

**Regular** paper

# Tissue variation of mitochondrial oxidative phosphorylation efficiency in cold-acclimated ducklings\*

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We investigated the oxidative phosphorylation efficiency of liver and gastrocnemius muscle mitochondria in thermoneutral and cold-acclimated ducklings. The yield of oxidative phosphorylation was lower in muscle than in liver mitochondria, a difference that was associated with a higher proton conductance in muscle mitochondria. Cold exposure did not affect oxidative phosphorylation efficiency or basal proton leak in mitochondria. We conclude that the basal proton conductance of mitochondria may regulate mitochondrial oxidative phosphorylation efficiency, but is not an important contributor to thermogenic processes in cold-acclimated ducklings.

Keywords: thermogenesis, liver, skeletal muscle, mitochondrial oxidative phosphorylation, proton conductance

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## INTRODUCTION

When chronically exposed to cold, the survival of some species of birds, including chickens, ducklings, and penguins, depends on their ability to develop non-shivering thermogenesis processes (review in Duchamp et al., 1999). In the absence of both mammalian thermogenic brown adipose tissue and UCP1, both skeletal muscle and liver appear as contributors to the non-shivering thermogenesis (Duchamp & Barre, 1993; Bedu et al., 2001). Two separate mechanisms have been proposed to explain how skeletal muscle may contribute to nonshivering thermogenesis. The first one is cytoplasmic and would generate heat from the futile cycling of calcium across the reticulum sarcoplasmic membrane (Dumonteil et al., 1995), consuming ATP and stimulating mitochondrial respiration. The second one is mitochondrial and involves loose coupling of oxidative phosphorylation (Skulachev & Maslov, 1960; Duchamp et al., 1992; Roussel et al., 1998), whereby a regulated leak of protons would partially dissipate the protonmotive force, producing heat instead of ATP. New evidence for this lies in the recent identification in the skeletal muscle of a functional avian uncoupling protein homologous to UCP1 (Raimbault et al., 2001; Vianna et al., 2001; Talbot et al., 2003), whose mRNA expression and activity are up-regulated during cold acclimation in birds (Raimbault et al., 2001; Toyomizu et al., 2002; Collin et al., 2003; Talbot et *al.*, 2004; Rey *et al.*, 2010; Teulier *et al.*, 2010). In liver, heat generation is ill defined and may be a by-product of the increased metabolic activity of the organ rather than

involve a specific thermogenic mechanism. Indeed, cold acclimation has been reported to stimulate hepatic energy-demanding pathways such as lipogenesis and glucagon-enhanced gluconeogenesis (Bedu et al., 2001; 2002).

As illustrated above, there are two fundamental ways of generating more heat for thermogenesis purpose, one is to increase the cellular ATP utilization and the other is to reduce the efficiency of mitochondrial ATP synthesis. A central requirement of these two mechanisms is the mitochondrial oxidative phosphorylation process, as mitochondria either supply the cellular energy-consuming mechanisms with ATP or undergo loose coupling, generating more heat per ATP synthesized. In the context of thermal physiology, the coupling efficiency of mitochondrial energy transduction is therefore of interest, because it describes how much energy mitochondria use for each ATP molecule they produce in the course of substrate oxidation and oxidative phosphorylation. Several studies have previously reported values of mitochondrial ADP/O ratio in muscle and liver from cold-acclimated birds, and found either a decrease or no changes when compared to the values found for control birds (Roussel et al., 1998; Chaînier et al., 2000; Toyomizu et al., 2002). However, in those studies the yield of ATP synthesis was calculated from the rates of oxygen consumption in mitochondria under fully phosphorylating active state (state 3, high concentration of ADP). In order to generate data with more physiological relevance, we initiated the study reported here. By measuring oxygen consumption and ATP synthesis at maximal and submaximal oxidative phosphorylation rates, we re-investigated the effect of cold acclimation on the energetic efficiency of skeletal muscle and liver mitochondria from ducklings. We found that the ATP yield of mitochondria was not affected by cold exposure but greatly differed between the muscle and liver mitochondria. This tissue variation was associated with a difference in the membrane proton conductivity of mitochondria.

