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Malekar VC, Morton JD, Hider RN, Cruickshank RH, Hodge S, Metcalf VJ. 2018. Effect of elevated temperature on membrane lipid saturation in Antarctic notothenioid fish. PeerJ 6:e4765  
<https://doi.org/10.7717/peerj.4765>

# Effect of elevated temperature on membrane lipid saturation in Antarctic notothenioid fish

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Homeoviscous adaptation (HVA) is a key cellular response by which fish protect their membranes against thermal stress. We investigated evolutionary HVA (long time scale) in Antarctic and non-Antarctic fish. Membrane lipid composition was determined for four Perciformes fish: two closely related Antarctic notothenioid species (*Trematomus bernacchii* and *Pagothenia borchgrevinki*); a diversified related notothenioid Antarctic icefish (*Chionodraco hamatus*); and a New Zealand species (*Notolabrus celidotus*). The membrane lipid compositions were consistent across the three Antarctic species and these were significantly different from that of the New Zealand species. Furthermore, acclimatory HVA (short time periods with seasonal changes) was investigated to determine whether stenothermal Antarctic fish, which evolved in the cold, stable environment of the Southern Ocean, have lost the acclimatory capacity to modulate their membrane saturation states, making them vulnerable to anthropogenic global warming. We compared liver membrane lipid composition in two closely related Antarctic fish species acclimated at 0 °C (control temperature), 4 °C for a period of 14 days in *Trematomus bernacchii* and 28 days for *Pagothenia borchgrevinki*, and 6 °C for 7 days in both species. Thermal acclimation at 4 °C did not result in changed membrane saturation states in either Antarctic species. Despite this, membrane functions were not compromised, as indicated by declining serum osmolality, implying positive compensation by enhanced hypo-osmoregulation. Increasing the temperature to 6 °C did not change the membrane lipids of *P. borchgrevinki*. However, in *T. bernacchii*, thermal acclimation at 6 °C resulted in an increase of membrane saturated fatty acids and a decline in unsaturated fatty acids. This is the first study to show a homeoviscous response to higher temperatures in an Antarctic fish, although for only one of the two species examined.

1 **Effect of elevated temperature on membrane lipid saturation**  
2 **in Antarctic notothenioid fish**

3

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## 15 Abstract

16 Homeoviscous adaptation (HVA) is a key cellular response by which fish protect their membranes  
17 against thermal stress. We investigated evolutionary HVA (long time scale) in Antarctic and non-  
18 Antarctic fish. Membrane lipid composition was determined for four Perciformes fish: two closely  
19 related Antarctic notothenioid species (*Trematomus bernacchii* and *Pagothenia borchgrevinki*); a  
20 diversified related notothenioid Antarctic icefish (*Chionodraco hamatus*); and a New Zealand  
21 species (*Notolabrus celidotus*). The membrane lipid compositions were consistent across the three  
22 Antarctic species and these were significantly different from that of the New Zealand species.  
23 Furthermore, acclimatory HVA (short time periods with seasonal changes) was investigated to  
24 determine whether stenothermal Antarctic fish, which evolved in the cold, stable environment of  
25 the Southern Ocean, have lost the acclimatory capacity to modulate their membrane saturation  
26 states, making them vulnerable to anthropogenic global warming. We compared liver membrane  
27 lipid composition in two closely related Antarctic fish species acclimated at 0 °C (control  
28 temperature), 4 °C for a period of 14 days in *Trematomus bernacchii* and 28 days for *Pagothenia*  
29 *borchgrevinki*, and 6 °C for 7 days in both species. Thermal acclimation at 4 °C did not result in  
30 changed membrane saturation states in either Antarctic species. Despite this, membrane functions  
31 were not compromised, as indicated by declining serum osmolality, implying positive  
32 compensation by enhanced hypo-osmoregulation. Increasing the temperature to 6 °C did not  
33 change the membrane lipids of *P. borchgrevinki*. However, in *T. bernacchii*, thermal acclimation  
34 at 6 °C resulted in an increase of membrane saturated fatty acids and a decline in unsaturated fatty  
35 acids. This is the first study to show a homeoviscous response to higher temperatures in an  
36 Antarctic fish, although for only one of the two species examined.

## 37 Introduction

38 When the cell membranes of fish and other poikilothermic organisms are subjected to thermal  
39 change, modifications in membrane lipids and fluidity may occur in order to maintain membrane  
40 properties and functions. Altered membrane composition in response to lower or higher  
41 temperature, known as homeoviscous adaptation (HVA), is observed across all poikilotherms  
42 (Hazel & Williams 1990). HVAs that occur over short periods during the lifetime of an individual  
43 are acclimatory adaptive changes; e.g. as observed in eurythermic temperate fish, which possess a  
44 broad thermal adaptable range. Acclimation to lower temperature results in increases in the  
45 proportion of unsaturated fatty acids in membranes, to allow optimal membrane fluidity to be  
46 maintained. This suggests a protective role of the homeoviscous response in short-term acclimation  
47 (Skalli et al. 2006; Snyder et al. 2012). In contrast to these acclimatory adaptive changes observed  
48 in non-Antarctic fish, the HVA response in Antarctic fish can be a long-term evolutionary adaptive  
49 change in response to the low temperatures experienced in the Southern Ocean. Antarctic  
50 notothenioid fish, for example, display an evolutionary adaptive mechanisms where their cell  
51 membranes possess an increased proportion of unsaturated fatty acids (Hazel 1995).

52 Marine Antarctic ectotherms are stenothermal as they experience negligible seasonal variation (-  
53 1.9 °C to approximately 1.8 °C), resulting in limited ability to adapt to temperature variation  
54 (Somero 2010), and increased vulnerability to climate change effects (Aronson et al. 2011). In  
55 temperate and tropical latitudes, marine ectotherms experience much greater seasonal variation in  
56 temperature, and are correspondingly more thermally tolerant or eurythermal (Aronson et al.  
57 2011). Evolution of stenotherms in a “stable ice bath” has involved many critical changes in the  
58 genome that facilitate life in extreme cold, such as losses of certain traits that are no longer required  
59 (Pörtner et al. 2007). Loss of heat shock response (HSR) has been observed in Antarctic fish  
60 (Hofmann et al. 2000), resulting in extreme stenothermality due to an incapability to minimise  
61 damage to their protein pool caused by elevated temperatures (Podrabsky 2009). Homeoviscous  
62 adaptation to the constant cold temperatures of the Southern Ocean is one of the key evolutionary  
63 adaptive changes in Antarctic notothenioid fish, but it is not known whether they have the capacity  
64 to change their membrane saturation states in response to warmer temperatures, as one trade-off  
65 cost for stenothermality may be reduced adaptive capacity.

66 Anthropogenic global warming (AGW) poses a threat to polar and especially stenothermal polar  
67 species, and there is a need to determine the impact of warmer temperatures on the acclimatory  
68 responses of these species, including cellular membrane remodelling. There is evidence that  
69 Antarctic fish may not exhibit an acclimatory HVA response to transient temperature changes, as  
70 unchanged membrane lipid saturation states were observed in Antarctic fish (*Trematomus*  
71 *bernacchii* and *T. newnesi*) acclimated to a temperature of 4 °C for five weeks (Gonzalez-Cabrera  
72 et al. 1995). However, these fish showed positive compensation with an increase in the Na<sup>+</sup>/K<sup>+</sup> (+)-  
73 ATPase activity and a decline in the serum osmolality, implying that membrane functions were  
74 not compromised in spite of the unmodified saturation states.

75 Temperature is also a major determinant of membrane-cholesterol levels, with high membrane  
76 cholesterol observed in warm-acclimated marine copepods (Hassett & Crockett 2009). Levels of  
77 cholesterol have been shown to increase at higher temperatures resulting in reduced membrane  
78 fluidity (Crockett 1998). It is not known whether Antarctic fish share this adaptive membrane  
79 cholesterol change in response to increased temperature.

80 This study aimed to investigate both evolutionary and acclimatory HVA responses in Antarctic  
81 fish, and brings together existing evidence, along with new experimental data, to understand the  
82 evolutionary adaptive response associated with cold tolerance. Firstly, to investigate evolutionary  
83 adaptive HVA in cold-water fish, we established the normal lipid saturation profile of liver tissue  
84 from three Antarctic fish species collected in their normal physiological temperature for  
85 comparison with a non-Antarctic New Zealand fish species. More specifically, we compared liver  
86 membrane lipid profiles of two closely related Antarctic species, *Pagothenia borchgrevinki* (PB)  
87 and *Trematomus bernacchii* (TB), and a more distantly related icefish species *Chionodraco*  
88 *hamatus* (CH) of Antarctic notothenioid fish, as well as the non-Antarctic Perciformes species  
89 *Notolabrus celidotus* (NC). Icefish have evolved a suite of physiological adaptations to account  
90 for their loss of haemoglobin (Kock 2005), following their diversification from the other Antarctic  
91 notothenioids, and we sought to determine whether CH had a different membrane lipid profile to  
92 the two closely related Antarctic fish species (PB and TB). Previous study indicates that  
93 erythrocyte membranes of icefish have fluidity consistent with those of TB, but with observed  
94 lipid differences (Palmerini et al. 2009). It is unknown whether the membrane lipid composition  
95 of other icefish tissues, especially liver, also differs from other notothenioids. Secondly, we

96 investigated the acclimatory response of Antarctic fish to higher temperatures by examining  
97 whether thermal stress at 4 °C and 6 °C resulted in membrane restructuring in two Antarctic fish  
98 species (PB and TB), as indicated by altered membrane saturation states and cholesterol content.  
99 We hypothesised that membrane saturation, the major thermal adaptive mechanism, would occur  
100 only at reduced levels in Antarctic notothenioid fish as a response to elevated temperatures due  
101 their stenothermal nature, and thus make them vulnerable to the effects of AGW.

