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Laboratory conditioning modifies properties of gills mitochondria from the Pacific oyster *Crassostrea gigas*

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Abstract :

Although laboratory experiments allow greater control of environmental conditions than field studies, they have several drawbacks. To analyze physiological responses to forcing environmental variables, experimental conditions should mimic natural conditions as closely as possible. For filter-feeding organisms in particular, diet quality and quantity is one of the environmental parameters that can differ markedly between experimental and field conditions. In the hatchery, Pacific oysters, Crassostrea aigas. commonly show good physiological performance and growth on a mixed algal diet of Tisochrysis lutea, formerly Isochrysis aff. galbana clone Tahiti (T-Iso), and Chaetoceros calcitrans, presumably as it provides a good supply of essential polyunsaturated fatty acids (PUFA) as 20:4n-6, 20:5n-3 and 22:6n-3. The present study tests whether the fluctuating biotic and abiotic conditions in the field modify the structure and function of oyster mitochondria. One group of oysters was maintained in the intertidal zone, and the other group was fed the mixed diet in a nearby experimental hatchery under salinity and temperature conditions equivalent to those in the field. After 4 weeks of conditioning, the functional capacities and membrane lipid composition of gill mitochondria were measured. For essential polyunsaturated fatty acids, only the proportion of 20:5n-3 differed between field and laboratory oysters. and confirmed the capacity of the mixed diet T-Iso + C. gracilis, to provide the essential PUFA. Nevertheless, proportions of other FA (e.g., 22:5n-6 and non-methylene-interrupted FA) differed markedly between laboratory and field-conditioned oysters. Mitochondrial oxygen uptake, cytochrome c oxidase activity, content of cardiolipin and concentration of cytochrome b were significantly lower in laboratory-conditioned than in field-conditioned oysters. These results indicate that laboratory conditioning, although allowing similar growth and gonad maturation, only partially mimics conditions that allow C. gigas to maintain mitochondrial function. Although our experimental design cannot ascertain what difference between experimental laboratory and field conditions led to changes in membrane composition and mitochondrial function, differences in nutritional quality (other than known essential PUFA) and abiotic factors (e.g., oxygen availability, emersion or daily temperature fluctuations) had a major impact on mitochondrial properties in oysters.

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Introduction

Although laboratory experiments permit much more control over environmental conditions than field studies, they have drawbacks, including the adequacy and realism of laboratory conditions. Experiments in artificial settings attempt to simplify natural complexity by limiting variation to a small number of factors. However, the constant environmental conditions used in laboratory experiments may be inappropriate for animals that have adapted to fluctuating conditions and may impair physiological and cellular processes (Guerra et al. 2012). Oysters generally inhabit the intertidal zone where they are exposed to large fluctuations in oxygen concentration, temperature, food availability and quality, on hourly, daily and seasonal basis. For marine bivalves, changes in abiotic conditions, such as temperature, oxygenation, salinity and desiccation lead to regulated physiological adjustments that can affect growth and reproduction. At the cellular level, changes in mitochondrial function and membrane composition in response to abiotic factors have been demonstrated in bivalves (Glémet and Ballantyne 1995; Gillis and Ballantyne 1999; Guderley 2004; Sussarellu et al. 2013; Dudognon et al. 2013). This ability to adjust mitochondrial structure and function is thought to be critical in controlling energy production under changing environmental conditions (Guderley 2004; Bremer and Moyes 2011). However, whether fluctuating abiotic and biotic conditions affect mitochondrial structure and function in marine bivalves is unknown.

The use of an appropriate diet in the laboratory is primordial, given the influence of diet on energy acquisition for physiological acclimation and growth. Formulating an adequate diet for filter feeding organisms in hatcheries represents a particular challenge, as filter feeders typically ingest living (mainly phytoplankton) and inert particles suspended in the water column. Cultured phytoplankton is generally used during laboratory conditioning of bivalves (Martínez et al. 2000; Farías and Uriate 2006). Algal shape, size, toxicity, digestibility and biochemical composition affect the nutritional value of microalgal species (Brown et al. 1997). Trial and error has shown that microalgal species used to rear bivalves in hatcheries do not have the same nutritional value and lead to different rates of growth, survival, gametogenesis, embryogenesis, hatching and metamorphosis (Epifanio 1979; Enright et al. 1986; Delaunay et al. 1993; Soudant et al. 1996a, 1996b; Utting and Millican 1997; Rico-Villa et al. 2006; Pronker et al. 2008; Marshall et al. 2010).

A major biochemical difference among algae lies in their lipid and, particularly, fatty acid (FA) composition that in turn, affects the composition of the bivalves feeding upon them.

Various studies have established that the polyunsaturated fatty acid (PUFA) composition of bivalve cell membranes changes with microalgal FA composition (Delaunay et al. 1993; Soudant et al. 1999; Pennarun et al. 2003; Delaporte et al. 2003, 2005, 2006; Marshall et al. 2010; Dudognon et al. 2014). Because of their structural role in membranes, modifications of membrane FA composition can modulate membrane protein activities and physiological parameters in mammals (Yamaoka et al. 1988; Senault et al. 1990; Hulbert and Else 1999; Leonard et al. 2001; Hirunpanich et al. 2007), fish (Moya-Falcón et al. 2004; Hamza et al. 2008), and bivalves (Soudant et al. 1996a, 1996b; Delaporte et al. 2003; Delaporte et al. 2006). Given the compositional variety of microalgal FA, a mixture of the microalgae Tisochrysis lutea, formerly Isochrysis aff. galbana clone Tahiti (T-Iso) and Chaetoceros calcitrans or Chaetoceros gracilis is often used as an optimal supply of essential FA that allows adequate performance of oysters under laboratory conditions (Rico-Villa et al. 2006; Dudognon et al., 2014). Nonetheless, comparison of adult oysters fed two algae as monospecific diets (i.e. T-Iso and C. gracilis), shows that although the FA composition of mitochondrial membranes is strongly modified by the FA composition of monoalgal diet, very few functional parameters are modified (Dudognon et al. 2014). Such insensitivity of the oxidative capacities of oyster gill mitochondria to major diet-induced changes of membrane composition suggests that regulated changes and homeostatic mechanisms maintain critical mitochondrial membrane characteristics despite major differences in nutritional quality.

