluation of IDH activity in the total liver content of a 40 g animal showed no differences between cold acclimated and control *Z. viviparus*, but again higher IDH activities in control (0°C) compared to 5°C acclimated Antarctic eelpout (Fig. 6b). Total IDH capacity was significantly higher in the liver of control 0°C Antarctic than of 5°C and 10°C exposed North Sea eelpout, whereas this difference fell at warmer temperatures and IDH capacity in

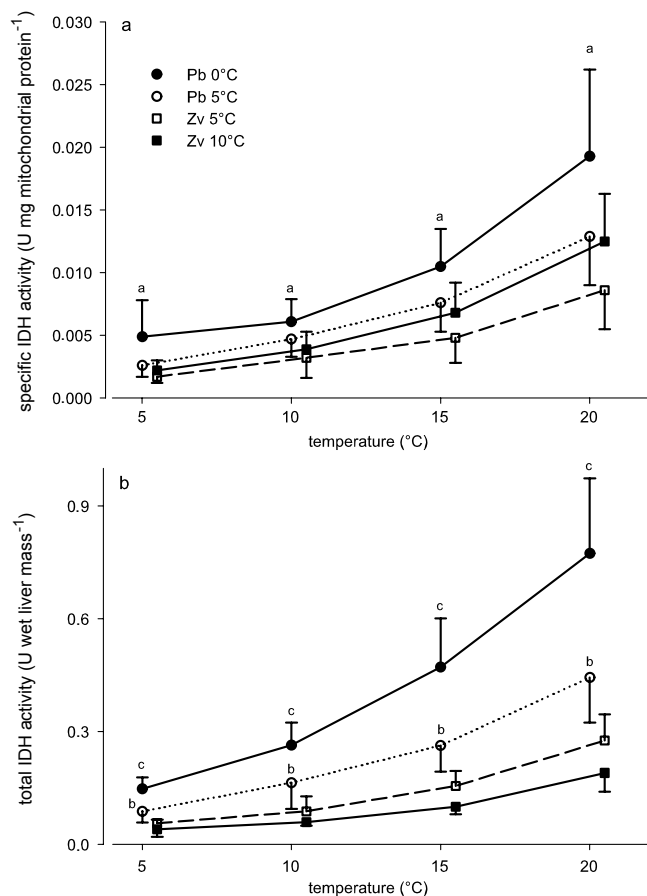
liver of 5°C acclimated *P. brachycephalum* was significantly higher than in control (10°C) *Z. viviparus* only. As for CS activities, no discontinuities were found in Arrhenius plots of IDH activities in the investigated temperature range (5°C – 20°C, data not shown) and  $Q_{10}$  values and Arrhenius activation energies ( $E_A$ ) did not differ between groups,  $Q_{10}=2.9\pm 0.3$  and  $E_A=72.4\pm 6.3$  kJ mol<sup>-1</sup> for *P. brachycephalum* and with  $Q_{10}=3.1\pm 0.3$  and  $E_A=76.5\pm 5.4$  kJ mol<sup>-1</sup> for *Z. viviparus* (means  $\pm$  standard deviation for control and acclimation groups of each species).

## Discussion

### Respiratory properties of liver mitochondria

Liver mitochondria of eelpout, *Z. viviparus* (North Sea, at 10°C and 5°C) and *P. brachycephalum* (Antarctica, at 0°C and 5°C) showed a clear negative correlation of respiratory coupling ratios with rising temperature (Fig. 3). The decrease of coupling ratios with temperature was elicited by different thermal sensitivities of oxidative phosphorylation capacity (state III) and proton leakage capacity (state IV<sub>oi</sub>) showing lower Arrhenius activation energies for state III than for state IV<sub>oi</sub>. Progressive uncoupling with rising temperature was also reported by Hardewig et al. (1999) and by Pörtner et al. (1999b) for Antarctic species, whereas studies carried out with temperate zone eurytherms showed coupling ratios to be stable and independent of temperature (Blier et al. 2001, Fischer 2002) or fairly insensitive to temperature (van den Thillart and Modderkolk 1978, Moyes et al. 1988, Blier and Guderley 1993).