## 102 **Materials and Methods**

### 103 **Fish samples and experimental design**

104 The fish species used in the study are described in Table 1, and details of the fish harvest and  
105 husbandry are provided in S1 (supporting information). The field study comprising thermal  
106 acclimation experiments were conducted in the laboratory facilities at Scott Base, Antarctica  
107 approved by Antarctica New Zealand (K058 - 2007/2008). The procedures of fish handling were  
108 approved by the Animal Ethics Committees at the University of Canterbury (AEC 2006/2R and  
109 2008/11R). Liver tissue from Antarctic notothenioid and non-Antarctic fish species sampled from  
110 their normal habitat were taken for the establishment of normal lipid profiles. PB and TB and CH  
111 were compared with the non-Antarctic fish NC, a common native New Zealand Perciformes  
112 species (Ayling & Cox 1982). NC is non-migratory and has a broad thermal range (eurythermal),  
113 experiencing daily and seasonal variations in temperature (Jones 1984). PB and TB samples  
114 comprised the pre-acclimation controls of the thermal acclimation experiment described in S1  
115 (supporting information), while sampling locations of CH and NC are provided in Table 1. Briefly,  
116 with respect to the thermal acclimation experiments, the fish were initially held at an ambient  
117 temperature of -1 °C. Fish were transferred to tanks for the experiment where the temperature was  
118 raised to the target temperature over 1 to 3 days and then maintained at that temperature. PB was  
119 acclimated for 28 days but TB species was only acclimated for 14 days primarily due to limitations  
120 in the Antarctic aquaria space and duration of the field season. Standard procedures were involved  
121 in liver tissue collection for all fish (S1 supporting information).

## 122 **Phospholipid fatty acid analysis**

### 123 **Lipid extraction**

124 The lipid extraction method followed that of Folch et al.(Folch et al. 1957) but was modified as  
125 follows. Total lipids were extracted from 0.2 g frozen liver tissue. Samples were ground under  
126 liquid nitrogen and suspended in 6 ml of dichloromethane/methanol 2:1, 0.01 % butylated  
127 hydroxytoluene (BHT). After sonication (W-225 from Watson Victor) for 5 minutes, 2 ml of 0.88%  
128 potassium chloride was added. The samples were vortexed for two minutes and centrifuged at  
129 1000 g for 5 minutes. The aqueous layer was re-extracted with 2 ml of dichloromethane/methanol  
130 2:1, 0.01 % BHT and the two organic layers combined and dried under nitrogen. Dried samples  
131 were then stored at 4 °C until fractionation.

### 132 **Lipid fractionation**

133 The lipid fractionation method followed that of Zelles (Zelles 1997), but was modified as follows.  
134 Phospholipids were separated by re-suspending the total lipid extracts in chloroform and loading  
135 on to a solid phase extraction column (Biotage isolate SI 500 mg 6 ml SPE). The sample was  
136 allowed to stand for 2 minutes in the column and the lipids sequentially eluted with 5ml of  
137 chloroform for elution of neutral lipids, 5ml of acetone for glycolipids and 5ml of methanol for  
138 phospholipids. The phospholipid fraction was dried with nitrogen then stored at 4 °C before  
139 proceeding with methylation.

### 140 **Methylation**

141 The tubes containing the evaporated samples were brought to room temperature and 1 ml of  
142 tetrahydrofuran: methanol (1:1v/v) was added, then vortexed for 30 seconds. 1 ml of 0.2M  
143 potassium hydroxide was added followed by 30 seconds vortex and incubation at 37 °C for 15  
144 minutes. After incubation, 2 ml of hexane: chloroform (4:1) plus 0.3 ml of 1M acetic acid and 2  
145 ml of deionised water were added and vortexed for 1 minute followed by centrifugation at 1000 g  
146 for 5 minutes. The top organic layer was transferred to a holding tube and 2 ml of hexane:  
147 chloroform (4:1) was added to the lower aqueous layer and vortexed for 1 minute followed by  
148 centrifugation at 1000 g for 5 minutes. The top organic layer was transferred to the holding tube  
149 containing the first organic fraction. The organic layer was evaporated under N<sub>2</sub> in a water bath at



150 37 °C. Hexane (50 µl) was added to the evaporated organic layer and this was then transferred to  
151 a 150 µl insert with a poly spring held in an amber vial for GC analysis.

## 152 **Gas chromatographic separation**

153 Fatty acid methyl esters were analysed on a Shimadzu GC-2010 Gas Chromatograph (Shimadzu,  
154 Tokyo, Japan) fitted with a silica capillary column (Varian CP7420, 100m, ID 0.25mm, film  
155 thickness 0.25µm, Serial # 6005241) and helium flow 0.96ml/minute. The split ratio was 15 to 1  
156 and the injector temperature was 250 °C. The initial column temperature was 45 °C for 4 minutes,  
157 then ramped at 13 °C/minute to 175 °C held for 27 minutes before another ramp of 4 °C/minute to  
158 215 °C. This temperature was held for 35 minutes before a final ramp 25 °C/minute to 245 °C for  
159 5 minutes. All GLC conditions were based on adapting the initial conditions indicated by Lee and  
160 Tweed (Lee & Tweed 2008). A flame ionisation detector was used at 310 °C and fatty acids were  
161 identified by comparison of retention times to standards (GLC 463, NuChek). Known fatty acids  
162 are reported as a percentage of total fatty acids and fatty acids less than 1% were not reported.

## 163 **Membrane cholesterol analysis**

164 Cholesterol was extracted with dichloromethane: methanol from 50 mg of liver tissue re-  
165 suspended in 1 ml of 2-methoxymethane, then stored at -80 °C (Gonzalez et al. 2013). The free  
166 cholesterol was measured using the cholesterol fluorometric assay 10007640 following the  
167 manufacturer's instructions (<https://www.caymanchem.com/pdfs/>, Kit item number 10007640)  
168 and read on a Fluorostar omega microplate reader (BMG Labtech).

## 169 **Plasma osmolality determination**

### 170 **Collection and storage of plasma samples**

171 Blood samples from the experimental Antarctic fish (TB, PB) were collected at Scott Base Wet  
172 Laboratory. The temperature of the Wet Laboratory was constantly below 5 °C. The experimental  
173 fish were anaesthetised for five minutes by administration of 0.1 g L<sup>-1</sup> solution of MS222 (ethyl  
174 m-amino benzoate methane sulphonate) dissolved in sea water. Blood samples were immediately  
175 drawn by cardiac puncture with a 25 gauge needle. Blood volume of 0.5 to 1.0 ml was collected  
176 into a tube containing anti-coagulant. The collected blood was centrifuged at 3000 g for two  
177 minutes for the plasma separation. The resultant blood plasma was collected and snap frozen in

178 liquid nitrogen and transported to New Zealand in an insulated container containing dry ice and  
179 then stored at  $-80\text{ }^{\circ}\text{C}$ . Plasma from fish samples from both the species acclimated to  $4\text{ }^{\circ}\text{C}$  and the  
180 control temperature of  $0\text{ }^{\circ}\text{C}$ , and collected at all the time-points, were taken for osmolality analysis.  
181 The plasma samples were thawed to room temperature and  $10\text{ }\mu\text{l}$  plasma aliquots were taken for  
182 osmolality determination. Osmolality was measured using a Wescor 5520 C vapour pressure  
183 osmometer, which was calibrated with standard solutions before the measurements.

## 184 **Calculations and statistics**

185 All statistical analysis was performed using Minitab v17 software. Comparison of lipid profiles of  
186 the different species was performed using principal component analysis (PCA) based on a  
187 correlation matrix. The raw data consisted of a matrix of the percent contribution of each  
188 phospholipid fatty acid in each sample. The data were not transformed prior to analysis. One way  
189 ANOVA followed by a Holm-Sidak post hoc test was performed to compare individual fatty acids  
190 among the four fish species.

191 Desaturase index (DSI) for  $\Delta 9$ -desaturase/Stearoyl-CoA desaturase (SCD) was calculated as the  
192 ratio of product to precursor of the individual fatty acids using the formula:  $\text{C16:1c9}/\text{C16:0}$  and  
193  $\text{C18:1c9}/\text{C18:0}$  (Cormier et al. 2014). Two particular unsaturated fatty acids ( $\text{C16:1c9}$  and  
194  $\text{C18:1c9}$ ) were used for DSI, as the ratio of ( $\text{C16:1c9}/\text{C16:0}$ ) and ( $\text{C18:1c9}/\text{C18:0}$ ) has been shown  
195 to correlate with Stearoyl-CoA desaturase activity, degree of desaturation and membrane fluidity  
196 in a previous study (Hsieh & Kuo 2005).

