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1	Mitochondrial threshold for H2O2 release in skeletal muscle of mammals
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21 Abstract

The aim of the study was to evaluate the interplay between mitochondrial respiration and 22 H₂O₂ release during the transition from basal non-phosphorylating to maximal 23 phosphorylating states. We conducted a large scale comparative study of mitochondrial 24 oxygen consumption, H₂O₂ release and electron leak (% H₂O₂/O) in skeletal muscle 25 mitochondria isolated from mammal species ranging from 7 g to 500 kg. Mitochondrial fluxes 26 27 were measured at different steady state rates in presence of pyruvate, malate, and succinate as respiratory substrates. Every species exhibited a burst of H₂O₂ release from skeletal muscle 28 mitochondria at a low rate of oxidative phosphorylation, essentially once the activity of 29 mitochondrial oxidative phosphorylation reached 26% of the maximal respiration. This 30 threshold for ROS generation thus appears as a general characteristic of skeletal muscle 31 mitochondria in mammals. These findings may have implications in situations promoting 32 33 succinate accumulation within mitochondria, such as ischemia or hypoxia.

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38 **1. Introduction**

In cells, ROS are continuously produced as a byproduct of the mitochondrial oxidative 39 phosphorylation process and more generally of aerobic metabolism (Kowaltowski et al., 2009; 40 Murphy, 2009). Because of its relationship with oxygen consumption, the link between ROS 41 production and metabolic rate has been extensively studied especially in the light of the free 42 radical theory of aging (Müller et al., 2007). Body mass dependence of respiration rates has 43 been largely reported in muscle and liver mitochondria from mammals and birds (Drabkin, 44 1950; Porter and Brand, 1993; 1995; Brand et al., 2003; Boël et al., 2019). An allometry of 45 46 mitochondrial oxidative activity is consistent with the lower activity and/or content of 47 mitochondrial oxidative enzymes reported in several tissues of large mammals compared with 48 small species (Drabkin, 1950; Fried and Tipton, 1953; Kunkel et al., 1956; Emmett and Hochachka, 1981; Phillips et al., 2012). Consequently, all studies found a negative correlation 49 50 between the specific oxygen consumption and body mass at all levels of organization, from 51 whole body to cell and mitochondria. However, regarding mitochondrial ROS production and 52 body mass, similar pattern is often observed in different taxa (Sohal et al., 1989; 1990; Ku et al., 1993; Sohal and Weindruch, 1996; Csiszar et al., 2012; Roussel et al., 2015), but not 53 54 systematically so (e.g. depending on the substrate used) (Lambert et al., 2007; Csiszar et al., 2012; Montgomery et al., 2012). One part of the discrepancies between studies could be 55 explained by the fact that mitochondrial generation of ROS is a complex process which 56 results in the sum of rates from up to eleven mitochondrial sites involved in substrate 57 oxidation and electron transport (Brand, 2016). 58

Notwithstanding such a complex picture of mitochondrial ROS production, the main 59 driver of ROS generation and electron leak (the ratio ROS/O) remains the reduction state of 60 the electron donors in the mitochondrial electron transport system (Quinlan et al., 2013). It 61 has been clearly shown that the ROS generation is non-linearly sensitive to the redox state of 62 respiratory chain intermediates but also to the proton motive force value and so the energetic 63 state of mitochondria (Korshunov et al., 1997; Votyakova and Reynolds, 2001; Starkov and 64 65 Fiskum, 2003; Kikusato and Toyomizu, 2013; Cortassa et al., 2014; Quinlan et al., 2012; Goncalves et al., 2015; Treberg et al., 2018). For instance, when mitochondria increase their 66 activity from basal respiration to maximal phosphorylating respiration, there is a 10% 67 decrease in membrane potential but a 10 fold reduction in ROS generation (Korshunov et al., 68 1997; Starkov and Fiskum, 2003; Kikusato and Toyomizu, 2013). Thereby, ROS generation is 69 maximal at the non-phosphorylating state when the high value of the proton motive force 70 71 restricts the respiration rate and corresponding electron flux, leading to increased electron leak

and ROS formation (Treberg et al., 2018). In contrast, ROS generation decreases as the
intensity of phosphorylating activity increases, reaching a minimum at maximal
phosphorylating activity, due to a higher oxidized state of electron transporter intermediates
(Starkov and Fiskum, 2003; Barja, 2007; Quinlan et al., 2012; Goncalves et al., 2015; Roussel
et al., 2019).