In the present study, we tested how laboratory conditions perform to mimic environmental conditions regarding oyster mitochondrial function. For this purpose, we compared physiological performance and mitochondrial properties of oysters kept under controlled laboratory conditions, with salinity and temperature in the experimental tanks similar to field conditions), and a same lot of oysters reared during the same period in the field where they were exposed to natural fluctuations of biotic and abiotic conditions. Physiological performance was assessed through growth and reproductive investment after 4 weeks of conditioning. Gill mitochondria were isolated from oysters before and after conditioning. We characterized mitochondrial oxidative capacities, cytochrome c oxidase (CCO) and citrate synthase (CS) activity, as well as contents of cytochrome a, b, c_1 and c, and membrane lipid composition.

Materials and methods

123124

125 Chemicals

- 126 All chemicals were purchased from Sigma (Saint Quentin Fallavier, France), unless
- mentioned otherwise.

128

129 Sampling procedures

- 130 In May 2010, one hundred Pacific oysters C. gigas, shell length ranging from 6 to 8.5 cm
- 131 (7.3 \pm 0.2, mean \pm standard error), originating from the same batch of spat, were obtained from
- a hatchery in Arcachon, France. After arriving in Brittany, oysters were placed in a mesh bag
- on a tray in the intertidal zone at Aber Benoît (North Brittany, 48° 32'N, 4° 30'W) near the
- 134 Argenton experimental station. One day after arriving, 50 were placed in a 800 L flow-
- through seawater tank at the Argenton experimental station (IFREMER, Argenton, France),
- with a water flow of 120 L.h⁻¹. In parallel, the other 50 oysters were kept in the mesh bag in
- the intertidal zone. The experimental tank was supplied with seawater pumped at mid-depth in
- a tidal dock and partly filtered (10 µm) through a bag filter. Thus, salinity and temperature in
- the experimental tank were similar to field conditions (from 12°C on May 20th, to 16°C on
- June 22nd). The tank was drained and cleaned three times a week.
- Samples were taken at the start of the experiment (T0, 20th of May 2010) and after 4 weeks of
- 142 conditioning (Tf, 22nd of June 2010). Total weight was measured on whole animals (flesh +
- shells) before dissection and mitochondrial preparation. Laboratory oysters were fed ad
- 144 libitum with a mix of two microalgae: Tisochrysis lutea formerly Isochrysis aff. galbana
- 145 clone Tahiti (T-Iso) and Chaetoceros gracilis, supplied in equal cell volumes. Cultures were
- produced in 300 L cylinders containing 1 µm filtered seawater enriched with Conway
- medium at 24±1°C, air-CO₂ (3%) and mix aerated, under continuous light. Microalgae were
- harvested at exponential growth phase (6–8 days).

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Reproductive activity

- 151 Qualitative analysis of gametogenic stage
- 152 A visceral mass slice (2-3 mm thick) was cut upstream of the adductor muscle, put in a
- histology cassette and transferred into a modified Davidson's fixative (Latendresse et al.
- 154 2002) for 48 h. Fixed tissues were dehydrated in ascending ethanol solutions, cleared with
- 155 Claral ® and embedded in paraffin wax. Five micrometer thick sections were cut, mounted on
- 156 glass slides, and stained with Harry's hematoxylin-eosin Y. Slides were examined under a
- light microscope to determine gametogenic stage according to the reproductive scale reported

- by Mann (1979): stage 0 (inactive), stage 1 (early gametogenesis), stage 2 (late
- gametogenesis) and stage 3 (ripe).

- 161 Quantitative analysis of gonad occupation area
- Percentage of gonad occupation area was determined on each histological section as described
- by Enríquez-Díaz et al. (2009). Briefly, slides were scanned with a digital scanner. Gonad
- area was measured by using image analysis software (Imaq Vision Builder, National
- 165 Instruments Corp., Austin, Texas, USA).

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Mitochondrial analysis

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- 169 Mitochondrial analyses were performed on mitochondria isolated from gills. For each group
- 170 (field and laboratory), six pools containing gills of five individuals were used (each pool
- 171 contained 2.5±0.1 g of gill tissue).

172

- 173 Isolation of mitochondria
- 174 Procedures for mitochondrial isolation are detailed in Dudognon et al. (2013). Oxygen
- 175 consumption was measured immediately on the fresh mitochondrial preparations.
- 176 A subsample of mitochondrial pellets was stored at -80°C for subsequent assays of enzymatic
- activities, cytochrome concentrations and lipid extraction.