197 Two-way ANOVA was used to assess the effects of temperature (control  $0\text{ }^{\circ}\text{C}$  and acclimated  $4$   
198  $^{\circ}\text{C}$ ), acclimation time (days) and the interaction between temperature and time on plasma  
199 osmolality. A Holm-Sidak post hoc test was subsequently used to determine which treatments  
200 differed significantly. Remaining data analysis in the  $4\text{ }^{\circ}\text{C}$  and  $6\text{ }^{\circ}\text{C}$  thermal acclimation trials was  
201 performed using an unpaired Student t test.

## 202 **Results and Discussion**

### 203 **Novelty of the study and key results**

204 This is the first study to show that higher temperature acclimation can induce a homeoviscous  
205 response in an Antarctic fish species; the response was dominated by changes in membrane

206 unsaturation while membrane cholesterol remain unchanged. Our results also reveal that the  
207 presence of a homeoviscous response can vary depending on the Antarctic fish species. Thermal  
208 acclimation to 4 °C did not induce the HVA response in either of the Antarctic species TB or PB.  
209 However, an HVA response was induced in one of the Antarctic fish species, TB, when it was  
210 acclimated to a warming temperature of 6 °C. In addition, apart from palmitic acid which had  
211 similar levels in icefish and the non-Antarctic fish species, the membrane fatty acid composition  
212 of Antarctic fish species was found to differ from that of a non-Antarctic fish species at their  
213 respective typical environmental temperatures.

### 214 **Distinct phospholipid fatty acid composition of Antarctic species**

215 The first two principal components (PCs) of the PCA of the phospholipid profiles explained 76.4  
216 % of the variance in the data matrix. The PCA clearly separated the phospholipid profiles of the  
217 three Antarctic fish species (PB, TB and CH) from the samples obtained from the non-Antarctic  
218 species (NC) along PC1 (Figure 1). The non-Antarctic fish NC was associated with high  
219 proportions of saturated fatty acids (C18:0) and the PUFA (C20:4), while the Antarctic species (all  
220 three) were associated with high MUFAs (C16:1c9, C18:1c11, C20:1c11 and an unknown  
221 MUFA), and PUFA (C20:5) (Figure 1; Table 2). Within the Antarctic species, the phospholipid  
222 profiles of closely-related species TB and PB were separated from those of the more distantly-  
223 related CH along PC2 (Figure 1). CH was associated with relatively high proportions of the SFA  
224 C16:0 and the MUFAs C18:1c9 and 20:1c11, whereas PB and TB had higher levels of the MUFA  
225 C16:1c9 and the unknown MUFA.

226 Generally the eurythermal NC had a significantly higher total SFA and lower total MUFA when  
227 compared to the Antarctic fish species, but this distinction was not specific for the total PUFA  
228 (Table 2). Our results suggest that the Antarctic fish species membrane fatty acid profiles are  
229 relatively consistent and distinct when compared to the eurythermal species (NC) (Figure 1; Table  
230 2). Stenothermal fish species such as Antarctic fish exist in constant cold and have a narrow  
231 thermal adaptable range, and have been reported to have higher percentages of unsaturated fatty  
232 acids than temperate fish or eurythermal fish (Logue et al. 2000). Similarly liver microsomes of  
233 Antarctic fish *Disostichus mawsoni* had higher percentages of MUFA when compared to the  
234 temperate fish such as trout and carp (Römisch et al. 2003). In our study it's the overall MUFA  
235 and some specific PUFA that are higher in Antarctic than the non-Antarctic fish species suggestive

236 of a central role of MUFA than PUFA in cold adaptation for Antarctic fish, and this phenomenon  
237 is considered as part of adaptive homeoviscous response in the fish acquired over their  
238 evolutionary history (Cossins 1977; Hsieh & Kuo 2005; Trueman et al. 2000; Williams & Hazel  
239 1995). A key question of this study was to determine whether the recently diversified icefish (CH)  
240 differ in their membrane lipids when compared to the other Antarctic fish species. This study  
241 shows that proportions of saturated fatty acids (SFAs), primarily palmitic acid (C16:0), were  
242 similar in the Antarctic species CH and the non-Antarctic species NC; both of these species had  
243 significantly higher levels of C16:0 compared to the Antarctic species TB and PB (Table 2). Higher  
244 proportions of C16:0 in the membranes of the icefish liver could be one feature acquired after  
245 diversification from the other Antarctic species. A previous study on the erythrocyte membrane  
246 lipids of CH showed higher levels of unsaturated longer chain fatty acids such as C:20–C:22,  
247 while shorter chain fatty acids such as C:16 and C:18 became unsaturated in TB, with both species  
248 having consistent membrane fluidity (Palmerini et al. 2009). The icefish species could thus have  
249 evolved specific adaptations in liver membrane lipids, such as higher C16:0 levels in liver  
250 membranes, as shown in the present study, and unsaturation of longer chain fatty acids in  
251 erythrocyte cell membranes (Palmerini et al. 2009).

252 Palmitic acid was significantly lower for the two closely related Antarctic species (TB and PB)  
253 than the New Zealand species NC, and the other Antarctic species CH (Table 2), and also formed  
254 the major fraction of the total saturated fatty acids in the Antarctic fish species. Stearic acid (18:0)  
255 was significantly lower in all three Antarctic species and formed the minor fraction. Palmitic acid  
256 has a role in cold adaptation of membranes (Farkas et al. 1994) and may be the reason for the  
257 predominance of palmitic acid among the saturated fatty acids in our study (Table 2). These results  
258 align with a study comparing the phospholipid compositions of muscle tissue in 15 marine species  
259 from the southeast Brazilian coast and two species from East Antarctica, where palmitic acid  
260 comprised 54–63% of the total SFA content (Visentainer et al. 2007); and another study examining  
261 the total fatty acid content for all organs in two Antarctic species, *Notothenia coriiceps* and  
262 *Notothenia rossii*, where palmitic acid represented 16–30% of the total FA content for all organs  
263 (Magalhaes et al. 2010). Apart from high palmitic acid in Antarctic fish, increases in palmitic acid  
264 due to cold acclimation was observed in a study comparing two confamilial species from different  
265 thermal habitats in the muscle of Antarctic eelpout, *Pachycara brachycephalum* in comparison to  
266 the temperate eelpout *Zoarces* (Brodte et al. 2008).

## 267 **Components of MUFA enhance membrane fluidity**

268 All three Antarctic fish species were associated with high levels of monounsaturated fatty acids  
269 (MUFA) associated with membrane fluidity, such as palmitoleic acid (C16:1c9), *cis*-vaccenic acid  
270 (C18:1c11), eicosenoate (C20:1c11) and total MUFA (Figure 1; Table 2). Other studies have  
271 reported high *cis*-vaccenic acid in membranes of the Antarctic fish *Pleuragramma antarcticum*  
272 (Mayzaud et al. 2011), high latitude fish of the sub-Arctic (Murzina et al. 2013), and  
273 *Caenorhabditis elegans* worms exposed to cold (Murray et al. 2007). *Cis*-vaccenic acid has been  
274 shown to enhance glucose transport in adipocytes (Pilch et al. 1980) and serotonin transport in  
275 endothelial cells (Block & Edwards 1987). The conformation of unsaturated *cis*-vaccenic acid  
276 presents a possible structural advantage and has a potential role in maintaining membrane fluidity,  
277 which may be the reason for its selective incorporation in the membranes of Antarctic fish. Lower  
278 growth temperature has also been shown to increase the amount of *cis*-vaccenic acid in *E.coli* and  
279 decrease the amount of palmitic acid incorporated in their membranes (Marr & Ingraham 1962).

## 280 **EPA could offer additional roles other than membrane fluidity**

281 Antarctic fish species had significantly lower levels of arachidonic acid (ARA, C20:4c5, 8, 11, 14)  
282 and higher levels of eicosapentaenoic acid (EPA, C20:5c5, 8, 11, 14, 17) than non-Antarctic  
283 species (Figure 1; Table 2). Levels of docosahexaenoic acid (DHA, C22:6c4, 7, 10, 13, 16, 19)  
284 were not significantly different between Antarctic and non-Antarctic species (Table 2). Higher  
285 EPA proportions in the Antarctic fish species included in our study is in alignment with high EPA  
286 levels observed in muscle phospholipids of Antarctic fish from the Weddell and Lazarev Seas  
287 (Hagen et al. 2000), in Antarctic silverfish, *Pleuragramma antarcticum* (Mayzaud et al. 2011), in  
288 cold acclimated fresh water alewives (*Alosa pseudoharengus*) (Snyder et al. 2012) and in cold  
289 acclimated *Caenorhabditis elegans* (Murray et al. 2007). Higher EPA in Antarctic species, and  
290 high EPA induced by cold acclimation in other species, suggest that EPA may play a role  
291 associated with cold tolerance, such as anti-inflammation or membrane stabilization. It has been  
292 suggested that DHA may possess a structural advantage over EPA in contributing to membrane  
293 fluidity due to the expanded molecular conformation of DHA (Hashimoto et al. 2006). We did not  
294 see an increase in DHA and perhaps MUFA perform this role in Antarctic species. EPA, but not  
295 DHA, has been shown to be a potent anti-inflammatory agent, whereas ARA is highly pro-  
296 inflammatory (Sears & Ricordi 2011; Seki et al. 2009). Hypercholesterolaemic rats, in whose

297 membrane fluidity is reduced, have been shown to display increased membrane fluidity in their  
298 platelets when fed DHA but not when fed EPA (Hashimoto et al. 2006). EPA may help in  
299 stabilization of hyper fluid membranes, as indicated by a study of the bacterium *Shewanella*  
300 *violacea* (Usui et al. 2012). EPA is one of the major (n-3) PUFAs present in the membranes of the  
301 Antarctic fish and contrary to other studies we do not observe correlation of DHA with membrane  
302 unsaturation, suggestive of modulation of particular fatty acids in HVA response. How these fatty  
303 acids (EPA, DHA and MUFA) contribute to fluidity and any other roles need further investigation  
304 in a larger range of fish species.