These findings, along with the present results of no clear acclimation effect on RCR (higher in *Z. viviparus* after cold acclimation but unchanged in *P. brachycephalum* after warm acclimation) suggest that there is no general pattern of changing mitochondrial coupling ratios with temperature and that patterns are species specific. Despite decreasing coupling ratios with increasing temperature in all groups, only mitochondria of the warm acclimated North Sea and Antarctic eelpout showed the expected concomitant decrease in phosphorylation efficiency (ADP/O ratio) at high temperatures. Pörtner et al. (1999b) found no change in P/O ratio with temperature by classical determination (from changes in respiration slopes) but significantly lower P/O ratios at high temperatures by biochemical P/O ratio determinations. These differences associated with different methods might indicate that the classical determination is not sensitive enough for small changes in P/O ratios. Hardewig et al. (1999) discussed that unchanged P/O ratios, despite progressive uncoupling, might become explainable in classical analyses of P/O ratios when maximal phosphorylation rates are determined at low proton motive force and low proton leak, and compared with the subsequent transition to maximum proton leakage at maximum proton motive force.



**Fig. 6** Temperature dependence of NADP<sup>+</sup>-dependent isocitrate-dehydrogenase activity (IDH), specific (a) and total (b, normalized to a 40 g animal), in liver of Antarctic and North Sea Zoarcidae, in *P. brachycephalum* (Antarctica) under control conditions (0°C) (filled circles  $n=6-8$ ) and acclimated to 5°C (open circles  $n=5-6$ ) and in *Z. viviparus* (North Sea) acclimated to 5°C (open squares  $n=3-5$ ) and 10°C (control conditions, filled squares  $n=4-6$ ). For better viewing squares were shifted to the right. **a** indicates data from *P. brachycephalum* at 0°C significantly different to data from *Z. viviparus* at 5°C and 10°C, when compared at the same temperature. **b** indicates data from *P. brachycephalum* at 5°C significantly different to data from *Z. viviparus* at 10°C, when compared at the same temperature. **c** indicates data from *P. brachycephalum* at 0°C significantly different to data from the other acclimation groups, when compared at the same temperature

Acclimation had no effect on specific maximal phosphorylation capacities (state III in  $\text{nmol O min}^{-1} \text{mg mitochondrial protein}^{-1}$ ) of liver mitochondria of both eelpout species but rates were lower in Antarctic than in North Sea eelpout. Johnston et al. (1994, 1998) and Guderley (1998) concluded that cold compensation of specific state III respiration does not occur in muscle of various fish from a latitudinal cline (largely Southern hemisphere species). In eurythermal fish from Northern latitudes, however, cold acclimation and adaptation are reflected in increased specific aerobic capacities of mitochondria (Fischer 2002), and it was suggested as an explanation that an increase in the specific phosphorylation capacity typically occurs in cold adapted eurytherms living at variable temperatures but may be negligible in stenothermal animals living at permanently low temperatures (cf. Pörtner et al. 2000). In that respect cold acclimated *Z. viviparus* are more eurythermal than *P. brachycephalum*.

Mitochondrial protein per g liver did not differ significantly between acclimation groups, and was only slightly higher in the cold. Mitochondrial proliferation as shown in muscle (Egginton and Sidell 1989; Johnston et al. 1998) did not thus occur during temperature compensation in liver. As found in previous studies, metabolic cold compensation in liver goes along with an increase in organ size (Kent et al. 1988; Seddon and Prosser 1997; Lannig et al. 2003) and one might expect a decrease in those warm acclimated. This is confirmed by the present paper (Fig. 1). By decreasing mitochondrial protein content with decreasing total liver size, warm acclimated *P. brachycephalum* showed similar patterns of temperature compensation as known for eurythermal fish species, resulting in a similar total hepatic mitochondrial content as found in *Z. viviparus* at the same temperature. This finding would suggest that *P. brachycephalum* has conserved some features of their eurythermal ancestors (see "Effect of warm acclimation on Antarctic eelpout"). Due to different mitochondrial protein contents in total liver, the picture of mitochondrial respiration rates changed after calculation for total liver. Mitochondrial state III respiration in total liver clearly showed cold or warm compensation (Fig. 4a). Our findings are in line with data by van Dijk et al. (1999) who found higher metabolic rates ( $\text{nmol O min}^{-1} 100 \text{ g eelpout}^{-1}$ ) in both cold adapted *P. brachycephalum* ( $0^\circ\text{C}$ ) and cold acclimated *Z. viviparus* ( $3^\circ\text{C}$ ) than in warm acclimated *Z. viviparus* ( $12^\circ\text{C}$ ). However, van Dijk et al. found similar standard metabolic rates in cold acclimated North Sea and cold adapted Antarctic eelpout, therefore, the overcompensation in state III respiration capacities of liver mitochondria of cold acclimated *Z. viviparus* found in the present work does not necessarily imply a concomitant rise in whole animal metabolic rate.