- 179 Measurement of oxygen consumption
- 180 Mitochondrial oxygen consumption was measured polarographically using a water-jacketed
- 181 O₂ monitoring system (Qubit System, Kingston, Ontario, Canada). Temperature was
- 182 controlled at 10°C by a circulating refrigerated water bath. For each assay, around 0.8 mg of
- mitochondrial protein (40 µl mitochondrial preparation) was added to 0.4 ml reaction buffer
- 184 containing 400 mM sucrose, 30 mM HEPES, 90 mM KCl, 10 mM KH₂PO₄, 50 mM taurine
- and 50 mM β -alanine, pH 7.5. On the day of the experiment, 0.5% BSA was added to the
- assay medium. For measurement of maximal oxidative capacities, glutamate (40 mM) or
- succinate (20 mM) was added to fuel oxygen consumption through complex I or II,
- respectively. The maximal respiration rate (state 3) was obtained after addition of ADP to a
- 189 final concentration of 600 µM. Preliminary experiments showed that glutamate alone
- stimulates respiration through complex I after addition of ADP (Dudognon et al. 2013). Non-
- 191 phosphorylating oxygen consumption (state 4) rate was measured after ADP depletion. Each
- measurement was performed in simultaneous triplicates using three polarographic chambers.

193 RCR (respiratory control ratio) was defined as the ratio between state 3 and state 4. Oxidative

phosphorylation efficiency (relation between ADP added and oxygen consumption) was

calculated as ADP/O ratio according to Estabrook (1967).

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- 197 Enzymatic activities
- 198 Cytochrome c oxidase was measured in mitochondrial preparations and citrate synthase (CS)
- activity was measured in gill extracts according to Dudognon et al. (2013) and Dudognon et
- 200 al. (2014), respectively.

201

- 202 Cytochrome concentrations
- 203 Cytochromes a, b, c_1 , and c concentrations were quantified by difference spectra according to
- Leary et al. (2003) except that mitochondrial suspensions (around 0.8 mg of mitochondrial
- protein) were diluted in phosphate buffer (NaH₂PO₄ and Na₂HPO₄ 50 mmol pH 7.8 at 25°C)
- and Triton-X was not used. Accordingly, the electron transport chain complex was reduced by
- adding a few grains of sodium dithionite and incubated for 15 min at room temperature. The
- 208 reduced samples were read against air-oxidized samples between 400 and 630 nm. We used
- 209 the solution to the simultaneous equations required to assess individual cytochrome
- 210 concentrations (Williams Jr. 1964).

211

- 212 Protein concentration
- 213 Aliquots of 20 µl of mitochondria preparations were suspended in reaction buffer without
- BSA and centrifuged for 10 min at 12,000 g at 4°C. The supernatant was discarded and the
- 215 pellet re-suspended, washed and centrifuged twice again to remove the BSA. Pellets were
- 216 maintained in 0.5 ml of ultra-pure water and frozen at -80°C until protein analysis. The
- 217 protein concentration was determined with the RC DC Protein Assay Kit (BIORAD) using
- BSA as standard.

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Membrane lipid analysis

- The membrane lipids of mitochondrial suspensions were extracted according to the method of
- Folch et al. (1957). Before lipid extraction, the aliquot of mitochondrial preparation was
- resuspended in reaction buffer minus BSA and centrifuged at 9000 g at room temperature for
- 224 10 min. The supernatant was discarded and the pellet resuspended, washed in reaction buffer
- 225 and centrifuged again two times. The final extract was stored at -80°C under nitrogen after
- adding 0.01% w/v butylated hydroxytoluene (BHT, antioxidant).

- 228 Isolation of membrane lipids on silica gel microcolumn
- 229 Lipid extract was evaporated to dryness under nitrogen, recovered with three washings of
- 230 500 μl of chloroform/methanol (98:2, v/v) and deposited at the top of a silica gel micro-
- column (30 mm x 5 mm i.d.), packed with Kieselgel 60 (70–230 mesh, Merck, Darmstadt,
- Germany) previously heated at 450°C and deactivated with 6 % H₂O (Marty et al. 1992).
- Neutral lipids were eluted with 10 ml of chloroform:methanol (98:2, v/v). The polar lipid
- fraction (membrane lipids) was recovered with 15 ml methanol, and C23:0 solution was
- added as internal standard for further fatty acid analysis.

- 237 Fatty acid analysis of membrane lipids
- The polar lipid fraction was evaporated to dryness under nitrogen and transesterified for 10
- 239 min at 100°C after adding 1 ml of methanol/BF₃ (10%). After cooling, 1 ml of ultra-pure
- 240 water and 1 ml of hexane were added. Tubes were mixed thoroughly for 2 min and
- 241 centrifuged at 1,000 g for 10 min. The organic phase containing fatty acid methyl esters
- 242 (FAME) was washed three times with 1 mL of water. Fatty acid methyl esters were quantified
- by gas chromatography and identified by comparing their retention times with those of a
- standard mixture containing 37 FAME (SUPELCO/Sigma-Aldrich, St-Quentin Fallavier,
- France), and other lab-made standard mixtures of marine bivalve. A total of 40 FA were
- 246 quantified and expressed as the molar percentage of the total FA content.

247

- 248 Separation of membrane lipid classes
- 249 Phospholipid classes were analyzed by high-performance thin-layer chromatography
- 250 (HPTLC) as described in details in Dudognon et al. (2014). Mass of each identified
- 251 phospholipid class was determined by comparison with standards of known amounts. The
- 252 technique used did not separate phosphatidylinositol (PI) and
- ceramideaminoethylphosphonate (CAEP) (unpublished preliminary results), and proportions
- of these phospholipids were presented as the sum PI + CAEP (CAEP being predominant over
- 255 PI in oyster gill mitochondria (data not shown)). The sum of PI+CAEP was quantified using
- sphingomyelin as standard because sphingomyelin has a similar structure to CAEP although
- is absent in bivalves (Le Grand et al. 2011). Results were expressed as mass% of total
- 258 phospholipids.