So far, previous works on mitochondrial ROS production have essentially reported on 77 basal non-phosphorylating and maximal phosphorylating states. Such double assessment is 78 useful to compare species, populations or individuals facing environment variations but does 79 80 not fully reflect the interplay between oxygen consumption and ROS formation particularly at 81 intermediate activity levels. Indeed, mitochondria working in the cells are never in a basal 82 non-phosphorylating state and are only rarely running ATP synthesis at full maximal capacity. ROS generation remains low until a threshold value of mitochondrial activity is reached, at 83 84 which point this production abruptly increases (Roussel et al., 2019). For instance, mitochondrial oxidative phosphorylation activity of skeletal muscles in ducklings had to be 85 86 reduced by 60-70% before mitochondrial ROS production sharply increased (Roussel et al., 2019). At some point, such a threshold value defines the frontier between two contrasted 87 88 functional states of mitochondria: i) one state characterized by a high ATP production and a moderate ROS production, and ii) another state characterized by a low ATP production and a 89 high ROS production. 90

On the whole, the variation of mitochondrial thresholds for H_2O_2 generation is 91 globally unknown in animals. Therefore, the aim of the present study was to i) evaluate the 92 interplay between mitochondrial respiration and H_2O_2 release or electron leak (% H_2O_2/O) 93 during the transition from basal non-phosphorylating (LEAK) to maximal oxidative 94 phosphorylating (OXPHOS) states in different mammals' skeletal muscle mitochondria, ii) 95 estimate the range of variation of the resulting threshold values, i.e. the mitochondrial activity 96 97 level at which H₂O₂ release from mitochondria start to increase dramatically and iii) assess the allometry of these parameters in mammal species ranging from 7 g to 500 kg. 98

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100 **2. Materials and methods**

101 2.1. Animals and tissue sampling

102 All experiments were conducted in accordance with animal care guidelines and were 103 approved by the Ethics Committee of Lyon and the Ministère de la Recherche et de 104 l'Enseignement Supérieur. All animals used in the present study were males. African pygmy 105 mice (*Mus minutoides*) and house mice (*Mus musculus*) were obtained from laboratories

(respectively ISEM, Montpellier, France and LBBE, Lyon, France) and killed by cervical 106 dislocation. Syrian hamsters (Mesocricetus auratus) and black rats (Rattus rattus) were 107 obtained from laboratories (Chronobiotron, Strasbourg, France for M. auratus; Jardin 108 Zoologique de la Citadelle, Besançon, France for *R. rattus*) and were killed under isoflurane-109 induced general anesthesia. Fresh tissues for bovines (Bos taurus) and sheep (Ovis aries) 110 were obtained from a local slaughter house (Cibevial, Corbas, France). Fresh tissues for all 111 other mammals were obtained from pest control (nutria - Myocastor coypus and muskrat -112 Ondatra zibethicus), local farmers (rabbits - Oryctolagus cuniculus) or Fondation Pierre 113 114 Vérot at Saint-André-de-Corcy, France (boar – Sus scrofa).

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116 2.2 Mitochondrial isolation

Fresh muscle tissue from every mammal studied was used for mitochondrial extraction. 117 118 Mitochondria were isolated from the hind-limb muscles for the two smallest species (M. minutoides and M. musculus), from a mixture of gastrocnemius and quadriceps muscles for 119 120 medium-size species (M. auratus, R. rattus, M. coypus, O. zibethicus, O. cuniculus), and from a sampling of the fore-leg muscles for the largest species (O. aries, S. scrofa, B. taurus). For 121 122 pygmy mice, skeletal muscle samples from three individuals were used for each mitochondrial preparation. Muscle mitochondria were isolated in an ice-cold isolation buffer 123 (100 mM sucrose, 50 mM KCl, 5 mM EDTA, 50 mM Tris-base, and pH 7.4). Briefly, the 124 mitochondrial isolation procedure involved Potter homogenization, protease digestion and 125 differential centrifugations, with all steps at 4°C as described previously (Boël et al., 2019). 126 Finally, skeletal muscle mitochondrial populations were pelleted at 8,700×g (10 min) and the 127 protein concentration of the mitochondrial suspension was determined by a Biuret method 128 129 with bovine serum albumin as standard.

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131 2.3 *Mitochondrial oxygen consumption rate*

Mitochondrial oxygen consumption rates were determined at 37°C in a respiratory buffer (120 132 133 mM KCl, 5 mM KH₂PO₄, 1 mM EGTA, 2 mM MgCl₂, 0.3% fatty acid-free bovine serum albumin, 1.6 U/mL hexokinase, 20 mM glucose, and 3 mM Hepes, pH 7.4). Mitochondria 134 were energized by adding pyruvate/malate (5/2.5 mM) followed by 5 mM succinate and the 135 rates of basal respiration recorded. Active phosphorylating respiration was initiated by the 136 addition of 4, 10, 20, 100 and 500 µM ADP (Boël et al., 2019). The rate of oxygen 137 consumption was recorded in a closed glass cell fitted with a Clark oxygen electrode (Rank 138 139 Brothers Ltd, Cambridge, UK). Respiratory control ratio (RCR) was calculated as the ratio

between the rates of ADP (500 μM)-induced phosphorylating respiration to basal nonphosphorylating respiration.