Since mitochondrial proton leakage rates correlate with the standard metabolic rate of an animal (Brookes et al. 1998) and comprise a constant fraction of 20%–50% of standard metabolism in endotherms and most

likely ectotherms (Brand 1990; Brand et al. 1994; Brookes et al. 1998), our results indicate that oxygen consumption due to proton leakage through the inner mitochondrial membrane (state  $\text{IV}_{\text{ol}}$ ) mirrors elevated metabolic rates in the cold. State  $\text{IV}_{\text{ol}}$  respiration in total liver of polar versus cold acclimated boreal eelpouts revealed no difference and thus, a similar pattern as found for the standard metabolism of eelpouts by van Dijk et al. (1999). Investigation of whole animal respiration rates also revealed similar values in *P. brachycephalum* acclimated to  $0^\circ\text{C}$  and  $5^\circ\text{C}$  (data not shown) despite different state III respiration capacities in the liver.

According to our earlier hypothesis, proton leakage is down-regulated in parallel with standard metabolism in cold adapted stenothermal Antarctic ectotherms (Pörtner et al. 2000), a conclusion supported by proton leakage rates found in bivalves and notothenioids (Pörtner et al. 1999b; Hardewig et al. 1999). However, in the case of the zoarcids, no differences were found in mitochondrial liver state  $\text{IV}_{\text{ol}}$  capacity and standard metabolic rate between cold acclimated boreal and polar ( $0^\circ\text{C}$  and  $5^\circ\text{C}$  acclimation) zoarcids. It is conceivable that the Antarctic zoarcid does not follow the trend found in invertebrates and notothenioids and shows no down-regulation of standard metabolism relative to its temperate zone congeners. This is also mirrored in the lack of down-regulation of the mitochondrial proton leak leading again to the conclusion that *P. brachycephalum* has kept some features of their eurythermal ancestor.

#### Effects of temperature on mitochondrial enzymes

In ectothermic animals long-term temperature shifts may cause compensatory changes in glycolytic and mitochondrial enzyme levels and capacities (Crockett and Sidell 1990; Torres and Somero 1988; Lannig et al. 2003). These findings are consistent with our present data (Figs. 5, 6). Warm acclimation led only to a slight reduction in specific CS and IDH activities ( $\text{U mg mitochondrial protein}^{-1}$ ) in *P. brachycephalum* but in parallel with reduced mitochondrial protein content in liver, total enzyme capacities were significantly decreased by a factor of 1.43 for CS and 1.69 for IDH (measured at  $5^\circ\text{C}$ ), when compared to *P. brachycephalum* at  $0^\circ\text{C}$ . Cold compensated CS activities after calculation for total liver were also found by Kent et al. (1988) and Lannig et al. (2003), though it is somewhat unexpected that the slightly lower specific enzyme (CS and IDH) activities in liver mitochondria of cold compared to warm acclimated North Sea eelpout remained uncompensated for despite the higher mitochondrial protein content in *Z. viviparus* acclimated to  $5^\circ\text{C}$ . In contrast to the present findings, Lucassen et al. (2003) found higher CS activity per g liver during cold acclimation ( $3.5^\circ\text{C}$ ) in North Sea eelpout, however, it might be that cold compensation of CS activities only starts when temperature falls below  $5^\circ\text{C}$ . Lucassen et al. found no differences in hepatosomatic



indices, whereas, in the present study, cold acclimation led to elevated  $I_H$  in *Z. viviparus*. It may well be that long-term cold acclimation (10 months in this study versus 25 days in the study by Lucassen et al. 2003) led to a down-regulation of specific CS capacities associated with an increase in  $I_H$ .

In general, the highly positive cold compensation of CS but not of IDH activities, shown by higher specific as well as total CS activities in *P. brachycephalum* compared to *Z. viviparus* both acclimated to 5°C, is likely due to the role of mitochondria not only in energy metabolism but also in anabolic processes, with CS providing excess citrate, for example for enhanced lipid synthesis in the cold (for review cf. Pörtner 2002a, b).