#### MATERIALS AND METHODS

Animals. Male muscovy ducklings (Cairina moschata L, pedigree R31, Institut National de la Recherche Agronomique, France) were obtained from a commercial

e-mail: damroussel@yahoo.fr This work is related to that presented in the poster and oral forms at the 16th European Bioenergetics Conference (Warsaw, 2010); abstract in Biochim Biophys Acta, 1797 (Suppl): 62 (2010).

Mitochondria isolation. Muscle and liver samples were rapidly dissected, weighed, placed in ice-cold isolation buffer and homogenized with a Potter-Elvehiem homogenizer. Liver mitochondria were isolated in a buffer containing 250 mM sucrose, 2 mM KH<sub>2</sub>PO<sub>4</sub>, 1 mM EGTA, 20 mM Tris/HCl, pH 7.3 at 4°C. The liver homogenate was centrifuged at  $700 \times g$  for 10 min. The resulting supernatant was centrifuged at  $1000 \times g$  for 10 min, filtered through cheesecloth and re-centrifuged at 8700×g for 10 min to pellet mitochondria. Muscle mitochondria were isolated from the internal part of the gastrocnemius muscle in a buffer containing 100 mM sucrose, 50 mM KCl, 5 mM EGTA, 50 mM Tris/HCl, pH 7.4 at 4°C. The muscle homogenate was centrifuged at  $800 \times g$  for 10 min. The pellet containing intermyofibrillar mitochondria was suspended in 40 ml of isolation buffer and then treated with nagarse (1 mg/g muscle wet weight) for 5 min in an ice bath. The mixture was diluted 1:2, homogenized and then centrifuged at  $1000 \times g$ for 10 min. The supernatant was filtered through cheesecloth and centrifuged at  $8700 \times g$  for 10 min. Finally, liver and muscle mitochondrial pellets were washed twice by suspension in their respective isolation buffer and centrifugation at  $8700 \times g$  for 10 min. The protein concentration of mitochondrial suspensions was determined by the biuret method with bovine serum albumin as a standard.

Mitochondrial respiration and ATP synthesis. Respiration was measured in a glass cell of 1.5 ml volume fitted with a Clark oxygen electrode (Rank Brothers Ltd, France), at 38°C and calibrated with air-saturated respiratory buffer (120 mM KCl, 5 mM KH<sub>2</sub>PO<sub>4</sub>, 1 mM EGTA, 2 mM MgCl<sub>2</sub>, 0.3% bovine serum albumin (w/v) and 3 mM Hepes, pH 7.4). Liver mitochondria (1 mg of protein/ml) and muscle mitochondria (0.3 mg of protein/ml) were incubated in the respiratory buffer supplemented with 20 mM glucose, 1.5 U/ml hexokinase, 5 µM rotenone and 5 mM succinate. Different steady states of phosphorylation were obtained by adding increasing amounts of ADP from 5 to 100 µM. The respiration rate was recorded for 2 min and four 300-µl aliquots of mitochondrial suspension were then withdrawn every 30 s and immediately quenched in perchloric acid solution (10% HClO<sub>4</sub>/25 mM EDTA). After centrifugation of the denatured protein  $(15000 \times g \text{ for } 5)$ min) and neutralization of the resulting supernatant with KOH (2 M KOH/0.3 M Mops), the ATP production was determined from the samples' glucose-6-phosphate content, which was measured enzymatically by spectrophotometry according to (Bergmeyer, 1974).

It is important to note that we made sure that the rates we measured were specific of the mitochondrial ATP synthase activity, by determining oxygen consumption and ATP synthesis rates in the presence of oligomycin (2  $\mu$ g/ml). Over the range of ADP concentrations used, an oligomycin-insensitive ATP synthesis activity was measurable at 100  $\mu$ M ADP in skeletal muscle

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mitochondria purified from both experimental groups  $(37 \pm 12 \text{ nmol ATP/min.mg of protein and } 35 \pm 3 \text{ nmol ATP/min} \cdot \text{mg of protein in control and cold-acclimated birds, respectively) and in liver mitochondria from cold-acclimated ducklings (5 \pm 1 \text{ nmol ATP/min} \cdot \text{mg of protein}). These values were taken into account to calculate the rate of mitochondrial ATP synthesis.$ 