### 305 **Lack of distinction of membrane cholesterol between Antarctic fish and a New** 306 **Zealand fish species.**

307 Membrane cholesterol was higher in the non-Antarctic New Zealand species NC than the Antarctic  
308 species PB, but not different to CH and TB (Figure 2). In general, ectotherms adapted to lower  
309 temperature have shown to have reduced cholesterol levels primarily for maintenance of fluid state  
310 of membranes (Crockett 1998). Contrary to the trend of a direct relationship with membrane  
311 cholesterol and habitat temperature, a higher percentage of cholesterol in muscle was observed in  
312 the higher Arctic fish species *Leptoclonus maculatus* in comparison to the related sub-Arctic  
313 species *Lumpenus fabricii* (Murzina et al. 2013). Currently there is limited data on the membrane  
314 cholesterol of Antarctic fish species. Our study showed cholesterol content varies with species,  
315 rather than the habitat temperature, a similar finding to those of (Palmerini et al. 2009) where  
316 cholesterol in erythrocyte ghost membranes was highest in CH, followed by the non-Antarctic  
317 species *Anguilla anguilla*, and then lower in other Antarctic and non-Antarctic species. Thus,  
318 membrane cholesterol from further Antarctic species and from different tissues needs to be  
319 determined to establish its role in homeoviscous adaptation.

### 320 **Lack of homeoviscous response in Antarctic species at 4 °C thermal acclimation**

321 Thermal acclimation at 4 °C did not induce the major common cellular homeoviscous response in  
322 either the pelagic species (PB) or the benthic species (TB) after 28 or 14 days, respectively ( Table  
323 3). There was no change in the desaturase index  $D(C16:1c9/C16:0)$  and  $(C18:1c9/C18:0)$  in either  
324 species (Table 3). In TB, thermal acclimation changed the PUFA profile with a decrease in EPA  
325 ( $C20:5c$  5, 8,11,14,17) levels and an increase in the amount of DHA ( $C22:6c$ 4,7,10,13,16,19)  
326 (Table 3). As explained above, EPA levels may have a specific function in the extreme cold,

327 perhaps in stabilizing membranes (Usui et al. 2012), or a protective role by reducing inflammation  
328 (Sears & Ricordi 2011; Seki et al. 2009). The present findings of unchanged saturation states for  
329 PB and TB align with previous thermal acclimation experiments at 4°C in the benthic Antarctic  
330 notothenioid species *T. bernacchii* and *Trematomus newnesi*, where membrane unsaturation states  
331 were unchanged and there was no sign of an HVA response in the membranes of gills, kidneys,  
332 liver and muscle (Gonzalez-Cabrera et al. 1995). Similarly, mitochondrial membrane saturation  
333 states were also unchanged upon thermal acclimation and acidification, in the Antarctic species  
334 *Notothenia rossii* acclimated at 7 °C and the sub-Antarctic species *Lepidonotothen squamifrons*  
335 acclimated at 9 °C (Strobel et al. 2013). Our findings have extended these observations to a  
336 cryopelagic species (PB), as well as confirming the lack of change in membrane saturation state  
337 in the benthic species TB.

### 338 **Thermal acclimation has no effect on membrane cholesterol in the Antarctic species**

339 Cholesterol is known to counter the effects of increased temperature on membrane lipids and an  
340 increase in cholesterol is often observed at high temperatures (Crockett 1998). The structure of  
341 cholesterol mimics phospholipid structure and intercalates in the phospholipid membrane bilayer,  
342 resulting in an increase in membrane order and a reduction in membrane fluidity (Crockett 1998).  
343 However, the membrane cholesterol in PB as well as TB was unaffected by thermal acclimation  
344 (Figure 3;  $P > 0.05$ ). This may be a tissue-specific effect as increased temperature resulted in a  
345 significant decline of cholesterol in the gill membranes of goldfish, but had no effect on the brain  
346 and liver cholesterol concentration (Gonzalez et al. 2013).

### 347 **Thermal acclimation results in a decline in plasma osmolality in both Antarctic** 348 **species**

349 Plasma osmolality gives an indication of the functioning of membranes. An inverse relationship  
350 exists between serum osmolality and water temperature. In an analysis of 11 teleost species, the  
351 serum concentration of Antarctic species was higher than the temperate species (Dobbs & DeVries  
352 1975). Fish inhabiting cold waters have high serum inorganic ion concentrations and these  
353 inorganic ions have been shown to have protective roles in freezing avoidance by decreasing the  
354 melting point (O'Grady & DeVries 1982). The plasma osmolality change over the 28 days of  
355 thermal acclimation at 4 °C in PB is presented in Figure 4. Overall, irrespective of days of  
356 acclimation the osmolality at 4 °C was significantly lower in PB ( $P < 0.01$ ), while a numerical but

357 non-significantly decline with temperature increase was observed for TB. At Day 2, the plasma  
358 osmolality was significantly higher in thermally acclimated PB, but not in TB. This transient  
359 increase in osmolality has been attributed to increased efflux of water and retention of ions, mainly  
360 due to the alteration of permeability to ions brought about by the release of stress hormones,  
361 cortisol and catecholamine (Lowe & Davison 2005). It is unclear why this transient increase in  
362 osmolality is seen in PB but not TB. The osmolality fell in both species after Day 3 of thermal  
363 acclimation and the reduction was significant at Day 7 ( $P < 0.01$ ). Plasma osmolality in PB at 0 °C  
364 over the 28 days of acclimation remained unchanged ( $P > 0.05$ ). The plasma osmolality showed a  
365 decreasing trend over the 14 day acclimation to 4 °C in TB, but this was not statistically significant  
366 (Figure 4). In our study, thermal acclimation caused a decline in serum osmolality for PB. Other  
367 studies have also shown reduced osmolality upon thermal acclimation (Gonzalez-Cabrera et al.  
368 1995; Guynn et al. 2002; Hudson et al. 2008; Lowe & Davison 2005) which in some cases has  
369 been attributed to increased Na<sup>+</sup>/K<sup>+</sup> -ATPase activity (Guynn et al. 2002). The ability of these  
370 fish to control osmolality indicated that membranes were still functioning at 4 °C.

### 371 **Thermal acclimation at 6 °C results in an HVA response in *T. bernacchii*, but not in** 372 **the pelagic species *P. borchgrevinki***

373 One of the key HVA responses is the change in the saturation states of membrane phospholipids  
374 (Hazel 1995). TB exhibited an HVA response at 6 °C (Figure 5), as shown by the increase in  
375 overall SFAs due to an increase in stearic acid, along with a decline in MUFA component  
376 eicosenoic acid (C20:1c11), total MUFAs and the PUFA component EPA (C20:5c5,8,11,14,17),  
377 while a significant increase in DHA (C22:6c4,7,10,13,16,19) was observed. Saturated fatty acids  
378 reduce membrane fluidity and offset the effects of increased temperature (Hazel 1995). Previous  
379 studies of non-Antarctic fish species have shown that warm acclimation resulted in increased SFA  
380 and a decline in PUFAs *viz.*, EPA, DHA and ARA in brain phospholipids of *Dicentrarchus labrax*  
381 (Skalli et al. 2006), which has also been seen in fresh water alewives (*Alosa pseudoharengus*)  
382 (Snyder et al. 2012). In yellow perch (*Perca flavescens*) warm acclimation resulted in decline of  
383 MUFA and PUFA in muscle phospholipids (Fadhlaoui & Couture 2016). Although, the  
384 mechanism of HVA response upon warm acclimation is primarily dominated by a decrease in  
385 unsaturation, within this we observed an increase in DHA upon warm acclimation in TB at 4 °C  
386 (Table 3) and at 6 °C (Figure 6), suggesting that particular fatty acids are modulated by temperature  
387 which could differ with tissue type and individual fish species. Tissue specific responses were also



388 observed when warm acclimation induced an increase in DHA and palmitic acid in goldfish liver,  
389 but not in brain, gill and muscle membrane lipids (Gonzalez et al. 2013) and also the role of DHA  
390 have shown to vary among the eurythermal and stenothermal fish (Brodte et al. 2008). In  
391 eurythermal fish DHA is involved in cold acclimation as seen by increase in DHA of mitochondrial  
392 phospholipids with cold acclimation in rainbow trout (Guderley et al. 1997), similarly cold  
393 acclimation in carp resulted in DHA increase in liver phospholipids (Farkas et al. 1980). While in  
394 this study, increase in DHA in stenothermal Antarctic fish TB with temperature increase suggest,  
395 DHA does not participate in cold adaptation. Similarly in the Antarctic fish *Pachychara*  
396 *bracycephalum*, high DHA correlated with temperature of highest growth in the muscle and liver  
397 tissue suggestive of a role in growth rather than with cold adaptation (Brodte et al. 2008). Thus  
398 our study supports the dual role of DHA depending on thermal environment of fishes.