- 260 Statistical analysis
- ANOVA was performed if normality and homogeneity of variances were respected. ANOVA
- was followed by post hoc Fisher's least significant difference (LSD) test. For data that did not

follow normality or meet homoscedasticity, non-parametric Kruskal-Wallis analysis was performed. For each test, the significance threshold was P<0.05. Percentage data were transformed (arcsin of the square root) before ANOVA but are presented in figures and tables as untransformed percentage values. All analyses were performed with the Statgraphics software, version Plus 5.1. (Manugistics, Inc, Dallas, USA). Results are expressed as mean \pm standard error (SE).

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Results

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- 273 Oyster weight
- 274 Initial oyster weight in trays was 41.7±2.6 g (T0) and increased significantly after 4 weeks
- 275 under both field and laboratory conditions (Figure 1). The two groups of oysters did not differ
- at the end of conditioning in June (mean values of 56.2±1.7 g and 61.0±2.3 g for field and
- laboratory oysters, respectively). No mortality occurred during the experiment.

278

- 279 Gonad maturation stages and gonad coverage area
- Oysters sampled in May were in late gametogenesis (stage 2), whereas those sampled in June
- were mostly (about 80%) reproductively ripe (stage 3) (data not shown). Similarly, percentage
- of gonad coverage area increased significantly (P<0.01) from initial values in May (43.4±3.5)
- to final values of 65.1±3.1% and 68.8±2.1% in oysters from laboratory and field conditions,
- 284 respectively. There was no difference in gonad occupation area between laboratory and field
- oysters (Figure 1).

- 287 Mitochondrial oxidative capacities
- Oxidative capacities of mitochondria isolated from oyster gills were assessed using glutamate
- or succinate as substrate. Overall, the capacities of mitochondria isolated from field oysters at
- 290 Tf were similar to those of oysters at T0; both were systematically higher than those of
- 291 laboratory conditioned oysters at Tf (Fig. 2). Maximal state 3 rates of glutamate and
- succinate oxidation were significantly lower in laboratory oysters than in field oysters for
- both substrates (around 40% lower) (Figure 2A). The same pattern was observed with the
- 294 non-phosphorylating rate (state 4) (Figure 2B). State 4 was significantly higher with succinate
- than with glutamate (Figure 2B, *P*<0.05).

- As a consequence of these modifications of state 3 and state 4 rates, RCR with glutamate was
- lower in laboratory than in field oysters at the end of the experiment (- 25%) (Table 3). On the
- other hand, RCR with succinate did not differ at Tf between field and laboratory oysters.
- 299 ADP/O did not differ between field and laboratory oysters for glutamate and succinate, but
- 300 values were significantly higher for field oysters at T0 compared to Tf for both substrates.
- 301 RCR and ADP/O were always significantly higher with glutamate than with succinate as
- 302 substrate (*P*<0.05).

- 304 Mitochondrial enzyme activities and protein composition
- 305 Cytochrome c oxidase and citrate synthase activity
- 306 CCO activity, expressed in U.mg⁻¹ mitochondrial protein, was significantly lower in
- laboratory conditioned oysters (-33%), than in field oysters sampled at T0 and Tf (Table 4).
- 308 As cyt a is located in complex IV (CCO) of the respiratory chain, activity of CCO was also
- 309 expressed per nmole of cytochrome a to reflect its catalytic activity. In that case, CCO
- activity did not differ between groups.
- 311 CS activity, measured on isolated gill tissue and expressed in U mg⁻¹ gill tissue, did not differ
- between the three groups of oysters.

313

- 314 Mitochondrial cytochrome concentrations
- 315 Cytochrome b concentration did not differ between field oysters at T0 and Tf; both groups
- 316 had significantly higher values than laboratory conditioned oysters (-24%) (Table 5).
- 317 Cytochrome a, c_1 and c concentrations did not differ between the 3 groups. There was no
- 318 difference between oyster groups when cytochrome quantities were expressed relative to
- 319 cytochrome c_1 . Similarly, there was no difference in the ratio $(b + c_1)/a$ between the three
- groups of oysters.

- 322 Fatty acid composition of phospholipids from gill mitochondria
- Four weeks under field and laboratory conditions significantly changed the FA composition
- of membrane phospholipids of oyster gill mitochondria compared to their initial status (T0)
- 325 (Table 1). More importantly, FA composition differed between rearing conditions at Tf.
- 326 The proportions of 20:5n-3 decreased throughout the experiment but was higher in
- 327 mitochondria of field-conditioned oysters than in laboratory-conditioned oysters at Tf. The
- proportions of 22:6n-3 and 20:4n-6 were also higher at T0, but did not differ between the two
- 329 groups of oysters at Tf. The proportion of 22:5n-6 was 0.4% at the start of the experiment and