Mitochondrial proton leak. Respiration and membrane potential were measured simultaneously using electrodes sensitive to oxygen and to the potential-dependent probe triphenyl-methyl-phosphonium (TPMP+), respectively. Liver mitochondria (1 mg of protein/ml) and muscle mitochondria (0.35 mg of protein/ml) were suspended in respiratory buffer supplemented with 65 ng/ ml nigericin, 1  $\mu$ g/ml oligomycin and 5  $\mu$ M rotenone. The TPMP electrode was calibrated by sequential 0.5 µM additions up to 2 µM TPMP+, and then succinate was added to start the reaction. Respiration and membrane potential were progressively inhibited through successive steady states induced by additions of malonate. Membrane potential was calculated as described in (Brand, 1995), assuming a TPMP binding correction of 0.42 and 0.35 (µl/mg of protein)<sup>-1</sup> for liver and skeletal muscle mitochondria, respectively (Rolfe et al., 1994).

**Statistical analysis.** The statistical significance of observed differences was assessed using two-way ANOVA. Differences between means were subsequently tested by Scheffe's test.

### **RESULTS AND DISCUSSION**

The relationships between the rates of ATP synthesis and oxygen consumption were linear in both liver and muscle mitochondria (Fig. 1A). The maximal ATP synthesis rate in muscle mitochondria (highest points to the right of the linear relationships) was significantly increased by 14% (P<0.01) in cold-acclimated ducklings compared to the control birds, while the corresponding rate of oxygen consumption was slightly increased but failed to reach statistical significance (+11%, P=0.06). In contrast, maximal rates of ATP synthesis and oxygen consumption were not affected by cold in liver mitochondria. Nevertheless, the relationship concerning the cold-acclimated group was superimposed on that of the thermoneutral group in both liver and skeletal muscle mitochondria (Fig. 1A). Hence, 4 weeks of cold exposure did not alter the amount of ATP synthesized from ADP per amount of oxygen consumed. This is better shown in Fig. 1B where the ATP/O ratio is plotted versus the oxygen consumption rate. Indeed, at any given rate of oxygen uptake, ATP/O was the same in the cold-acclimated and thermoneutral groups. The present observation that cold acclimation specifically increased the maximal capacity of skeletal muscle mitochondria to synthesize ATP with no alterations in the efficiency of oxidative phosphorylation reinforce previously published data (Roussel et al., 1998; Teulier et al., 2010).

Moreover, liver mitochondria displayed higher oxidative phosphorylation efficiency than skeletal muscle mitochondria (Fig. 1). Indeed, the linear relation in liver mitochondria was shifted to the left compared to muscle mitochondria, indicating that to produce a given amount of ATP, less oxygen was consumed by liver mitochondria (Fig. 1A). This remains true even though Fig. 1B shows that at maximal rates of ATP synthesis and oxygen consumption (the highest points to the right of the relationships), ATP/O was slightly higher in skeletal

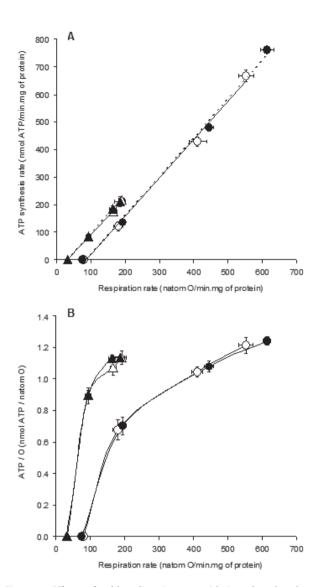


Figure 1. Effects of cold acclimation on oxidative phosphorylation efficiency

Liver (triangles) and skeletal muscle mitochondria (circles) from thermoneutral (open symbols) and cold-acclimated ducklings (filled symbols) were incubated in respiratory buffer supplemented with 20 mM glucose and 1.5 U/ml hexokinase, as described in Materials and Methods. Oxygen consumption and ATP synthesis rates were measured in the presence of increasing concentrations of ADP (0–100  $\mu$ M). Values are means  $\pm$  S.E.M. of four to six independent preparations.

muscle (+8%) than in liver mitochondria from both thermoneutral ( $1.21\pm0.05$  versus  $1.12\pm0.05$ , P=0.27) and cold-acclimated ducklings ( $1.24\pm0.02$  versus  $1.14\pm0.01$ , P<0.05). However, it remains that for the same fluxes, i.e., at any given rate of oxygen consumption, ATP/O was higher in liver mitochondria (Fig. 1B), indicating that liver mitochondria are set to minimize the cost of ATP production thereby optimizing ATP synthesis for cellular processes. In turn, it appears that mitochondria are more "thermogenic" in the skeletal muscle than in the liver, a characteristic that would be further pronounced in resting skeletal muscle, when cellular ATP demand and mitochondrial phosphorylation activity are low (Fig. 1B).