Isocitrate dehydrogenase displayed similar  $Q_{10}$  and  $E_A$  values as proton leakage rates (state  $IV_{ol}$ ) and might be involved in setting the mitochondrial proton leakage rate as discussed by Pörtner et al. (2000). High  $Q_{10}$  and  $E_A$  values indicate that proton leakage might involve an enzyme catalyzed process (Ellory and Hall 1987; Sazanov and Jackson 1994; Pörtner et al. 1999b). However, further investigations of the mechanism of proton leakage are required. The present data indicate that IDH and proton leakage capacities are, if at all, only loosely correlated since the changes in IDH activity did not correlate with the observed changes in  $IV_{ol}$  state.

#### Effect of warm acclimation on Antarctic eelpout

To our knowledge this is the first long-term warm acclimation experiment for a fish species from the stable cold Antarctic ecosystem (Eastman 1993). In former studies *Trematomus bernachii* and *T. newnesi*, both species belonging to the Antarctic endemic Nototheniidae, were acclimated to 4°C during 2–5 weeks only and this temperature is the highest to which these fish can be warm acclimated for several weeks (cf. Weinstein and Somero 1998). Some adjustments in metabolism due to temperature changes can occur in notothenioids. Gonzalez-Cabrera et al. (1995) found temperature-induced capacity changes for osmoregulation in Antarctic Notothenioids, *T. bernachii* and *T. newnesi*. Warm acclimation (4°C) led to decreased serum osmolality, which was a result of the positive compensation of  $Na^+ / K^+$ -ATPase in osmoregulatory tissues. In our study, *P. brachycephalum* was acclimated to 5°C for at least 10 months and during this time none of the fishes died. They even showed positive growth (data not shown). Furthermore, the lack of significant differences in condition factors and gonadosomatic indices (Table I) between the different species and acclimation groups, indicate that acclimation to 5°C does not stress *P. brachycephalum*. Evidently, *P. brachycephalum* is still able to undergo temperature compensation despite its life at stable cold temperatures. As in the study of Weinstein and Somero (1998), no temperature-induced changes were found in specific mitochondrial respiration rates but warm acclimation led to decreased aerobic

capacities in total liver. Mitochondrial oxidative phosphorylation (state III) and enzyme activities, CS and  $NADP^+$ -dependent IDH, were lower in total liver of warm versus cold acclimated *P. brachycephalum* due to decreased liver size and mitochondrial protein content in total liver. Since no data are available on liver size and mitochondrial liver protein of *T. bernachii*, it remains unclear whether Notothenioids also show adjustments in aerobic capacities at the whole organ level after warm acclimation.

Zoarcids are most abundant in the North Pacific and North Atlantic as well as in the polar regions of Arctica and Antarctica. They originated in the Eocene about 50 million years ago in the Northern Pacific. Around the Miocene they radiated into the polar waters of Antarctica (Anderson 1994). The confamilial species, the Antarctic eelpout *P. brachycephalum* and the temperate common eelpout *Z. viviparus* show very low levels of standard metabolic rate and a benthic sluggish lifestyle which generally characterizes this family. It might therefore be that in this group, the pressure is less to adjust aerobic metabolism to the permanent cold than in the more active notothenioids. Furthermore, it might be that *P. brachycephalum* has retained some features of a cold adapted but hypometabolic eurytherm, with elevated CS activities but mitochondrial protein contents similar to boreal eelpout acclimated to 5°C. The finding that *P. brachycephalum* permanently tolerated and grew at 5°C and, evidently, retained the capacity to warm acclimate by reducing liver aerobic capacity may relate to the presence of this species in warmer bottom waters of Antarctica. Zoarcids were not found at the same cold temperatures as the notothenioids (F. Mark, personal communication). Further investigations are required to reveal to what extent aerobic metabolism can be adjusted in other Antarctic ectotherms, or whether down-regulation of aerobic capacities during warm acclimation only exists in Antarctic Zoarcidea.

**Acknowledgements** The authors would like to thank M. Lucassen for assistance in developing the stabilisation buffer for analysis of  $NADP^+$ -dependent isocitrate dehydrogenase activity, T. Hirse for support in protein determinations, and E. Brodte for her helpful comments on the manuscript.

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