- did not change in oysters after 4 weeks in the field. However, it rose by more than 5 fold
- 331 (2.1%) in oysters after laboratory conditioning.
- 332 After 4 weeks conditioning, total NMI (non-methylene interrupted) FA were more abundant
- in laboratory-conditioned oysters than in field oysters. This difference mainly reflected higher
- levels of 22:2NMI(7,15) in laboratory-conditioned oysters.
- The proportion of 18:1n-7 (NMI FA precursor) reached 5.6% in oysters after 4 weeks in the
- laboratory while this FA was lower (3.4%) in oysters from the field at the same time. There
- was no difference in the proportion of the monounsaturated FA (MUFA) 16:1n-7 between the
- three groups of oysters. The level of 18:1n-9, 20:1n-9 and 20:1n-7 reached similar levels
- between laboratory and field oysters after 4 weeks conditioning.
- 340 The proportion of total saturated FA (SFA), monounsaturated FA (MUFA) and
- polyunsaturated FA (PUFA) changed markedly throughout the experiment. SFA were higher
- in mitochondria of oysters from the field than from the laboratory at Tf. At Tf, MUFA were
- 343 higher in mitochondria of laboratory oysters as compared to field oysters. On the other hand,
- total PUFA did not differ between the two groups sampled at Tf. Within PUFA, mitochondria
- of oysters from the laboratory showed the highest proportion of n-6 PUFA but the lowest
- proportion of n-3 PUFA at the end of the experiment.
- 348 Phospholipid classes of mitochondrial membranes
- Phosphatidylcholine (PC) and phosphatidylethanolamine (PE) were the main phospholipid
- 350 classes in mitochondrial membranes of oysters accounting for 64-66.8% of total
- 351 phospholipids (Table 2). Phosphatidylinositol (PI) and ceramideaminoethylphosphonate
- 352 (CAEP) together represented around 20% and phosphatidylserine (PS) accounted for about
- 353 8% of the total phospholipids. The phospholipid-to-protein ratio (i.e. total phospholipid
- content expressed per mg of mitochondrial proteins), as well as proportions of PE, PC and PS
- 355 did not differ among treatments at Tf. The proportions of PI + CAEP increased slightly but
- 356 significantly after 4 weeks of experimentation and oysters from the laboratory showed a
- 357 higher proportion of PI + CAEP than oysters from the field. The proportions of CL changed
- 358 significantly between rearing conditions and laboratory oysters showed lower levels in
- mitochondria than field oysters at Tf (by about -25%).
- 360 PC/PE ratio decreased after 4 weeks conditioning and reached significantly lower values in
- laboratory as compared to field oysters, whereas the ratio PE/CL was higher in mitochondria
- of oysters from the laboratory than field oysters.

Discussion

The similar increase in weight and gonadal occupation between oysters grown in the laboratory and those left in a mesh bag in their natural environment confirmed the suitability of laboratory conditioning for oyster growth and gonad development. Most feeding studies on bivalves use such overall criteria, while few investigate cellular or sub-cellular responses. The present results showed that rearing oysters in laboratory-controlled conditions rather than under fluctuating field conditions decreased the capacities of gill mitochondria, while overall growth and gonad development did not differ.

Major differences of mitochondrial capacities were observed between the groups. Phosphorylating rates of oxygen consumption (state 3) were almost two fold lower in gill mitochondria from laboratory-conditioned oysters. This was true both when isolated mitochondria were fueled with glutamate through complex I or with succinate through complex II. Non-phosphorylating oxygen consumption (state 4) which partly reflects proton leak (St. Pierre et al. 2000) was also consistently lower in laboratory-conditioned oysters.

Differences in mitochondrial oxidative capacities can reflect changes in concentrations or relative levels of electron transport chain complexes (Guderley et al. 2005). Cytochrome *b* was 24% less abundant in gill mitochondria from oysters reared in laboratory tanks than in the field. A decrease in cytochrome *b*, a component of complexes II and III of the respiratory chain, may be implicated in the diminished mitochondrial oxidative capacities in laboratory oysters. A concomitant decrease in mitochondrial respiration and in CCO activity is observed in response to diet in rats (Yamaoka et al. 1988), temperature in fish (Blier and Lemieux 2001); (Kraffe et al. 2007), and hypoxic hibernation in frogs (Boutilier and St-Pierre 2002). CCO is the last complex of the respiratory chain and can control, at least in part, rates of electron transport through the electron transport chain (Groen et al. 1982). Therefore, diminished CCO activity could be one cause of reduced mitochondrial oxidative capacity in gills of laboratory conditioned oysters.

Phospholipid classes and their FA composition influence the activity of membrane-bound proteins, such as respiratory chain complexes in mitochondria (Robinson 1993; Stuart et al. 1998; Frick et al. 2010). Phospholipid proportions in mitochondrial membranes differed between laboratory and field oysters. Among these differences, the lower levels of CL in gill mitochondria of laboratory oysters could explain the lower activity of CCO. Indeed, CL is specifically located in the inner membrane of mitochondria and its association with respiratory chain enzymes is of major importance for regulation of their activity and the generation of the inner membrane potential (Schlame et al. 2000; Haines and Dencher 2002;

Sedlák et al. 2006; Zhou et al. 2011; Böttinger et al. 2012; Kagan and Epand, 2014). A reduced level of CL is associated with decreased mitochondrial activity in mammals (Paradies et al. 1997), snails (Stuart et al. 1998) and fishes (Frick et al. 2010). CL deficiency also induces destabilization, decreased activity and loss of complex III (Schlame et al. 2000; Lange et al. 2001; Petrosillo et al. 2003; Böttinger et al. 2012), potentially explaining the lower cytochrome *b* content in laboratory conditioned oysters

The FA composition of phospholipids influence the activity of membrane-bound proteins, either through overall or micro-environment localized effects (Robinson 1993; Phillips et al. 2009; Hulbert and Else 1999). Proportions of SFA were higher and those of MUFA lower in gill mitochondria from field oysters than laboratory conditioned oysters. These overall differences may well have changed mitochondrial oxygen uptake and CCO activity. The results of a partner study indicate that maintenance of overall aspects of membrane FA composition (total SFA, MUFA and PUFA) despite marked changes in the levels of individual FA may be key for the maintenance of mitochondrial function in oyster gills (Dudognon et al., 2014) as reported in mammals (Astorg and Chevalier 1991; Lemieux et al. 2008; Abbott et al. 2012) and fish (Guderley et al. 2008; Martin et al. 2013).