As suggested from Fig. 2 in (Roussel *et al.*, 2002), a difference in the membrane proton conductivity between liver and muscle mitochondria might explain

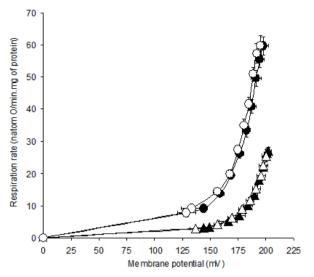


Figure 2. Effect of cold acclimation on basal proton leak Liver (triangles) and skeletal muscle mitochondria (circles) from thermoneutral (open symbols) and cold-acclimated ducklings (filled symbols) were incubated in respiration buffer supplemented with 65 ng/ml nigericin and 1  $\mu$ g/ml oligomycin, as described in Materials and Methods. Oxygen consumption rate and membrane potential were measured in the presence of increasing concentrations of malonate (0–3 mM). Values are means ±S.E.M. of seven independent preparations.

such a difference in ATP yield. The reason for this being that in the effective ATP/O values reported in Fig. 1B, a fraction of the oxygen consumed is associated with proton leaks in the inner membrane and not directly involved in the production of ATP (Stuart et al., 2001). We therefore measured the activity of basal proton leak in both liver and skeletal muscle mitochondria from thermoneutral and cold-acclimated ducklings (Fig. 2). It clearly appeared that the liver inner mitochondrial membrane displayed a lower leak than the skeletal muscle inner mitochondrial membrane. Indeed, at any given membrane potential, the rate of oxygen consumed to counteract the proton leak (mitochondria respiring in the presence of oligomycin) was lower in the liver than in skeletal muscle mitochondria. Interestingly, cold exposure did not alter the basal proton leak of either mitochondrial type (Fig. 2), which may explain why the mitochondrial ATP yield was not different in thermoneutral and cold-acclimated ducklings. Interestingly, it was found previously that cold exposure altered the basal proton leak in muscle and liver mitochondria from rats (Mollica et al., 2005), underlining some species specificities in mitochondrial adaptations toward cold exposure. Finally, it is important to keep in mind that the basal proton leak measured in the present study was not related to the activity of avian uncoupling protein, which requires specific activators (free fatty acids and superoxides) and inhibitor (GDP) to function (Talbot et al., 2003). Alternatively, it has been suggested that the basal proton conductance of mitochondria might depend on the ADP/ATP carrier content (Brand et al., 2005). Since skeletal muscle is well known to have a higher content of the ADP/ATP carrier than liver mitochondria (Rossignol et al., 2000; Mollica et al., 2005; Hulbert et al., 2006), the lower coupling of oxidative phosphorylation reported in duckling skeletal muscle mitochondria would be explained by their higher content of ADP/ATP carrier. This assumption deserves further investigation.

The slope values of the linear relationships shown in Fig. 1A describe the amount of extra oxygen that mitochondria have to consume in order to sustain an additional ATP production imposed by a change in the activity of the phosphorylating system, and so independently of the activity of ATP synthesis-independent proton gradient-consuming processes (e.g., proton leak). As such, these values would be close to, if any a measure of, the mechanistic stoichiometry of mitochondrial oxidative phosphorylation. Notwithstanding, these slopes displayed an average value of  $1.38 \pm 0.02$  and were not significantly different between liver and skeletal muscle mitochondria or between control and cold-acclimated ducklings. Surprisingly, this is a good estimate of the maximal P/O ratio for succinate if we assume that duckling mitochondrial ATP synthase is comprised of three  $\alpha\beta$  pairs, hence three catalytic sites, in its catalytic sector  $F_1$ , and ten c-subunits in its membrane embedded sector,  $F_0$  (Stock *et al.*, 1999), which would predict an H<sup>+</sup>/ATP ratio of 10/3 and an overall coupling stoichiometry of 4.33 (Brand, 2005)

In conclusion, the basal proton conductance of mitochondria may be a determinant of the effective ATP/O ratio but it does not play an important role in the development of non-shivering thermogenesis in cold-acclimated ducklings.

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