399 In PB we found that 6 °C did not induce a significant HVA response (Figure 6, although there was  
400 a decline in the MUFA component eicosenoic acid. Warm acclimation in both TB and PB caused  
401 a significant reduction of eicosenoic acid (Figure 5 and 6). At their normal environmental  
402 temperature, these fish are found to have high proportions of eicosenoic acid in their membranes,  
403 as shown in analysis of the general phospholipid profile (Figure 1), when compared to the New  
404 Zealand species, in which it was not detected. A similar role of eicosenoic acid in HVA response  
405 was observed in warm acclimated goldfish, with a decrease in the percent eicosenoic acid of brain  
406 and muscle phospholipids (Gonzalez et al. 2013). Apart from the reduction in eicosenoic acid, a  
407 major HVA response was not seen in PB. Other tissues may need to be analysed to confirm the  
408 apparent lack of a significant HVA response in PB. For example, the warm acclimation of  
409 *Dicentrarchus labrax* resulted in an HVA response in the brain, rather than the liver (Skalli et al.  
410 2006). In another study, warm acclimation of the Antarctic species *Notothenia rossii* at 7 °C and  
411 *Lepidonotothen squamifrons* at 9 °C did not result in an HVA response in mitochondrial  
412 membranes (Strobel et al. 2013). PB has a higher degree of thermal plasticity (Franklin et al. 2007)  
413 and higher upper lethal temperature compared with TB (Somero & DeVries 1967). Thus,  
414 temperatures greater than 6 °C may be required to induce an HVA response in PB.

#### 415 **Desaturase index correlates with membrane saturation state**

416 In the present study the desaturase index (DSI) (C16:1c9/C16:0) and (C18:1c9/C18:0) were shown  
417 to correlate with the saturation states of the membrane and DSI has been used as a surrogate for  
418 the measurement of stearyl-CoA desaturase (SCD) enzyme activity in membrane remodelling in

419 response to temperature (Fadhlaoui & Couture 2016). The enzyme SCD plays a key role in  
420 unsaturation of SFA by catalysing the synthesis of MUFA, primarily by the introduction of the  
421 first double bond between the C9 and C10 position of the fatty acid which results in increased  
422 membrane disorder and enhanced fluidity (Paton & Ntambi 2009). The Antarctic species had a  
423 high desaturase index (C16:1c9/C16:0) compared to the non-Antarctic species, whereas this trend  
424 is not specific for the DSI (C18:1c9/C18:0) (Figure 7). High DSI (C16:1c9/C16:0) in the Antarctic  
425 fish species could be attributed to an increase in the MUFA palmitoleic acid C16:1c9 reflecting  
426 higher desaturation of palmitic acid by SCD. Furthermore, in this study there was significant  
427 decline in DSI (C16:1c9/C16:0) upon thermal acclimation at 6 °C in *T. bernacchii* (Figure 8). A  
428 positive correlation does exist with the desaturase index and membrane saturation states, as  
429 previously established in two fish species, milk fish and the grass carp when subjected to cold  
430 acclimation from 25 °C to 15 °C over 21 days (Hsieh & Kuo 2005). Similarly higher desaturase  
431 index for SCD was observed in yellow perch (*Perca flavescens*) acclimated at 9 °C than at 28°C  
432 (Fadhlaoui & Couture 2016).

433

#### 434 **Conclusions and perspectives**

435 This study has established a consistent membrane lipid profile across three notothenioid Antarctic  
436 species, in contrast to a varying membrane lipid composition between Antarctic species and a non-  
437 Antarctic New Zealand species. The Antarctic fish exhibit an evolutionary HVA response, as  
438 reflected by high levels of unsaturated fatty acids and selective dominance of *cis*-vaccenic acid  
439 and EPA in their membranes. This calls for further analysis of a wide range of fish species from  
440 different thermal habitats to decipher the specific roles of *cis*-vaccenic acid and EPA in cold  
441 adaptation. Previously undetermined is whether Antarctic fish can protect their membranes by  
442 exhibiting the acclimatory HVA response, which may make them less vulnerable to the effects of  
443 AGW. Our findings suggest that at 4 °C neither of the closely related Antarctic species exhibited  
444 any significant HVA response either with phospholipid unsaturation or with membrane  
445 cholesterol, but membrane-associated functions such as osmoregulation remain uncompromised.  
446 Furthermore, acclimatory HVA response of membrane unsaturation was detected at 6 °C in the  
447 liver of the benthic species TB while this response was lacking in liver membranes of the  
448 cryopelagic species PB. In the present study HVA response was dominated by phospholipid

449 unsaturation with no change in membrane cholesterol and the potential role of cholesterol in HVA  
450 response in Antarctic fish still remain unclear. Future studies especially at higher temperature  
451 acclimation as well as in other tissues are needed to determine the role of membrane cholesterol  
452 to HVA response in Antarctic fish. In conclusion it appears that some Antarctic fish species can  
453 exhibit a limited HVA response to warming temperatures after a given acclimation period.  
454 However, this study has reinforced the need for further experimental work involving more species,  
455 over a wider range of acclimation temperatures and assaying multiple tissue types in order to  
456 ascertain the generality or specificity of acclimatory HVA responses in Antarctic fish.

457

458

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460

#### 461 **Acknowledgements:**

462 We thank Dr Adrian Paterson for input to manuscript preparation.

463

#### 464 **Data Availability**

465 Raw data has been supplied as a supplementary file

#### 466 **Supplemental Information**

467 Supporting information S1 Thermal acclimation experiment and sampling

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**Table 1** (on next page)

Fish species sampled and collection location

\* These fish were used for establishment of membrane lipid profiles (pre acclimation controls) and for thermal acclimation studies.

1

<b>Fish species</b>	<b>Family</b>	<b>Location</b>	<b>Adaptation Temperature(°C)</b>
<i>Trematomus bernacchii</i>	Nototheniidae	McMurdoSound, Antarctica*	-1- 1.9
<i>Pagothenia borchgrevinki</i>	Nototheniidae	McMurdoSound, Antarctica*	-1- 1.9
<i>Chionodraco hamatus</i>	Channichthyidae	Terra Nova Bay, Antarctica	-1- 1.9
<i>Notolabrus celidotus</i>	Labridae	Kaikoura, New Zealand	9-13

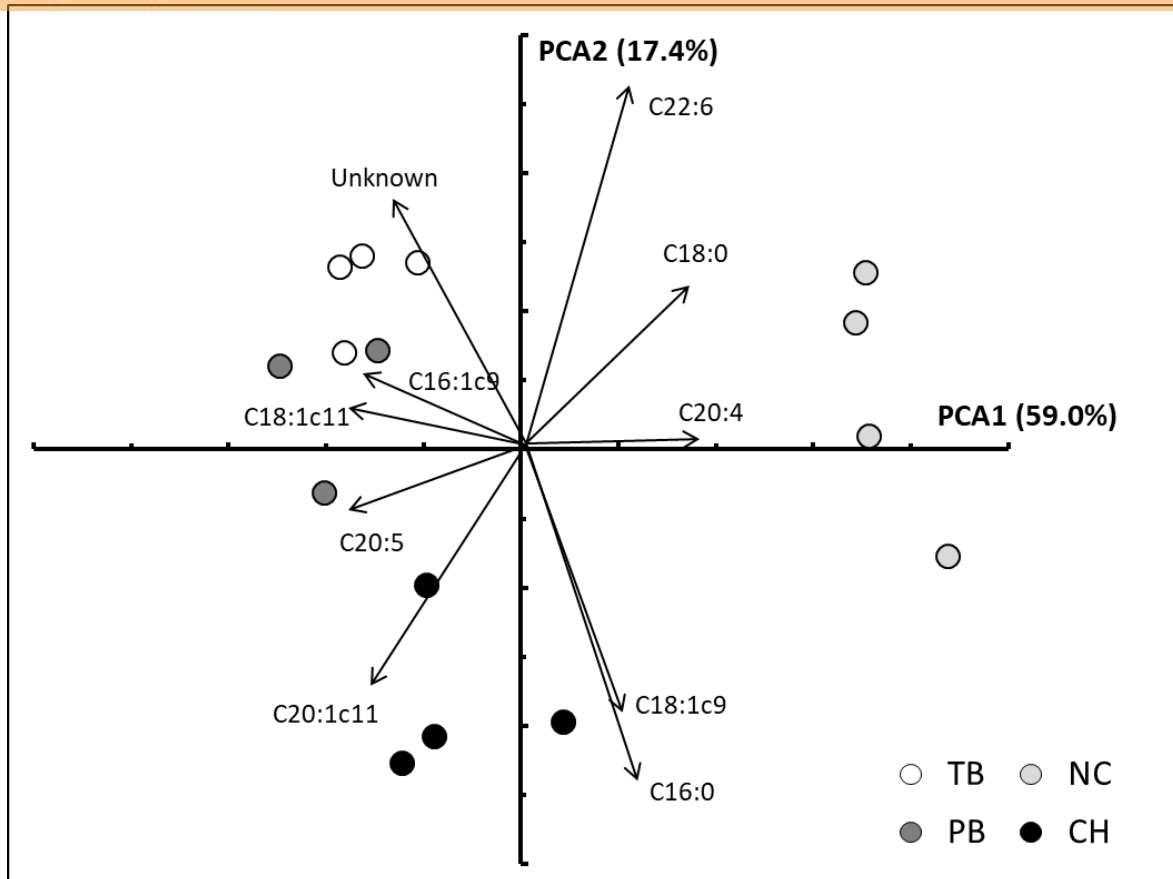
2



**Figure 1**(on next page)

PCA plot of the contribution of the phospholipid fatty acids to the principal components in liver tissue of the Antarctic species and non-Antarctic species.