In the experimental tank, feeding oysters ad libitum with a mixture of the two microalgae T.Isochrysis lutea, and Chaetoceros gracilis permited a balanced supply of essential FA as 20:4n-6, 20:5n-3 and 22:6n-3 for oysters (Delaunay et al., 1993; Soudant et al. 1996a; Delaporte et al., 2005; Rico-Villa et al. 2006; Marshall et al., 2010; Dudognon et al. 2014). Laboratory and field conditioned oysters had indeed similar levels of the essential FA in gill mitochondria, with the exception of 20:5n-3, confirming that the mixed diet T-Iso + C. gracilis provided sufficient essential FA. Nevertheless, differences of some particular FA (i.e. 22:5n-6 and NMI FA) between laboratory and field oysters may be important for the regulation of mitochondrial membrane protein activities (Zhou et al. 2011; Khairallah et al. 2012). Whereas 22:5n-6 appears to be preferentially incorporated from the microalgae T. Iso (Soudant et al., 1996a; Soudant et al., 1996c; Dudognon et al., 2014), NMI FA are the only long chain PUFA that bivalves synthesize de novo (Zhukova 1991). Both 22:5n-6 and NMI FA have been identified as potentially significant for the physiology of bivalve larvae and the structure and fluidity of membranes in female gametes (Soudant et al. 1996a; Soudant et al., 1996c; Pernet et al. 2005; Mike et al. 2008), for membrane functions of bivalve hemocytes (Le Grand 2011, 2013; Dudognon et al. 2014) as well as for peroxidation resistance and activities of gills and gill mitochondria of bivalves (Dudognon et al. 2014; Munro and Blier 2012; Barnathan 2009; Delaporte et al. 2005; Kraffe et al. 2004).

Changes in several characteristics, ranging from cytochrome contents to phospholipid proportions and FA characteristics, could explain the loss of capacity of gill mitochondria from oysters conditioned in the laboratory. A loss of capacity after laboratory conditioning also occurred in mitochondria from various tissues of the scallop, Argopecten purpuratus (Guderley et al. 2011). One hypothesis questions whether the bi-specific algal diet contains what oysters need to maintain mitochondrial activity. Beside their content in FA, carbohydrate or amino acids, microalgal species also vary significantly in contents of vitamins, minerals and trace elements (Brown 2002). In mammals, these compounds are critical cofactors that support mitochondrial functions. Deficiencies in some minerals and vitamins can reduce activities of electron transport complexes and cardiolipin content in mammals (Ames et al. 2005; Oliveros et al. 2007). As growth rates and gametogenesis did not differ between laboratory and field oysters, the bispecific algal diet should have provided enough energy for somatic growth and gametes maturation, but may have been lacking materials for mitochondrial activity. Finally, the continuous ad libitum feeding we used in the laboratory can be unfavourable for intertidal bivalves such as oysters (Utting et Spencer 1991; Racotta et al. 1998; Hurtado et al. 2009), and may have decreased mitochondrial performance.

Nutrient supply may not be the only environmental factor that could have impacted mitochondrial functions and compositions in oysters. Laboratory conditioned oysters were constantly immersed in well-aerated filtered seawater whereas field oysters had to cope with marked variations in physical, chemical and biological parameters (e.g. temperature, oxygenation and feeding activity during emersion and immersion cycles), linked to tidal cycles. Modifications of environmental conditions, including salinity, hypoxia and temperature, change the structure and function of bivalve mitochondria (Glémet and Ballantyne 1995; Gillis and Ballantyne 1999; Guderley 2004; Sussarellu et al. 2013; Dudognon et al. 2013). Our observations suggest that variable environmental conditions in the field may have stressed changes in specific lipid constituents of mitochondrial membranes and modified mitochondrial capacities.

Conclusion

Standardized conditions and *ad libitum* feeding of oysters with a mixed plankton diet do not fully mimic field conditions. Changes of specific FA and phospholipid classes in mitochondrial membranes may explain the lower mitochondrial capacities in laboratory-conditioned oysters. Field conditions impose fluctuations and differences in abiotic and biotic

factors that may have led structural and functional properties of gill mitochondria to differ from those of laboratory conditioned oysters. Nonetheless, overall "fitness" was not affected as oysters showed similar growth and gonadal development in the field and laboratory. Nevertheless, the changed mitochondrial capacities suggest that cellular processes in oysters held in constant laboratory conditions are reduced compared to those of oysters living in the intertidal zone. This could impair other physiological and cellular processes. It may be for example questionable why oysters fed *ad libitum* and probably having access to more food and for a longer period of time did not grow better in the laboratory.

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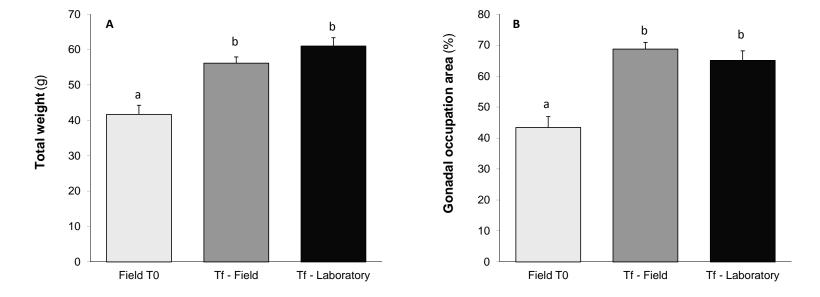


Figure 1: Total weight (expressed as g) (A) and gonad occupation area (expressed as %) (B) of oysters (n = 30) grown on their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf - Field, Tf - Laboratory). Values are mean \pm SE. Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups.

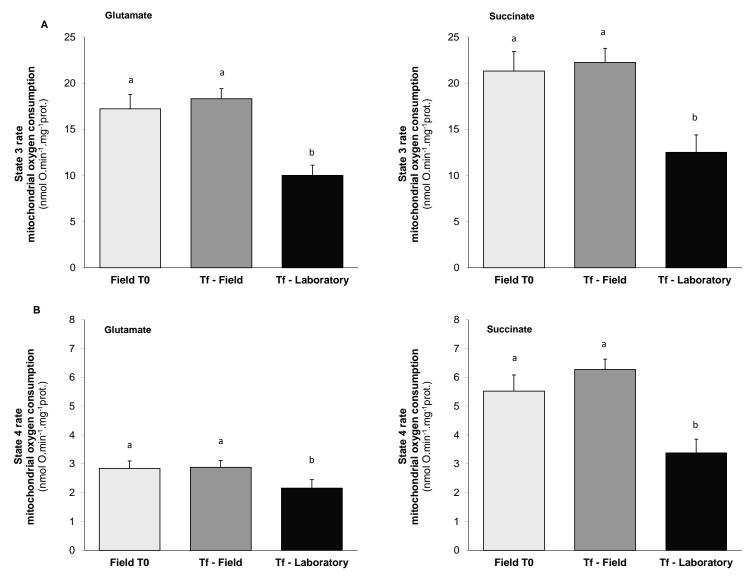


Figure 2: Rates of oxidative phosphorylation (state 3) (A) and non-phosphorylating oxygen consumption (state 4) (B) with glutamate 40 mM or succinate 20 mM on mitochondria isolated from oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf. Field, Tf. Laboratory). Assay temperature was 10°C. Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean ± SE, n = 6 pools of 5 oysters).

Table 1: Fatty acid composition of gill mitochondria phospholipids of oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf – Field, Tf – Laboratory). The results are expressed as mol percentages of total fatty acids. Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean \pm SE, n = 6 pools of 5 oysters).

	Field T0		Tf - Field		Tf - Laboratory		
_	Mean	SE	Mean	SE	Mean	SE	
14:0	1.4 a	0.0	1.5 a	0.1	1.8 b	0.1	
16:0	10.3 ^a	0.1	12.2 b	0.4	11.1 °	0.1	
18:0	5.7 ^a	0.1	9.5 ^b	0.6	5.5 ^c	0.0	
16:1n-7	2.2	0.1	1.9	0.1	2.1	0.1	
18:1n-9	1.3 ^a	0.0	1.6 b	0.1	2.0 b	0.1	
18:1n-7	4.2^{a}	0.1	3.4 b	0.1	5.6 °	0.1	
20:1n-11	1.5 a	0.0	2.1 b	0.0	2.6 °	0.1	
20:1n-9	1.7 a	0.0	1.4 ^b	0.0	1.4 ^b	0.1	
20:1n-7	5.4 ^a	0.0	4.7 ^b	0.1	5.0 b	0.0	
18:2n-6	0.6^{-a}	0.0	0.8 b	0.0	1.2 °	0.1	
18:4n-3	1.3 ^a	0.1	2.1 b	0.3	1.0 ^c	0.0	
20:4n-6	5.6 ^a	0.1	4.6 b	0.1	4.4 ^b	0.1	
20:5n-3	19.4 ^a	0.2	16.4 ^b	0.3	14.9 ^c	0.5	
22:2NMI(7,13)	1.9	0.1	1.9	0.1	2.2	0.1	
22:2NMI(7,15)	5.9 ^a	0.2	5.5 ^b	0.1	7.4 ^c	0.1	
22:3NMI(7,13,16)	1.2 a	0.0	1.0 b	0.1	0.9 ^b	0.0	
22:4n-6	0.6^{-a}	0.0	0.5 b	0.0	0.5 ^b	0.0	
22:5n-6	0.4^{-a}	0.0	0.4^{-a}	0.0	2.1 b	0.2	
22:5n-3	1.7 ^a	0.0	1.7 ^a	0.0	1.0 b	0.0	
22:6n-3	20.9^{-a}	0.3	18.7 ^b	0.6	20.1 ^{ab}	0.3	
Others*	3.8 ^a	0.1	4.9 b	0.2	3.7 ^a	0.1	
Total BR	0.5 a	0.0	0.5 a	0.0	0.3 b	0.0	
Total SFA	17.6 ^a	0.2	23.4 ^b	1.0	18.5 ^c	0.2	
Total MUFA	16.5 a	0.1	15.4 b	0.3	19.1 ^c	0.1	
Total n-9	3.2 a	0.1	3.1 ^a	0.2	3.7 b	0.2	
Total n-7	11.7 ^a	0.1	10.0 b	0.2	12.7 ^c	0.1	
Total PUFA	65.4 ^a	0.3	60.6 b	0.8	62.1 ^b	0.3	
Total n-4	0.3^{a}	0.0	0.3 a	0.0	0.2 b	0.0	
Total n-6	8.4 ^a	0.1	7.7 b	0.2	9.5 ^c	0.3	
Total n-3	47.3 ^a	0.3	43.7 b	0.7	41.1 °	0.5	
Total NMI	9.4 ^a	0.2	9.0 ^a	0.1	11.2 b	0.2	