Antarctic species *C. hamatus* (CH), *P. borchgrevinki* (PB), and *T. bernacchii* (TB) and the non-Antarctic species *N. celidotus*(NC).



**Table 2** (on next page)

Fatty acid composition of phospholipids in liver of Antarctic (CH, PB, TB) and non-Antarctic fish (NC) expressed as % of total phospholipid fatty acids.

Values are mean  $\pm$  SEM (n=4), except for *Pagothenia borchgrevinki* (n=3), nd=not detected. Significant differences among the species for each particular fatty acid are indicated by different letter codes (P<0.05).

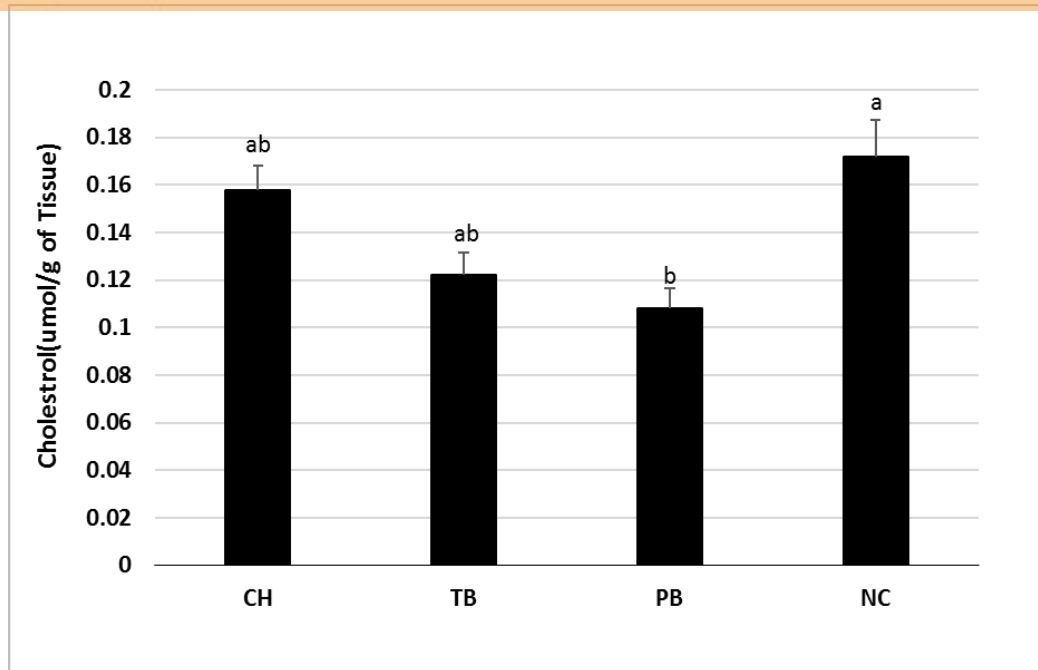
	<i>N. celidotus</i>	<i>C. hamatus</i>	<i>P. borchgreviniki</i>	<i>T. bernacchii</i>
C16:0	20.05 ± 0.51 <sup>a</sup>	20.43 ± 0.69 <sup>a</sup>	13.16 ± 1.11 <sup>b</sup>	13.08 ± 0.52 <sup>b</sup>
C18:0	12.64 ± 1.16 <sup>a</sup>	3.05 ± 0.56 <sup>b</sup>	4.16 ± 0.64 <sup>b</sup>	5.01 ± 0.29 <sup>b</sup>
<b>ΣSFA</b>	<b>32.69 ± 1.45<sup>a</sup></b>	<b>23.47 ± 0.40<sup>b</sup></b>	<b>17.32 ± 1.55<sup>c</sup></b>	<b>18.09 ± 0.68<sup>c</sup></b>
C16:1c9	0.48 ± 0.48 <sup>b</sup>	3.30 ± 0.73 <sup>a</sup>	4.01 ± 0.41 <sup>a</sup>	4.97 ± 0.35 <sup>a</sup>
C18:1c9	11.07 ± 1.70	10.51 ± 1.99	9.64 ± 1.12	7.42 ± 0.28
C18:1c11	3.22 ± 0.50 <sup>b</sup>	7.69 ± 0.59 <sup>a</sup>	9.84 ± 0.74 <sup>a</sup>	9.11 ± 0.49 <sup>a</sup>
C20:1c11	nd	5.04 ± 0.98	2.78 ± 0.14	3.62 ± 0.38
Unknown	nd	0.87 ± 0.50	2.93 ± 1.83	4.24 ± 0.81
<b>ΣMUFA</b>	<b>14.77 ± 1.39<sup>b</sup></b>	<b>27.40 ± 1.98<sup>a</sup></b>	<b>29.20 ± 0.38<sup>a</sup></b>	<b>31.99 ± 0.72<sup>a</sup></b>
C18:2c9, 12	0.62 ± 0.62	1.17 ± 0.41	nd	nd
C20:4c5, 8, 11, 14	9.08 ± 0.50 <sup>a</sup>	4.97 ± 0.35 <sup>b</sup>	3.13 ± 0.24 <sup>b</sup>	4.79 ± 0.33 <sup>b</sup>
C20:5c 5, 8, 11, 14, 17	14.00 ± 0.75 <sup>b</sup>	19.83 ± 0.80 <sup>a</sup>	22.52 ± 1.92 <sup>a</sup>	19.63 ± 1.08 <sup>a</sup>
C22:5c5	1.41 ± 0.47	nd	nd	nd
C22:6c 4, 7,10, 13, 16,19	28.06 ± 1.53 <sup>a</sup>	20.84 ± 1.44 <sup>b</sup>	23.54 ± 1.36 <sup>ab</sup>	25.50 ± 0.32 <sup>ab</sup>
<b>ΣPUFA</b>	<b>53.16 ± 1.96<sup>a</sup></b>	<b>46.80 ± 1.13<sup>b</sup></b>	<b>49.18 ± 0.50<sup>ab</sup></b>	<b>49.91 ± 0.73<sup>ab</sup></b>

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**Figure 2**(on next page)

Membrane Cholesterol concentration in the livers of Antarctic species *C. hamatus* (CH), *P. borchgrevinki* (PB), *T. bernacchii* (TB) and non-Antarctic species *Notolabrus celidotus* (NC).

Values are mean  $\pm$  SEM (n=4). Significant effects among species are indicated by different letters (P<0.05).



**Table 3**(on next page)

Fatty acid composition of phospholipids in the liver of *Trematomus bernacchii* (14 days acclimation) and *Pagothenia borchgrevinki* (28 days acclimation) acclimated at 0 °C and 4 °C.

Values are mean  $\pm$  SEM (n=4) and expressed in % of total phospholipid fatty acids. Significant effects of thermal acclimation are indicated by asterisks (P<0.05)

	<i>Trematomus bernacchii</i>			<i>Pagothenia borchgrevinki</i>		
	T0	T4	P-value	T0	T4	P-value
C16:0	16.01 ± 0.76	14.30 ± 0.53	0.13	13.79 ± 1.00	14.18 ± 0.34	0.74
C18:0	6.49 ± 0.30	7.71 ± 0.47	0.08	5.33 ± 0.86	4.20 ± 0.37	0.29
<b>ΣSFA</b>	<b>22.50 ± 1.00</b>	<b>22.01 ± 0.54</b>	<b>0.69</b>	<b>19.12 ± 1.5</b>	<b>18.38 ± 0.58</b>	<b>0.67</b>
C16:1c9	4.42 ± 0.78	3.09 ± 0.47	0.22	3.66 ± 0.38	4.17 ± 0.53	0.47
Unknown	2.44 ± 0.87	2.68 ± 0.11	0.81	0.43 ± 0.43	1.81 ± 0.67	0.15
C18:1c9	5.68 ± 0.26	5.98 ± 0.87	0.76	10.05 ± 0.68	11.27 ± 0.32	0.18
C18:1c11	8.55 ± 0.23	9.03 ± 0.83	0.61	9.50 ± 0.50	9.69 ± 0.39	0.78
C20:1c11	3.07 ± 0.27	3.26 ± 0.31	0.66	2.11 ± 0.20	2.05 ± 0.23	0.86
Unknown	1.93 ± 1.20	2.83 ± 1.10	0.59	0.72 ± 0.72	1.17 ± 0.70	0.67
<b>ΣMUFA</b>	<b>26.08 ± 1.20</b>	<b>26.87 ± 1.00</b>	<b>0.64</b>	<b>26.47 ± 0.83</b>	<b>30.15 ± 1.40</b>	<b>0.09</b>
C20:4c5,8,11,14	4.75 ± 0.73	4.68 ± 0.34	0.94	2.56 ± 0.10	3.49 ± 0.36	0.09
C20:5c5,8,11,14,17	24.24 ± 1.10	19.05 ± 0.72	0.01*	18.40 ± 1.30	17.64 ± 2.00	0.76
C22:6c4,7,10,13,16,19	22.43 ± 0.64	27.38 ± 0.93	0.01*	29.51 ± 2.90	30.35 ± 3.20	0.85
<b>ΣPUFA</b>	<b>51.42 ± 2.00</b>	<b>51.11 ± 0.55</b>	<b>0.89</b>	<b>50.47 ± 2.30</b>	<b>51.47 ± 1.50</b>	<b>0.73</b>
<b>DSI (Desaturase Index)</b>						
C16:1c9/C16:0	0.27 ± 0.04	0.21 ± 0.03	0.24	0.27 ± 0.02	0.30 ± 0.04	0.564
C18:1c9/C18:0	0.88 ± 0.06	0.78 ± 0.11	0.46	2.11 ± 0.52	2.76 ± 0.31	0.341

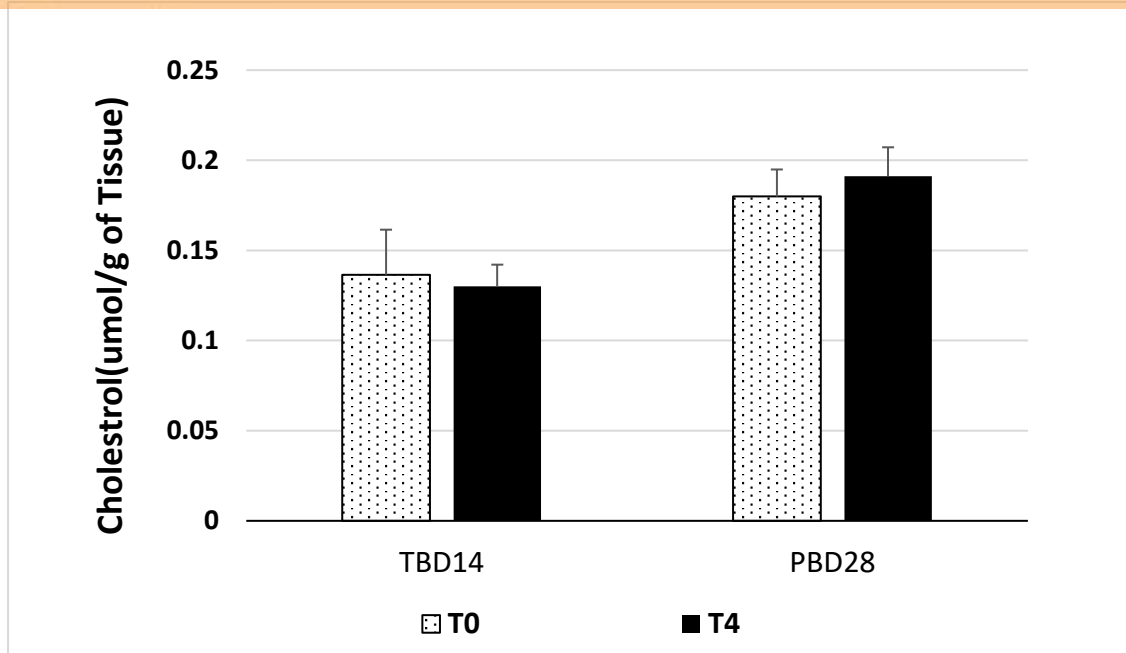
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**Figure 3**(on next page)

Effect of thermal acclimation on membrane cholesterol concentration in the livers of *T.bernacchii* (TB) and *P.borchgrevinki* (PB).

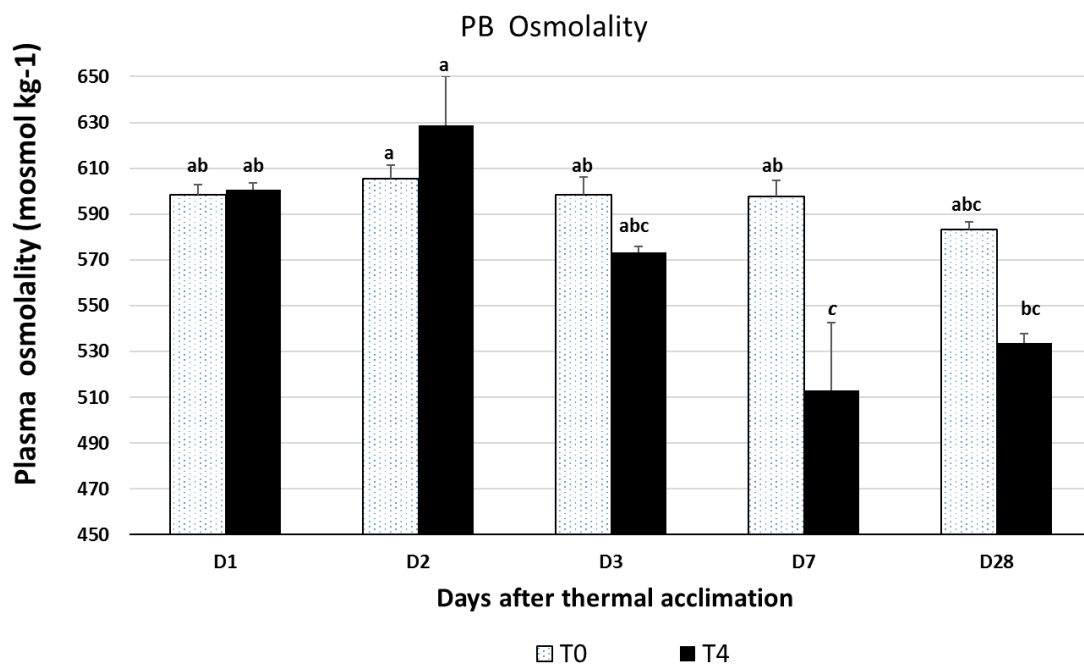
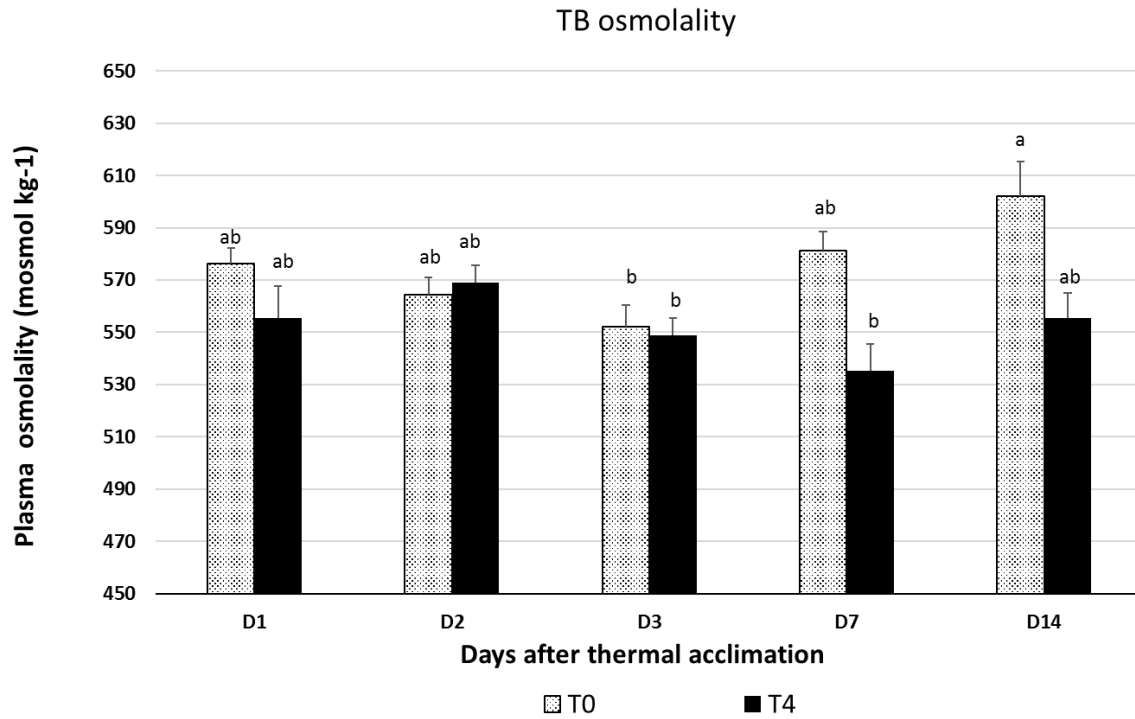
Membrane cholesterol was determined 14 days after thermal acclimation in TB and 28 days in PB. Values are means  $\pm$ SEM (n=4) for control temperature (T0: 0°C) and warm (T4: 4°C) acclimation.