*Others: total of 21 fatty acids detectable (iso17:0, ant17:0, 20:0, 22:0, 24:0, 18:1n-11, 22:1n-9, 16:2n-7, 16:2n-4, 16:3n-6, 16:4n-3, 18:2n-4, 18:3n-6, 18:3n-3, 20:2NMI(5,11), 20:2NMI(5,13), 18:5n-3, 20:2n-6, 20:3n-6, 20:4n-3, 21:5n-3), none of which were more than 1%.

BR: branched FA; SFA: saturated FA; MUFA: monounsaturated FA; PUFA: polyunsaturated FA; NMI: non-methylene-interrupted FA

Table 2: Content of total phospholipids and proportion of phospholipid classes in gill mitochondria of oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf – Field, Tf – Laboratory). The results are expressed as mol percentages of total fatty acids. Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean \pm SE, n = 6 pools of 5 oysters).

	Field T0		Tf - F	Tf - Field		Tf - Laboratory			
_	Mean		SE	Mean		SE	Mean		SE
PE	29.8		0.3	30.3		0.2	30.6		0.3
PI + CAEP	19.3	a	0.2	20.1	b	0.2	21.9	c	0.4
CL	6.8	a	0.3	6.4	a	0.2	5.0	b	0.2
PC	37.0	a	0.5	35.2	ab	0.6	33.4	b	0.9
PS	7.1		0.3	8.0		0.7	9.0		1.0
PC/PE	1.24	a	0.02	1.16	b	0.03	1.09	c	0.02
PE/CL	4.4	a	0.2	4.8	a	0.2	6.1	b	0.2
Total phospholipid content (μg.mg ⁻¹ prot.)	471.3		39.2	450.4		7.9	394.7		32.5

PE: phosphatidylethanolamine; PI: phosphatidylinositol; CL: cardiolipin; CAEP: ceramide aminoethylphosphonate; PC: phosphatidylcholine; PS: phosphatidylserine

Table 3: Respiratory control ratio (RCR) and oxidative phosphorylation efficiency (ADP/O), with glutamate 40 mM or succinate 20 mM as substrate and ADP 0.6 mM of oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf – Field, Tf – Laboratory). Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean \pm SE, n = 6 pools of 5 oysters).

	Field T0		Tf - Fie	ld	Tf - Laboratory		
_	Mean	SE	Mean	SE	Mean SE		
RCR glutamate	6.2 ^a 3.9 ^a	0.5	6.4 a	0.2	4.8 b 0.4		
RCR succinate		0.1	3.6 b	0.1	3.3 b 0.2		
ADP/O glutamate	2.6 a	0.2	2.0 b	0.0	2.3 ab 0.3		
ADP/O succinate	1.9 a	0.1	1.6 b	0.1	1.8 ab 0.2		

Table 4: CCO activity, expressed per mg of mitochondrial proteins or per nmol of cytochrome a, and CS activity, expressed in per mg of gill tissue of oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf – Field, Tf – Laboratory). Assay temperature was 25°C. Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean \pm SE, n = 6 pools of 5 oysters).

	Field T0		Tf - Fi	ield	Tf - Laboratory		
	Mean	SE	Mean	SE	Mean	SE	
CCO activity							
(U.mg ⁻¹ prot.)	0.3 a	0.0	0.3 a	0.0	0.2^{-b}	0.0	
$(U.nmol^{-1}cyt a)$	4.4	0.9	4.5	0.7	3.2	0.4	
CS activity (U.mg ⁻¹ tissue)	4.4	0.4	4.7	0.3	5.0	0.2	

Table 5: Concentrations of mitochondrial cytochromes a, b, c_1 and c, expressed as nmol of cytochrome per mg of mitochondrial proteins, relative cytochrome ratios related to cytochrome c_1 and ratio $(b + c_1)/a$ of oysters grown in their natural environment or in the laboratory, and sampled before (Field T0) and after 4 weeks of conditioning (Tf – Field, Tf – Laboratory).

Different superscript letters indicate values that differ significantly (p<0.05) between oyster groups (mean \pm SE, n = 6 pools of 5 oysters).

	Field T0		Tf - Field		Tf - Laboratory	
	Mean	SE	Mean	SE	Mean	SE
Cytochrome concentrations (nmol.mg ⁻¹ prot.)						
a	0.07	0.01	0.06	0.01	0.05	0.01
b	0.21 a	0.02	0.21 a	0.02	0.16 b	0.01
c_1	0.12	0.01	0.13	0.01	0.12	0.02
c	0.06	0.01	0.08	0.01	0.06	0.01
Total	0.45	0.03	0.48	0.04	0.38	0.03
Relative ratios						
a/c_1	0.60	0.11	0.44	0.04	0.46	0.08
b/c_1	1.85	0.22	1.62	0.11	1.48	0.15
c/c_1	0.55	0.07	0.57	0.04	0.48	0.07
$(b+c_1)/a$	5.2	0.80	6.1	0.60	6.1	1.10