**Figure 4**(on next page)

Plasma osmolality determined at various time points in *T.bernacchii* (TB) and *P.borchgrevinki* (PB).

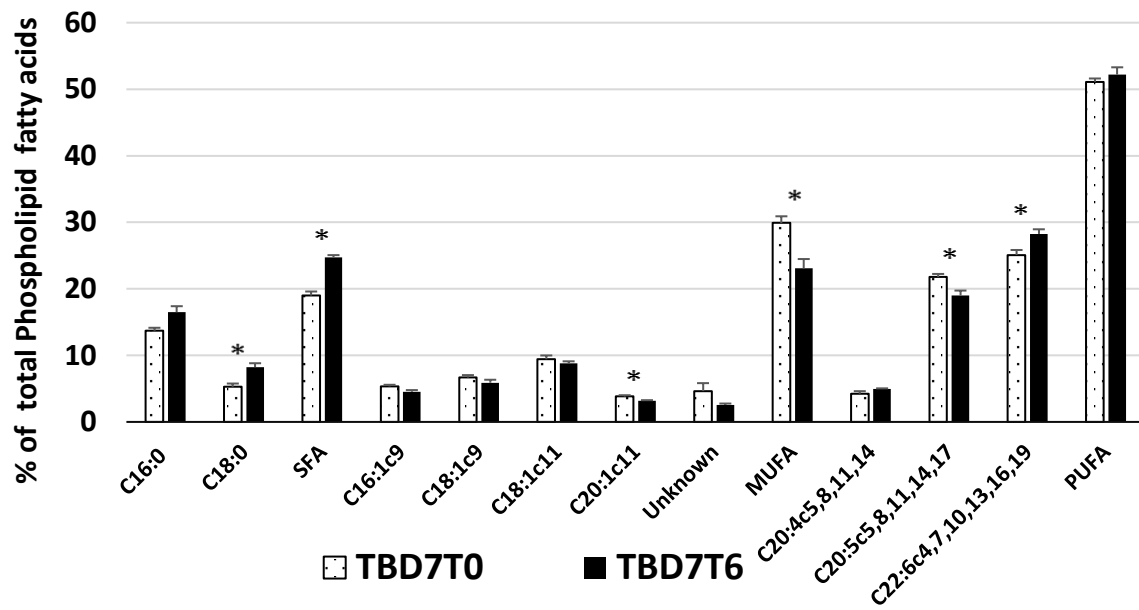
Plasma osmolality was determined 14 days after thermal acclimation in TB and 28 days in PB. Days after thermal acclimation D1, D2, D3, D7, D14 and D28 at 4 °C (T4) and the control temperature of 0 °C (T0). Values are  $\pm$ SEM (n=4). Significant effects of the interaction of thermal acclimation and days of acclimation are indicated by different letters.



**Figure 5**(on next page)

Phospholipid profile of *T. bernacchii* (TB) in liver after 7 days (D7) of thermal acclimation at 6 °C.

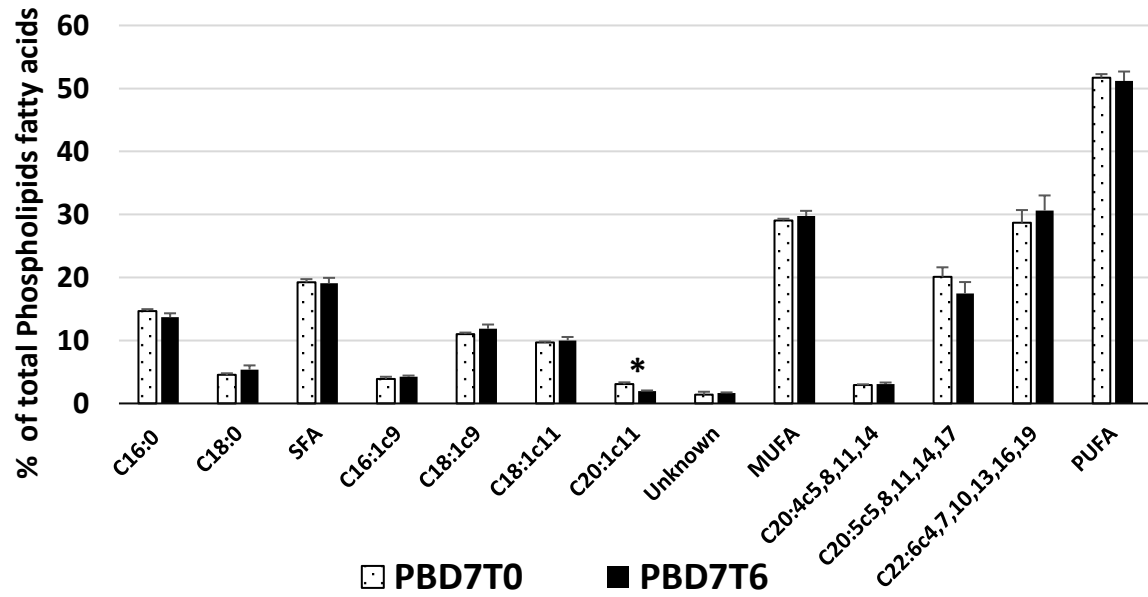
Values are means  $\pm$ SEM (n=4) for control temperature (T0: 0 °C) and warm (T6: 6 °C) acclimation (n=3). Significant effects of thermal acclimation are indicated by asterisks (P<0.05).



**Figure 6**(on next page)

Phospholipid profile of *P. borchgreviniki* (PB) in liver after 7 days (D7) of thermal acclimation at 6 °C.

Values are means  $\pm$  SEM (n=4) for control temperature (T0: 0 °C) as well as warm (T6: 6 °C) acclimation. Significant effects of thermal acclimation are indicated by asterisks (P<0.05).

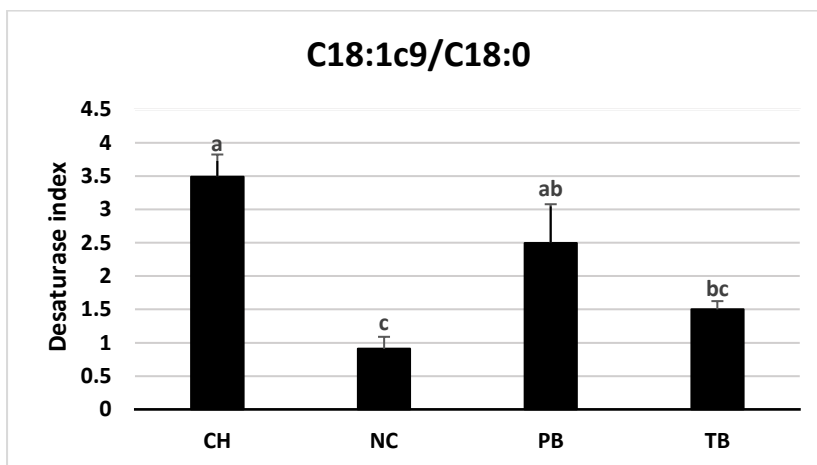
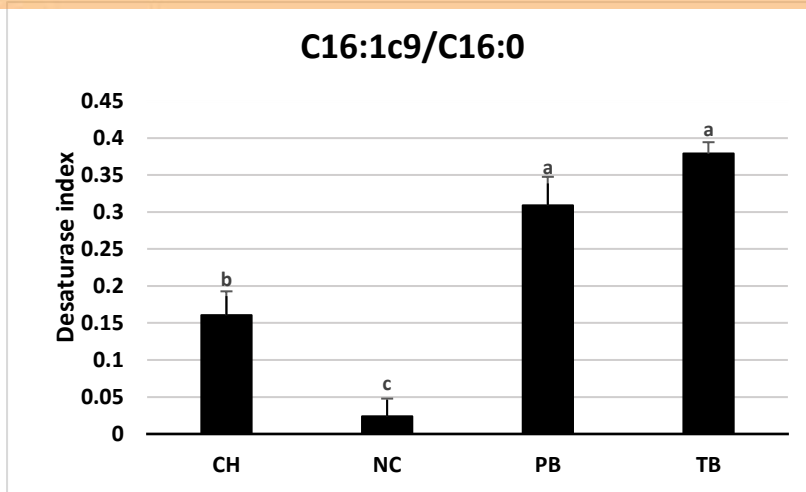




**Figure 7** (on next page)

Desaturase Index [C16:1c9/C16:0 (above), C18:1c9/C18:0 (below)] in livers of Antarctic species *C. hamatus* (CH), *P. borchgrevinki* (PB), and *T. bernacchii* (TB) and the non-Antarctic species *N. celidotus*(NC).

Values are mean  $\pm$  SEM(n=4). Significant effects among species are indicated by different letters (P<0.05).



**Figure 8**(on next page)

Changes in the Desaturation Index [C16:1c9/C16:0 (above), C18:1c9/C18:0 (below)] in the livers of *P. borchgrevinki* (PB) and *T. bernacchii* (TB) acclimated at 6 °C for 7 days.

Values are means $\pm$ SEM (n=4) for control temperature (T0: 0 °C) as well as warm (T6: 6 °C) acclimation. Significant effects of thermal acclimation are indicated by asterisks (P<0.05).