142

143 2.4 Mitochondrial reactive oxygen species production

The rate of H₂O₂ released by isolated mitochondria was measured in the respiratory buffer at 144 37°C using a fluorometer (SFM-25, Kontron Instrument, Augsburg, Germany) at excitation 145 and emission wavelengths of 560 nm and 584 nm, respectively. The respiratory buffer was 146 supplemented with 5 U/mL horseradish peroxidase and 1 µM Amplex Red fluorescent dye. 147 148 Respiratory substrates were successively added with 5 mM pyruvate/2.5 mM malate followed 149 by 5 mM succinate and the rates of H_2O_2 generation recorded. Thereafter, H_2O_2 generation 150 was titrated with the sequential addition of ADP at final concentrations of 4, 10, 20, 100, and 500 µM. The fluorescent signal was finally calibrated using a standard curve obtained after 151 152 successive addition of H₂O₂ (up to 35 pmoles). Electron leak (%H₂O₂/O) was calculated as the fraction (%) of the total electron flow that reduces oxygen into H_2O_2 instead of reaching 153 154 cytochrome-c oxidase to reduce oxygen into water (Rey et al., 2013).

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156 2.5. Statistical analyses

157 The individual breakpoint values of oxygen consumption for significant H₂O₂ release have been determined using the package 'segmented' of R core Team. Subsequently, the threshold 158 values for mitochondrial H₂O₂ release representing the percentage maximal oxidative 159 phosphorylation rate at which H₂O₂ release started to increase sharply were determined 160 individually. Following the determination of these parameters, all statistical analyses were 161 conducted on data corrected by the Phylogenetic Independent Contrast model in order to get 162 them phylogenetically independent. From the mammal phylogenetic super-tree proposed by 163 Fritz et al. (2009), a phylogenetic tree with the ten species chosen in the present study was 164 created. Thereafter, the data were corrected by the PIC function that applies the Phylogenetic 165 Independent Contrast method described by Felsenstein (1985), assuming that the evolution 166 167 model for life-history traits follows Brownian motion. After transforming body mass and H_2O_2 release by the common logarithm (log 10), the relationship between these two variables 168 was analyzed with the linear model, with $log_{10}(body mass)$ as a fixed factor. The same 169 170 statistical analyses were made to investigate the impact of body mass on oxygen consumption rate, electron leak, or threshold value. Fits were performed on log₁₀-transformed data with the 171 linear model, with log₁₀(body mass) as fixed factor. For all these allometric relationships, 172 173 normality and homoscedasticity criteria for the model's residues were checked by the

Shapiro-wilk normality test coupled to the Plot Diagnostics for a lm object. Statistical
analyses were done with the R Core Team (R Core Team, 2018) and the packages 'phytools',
'ape', 'multcomp', 'mvtnorm', 'survival', 'TH.data', and 'MASS'.

177

178 **3. Results**

179 *3.1. Mitochondrial bioenergetics parameters*

In the present protocol, mitochondria were energized with pyruvate/malate and the 180 resulting rates of oxygen consumption and H₂O₂ release were recorded before succinate was 181 added in the respiratory chamber. The basal non-phosphorylating respiration rate of skeletal 182 muscle mitochondria respiring on pyruvate/malate was significantly decreased with an 183 increasing body mass of mammals ($F_{(1,8)} = 10.38$; p < 0.05; Fig. 1). In contrast, the rate of 184 H₂O₂ release from mitochondria respiring on pyruvate/malate ($F_{(1,8)} = 2.55$; p = 0.15; Fig. 2) 185 and the corresponding % H₂O₂/O (2.30 × mass^{0.013}; $r^2 = 0.011$; $F_{(1.8)} = 0.23$; p = 0.64; data not 186 shown) were not significantly related to body mass. 187

188 Table 1 describes several bioenergetics parameters of mitochondria isolated from the skeletal muscle of the ten mammalian species and respiring on pyruvate/malate/succinate in 189 190 both basal and ADP-induced active phosphorylating state. The rates of mitochondrial oxygen consumption significantly decreased with increasing body mass in both the basal state (J_0 = 191 157 x mass^{-0.12}; $r^2 = 0.85$; $F_{(1,8)} = 13.94$, p < 0.01) and in the active state, i.e. at maximum rates 192 of ATP synthesis ($J_0 = 761 \text{ x mass}^{-0.10}$; $r^2 = 0.67$; $F_{(1,8)} = 7.77$, p < 0.05). The rate of H₂O₂ 193 release was also related to body mass but only in the active state ($J_{H2O2} = 660 \times mass^{-0.091}$; $r^2 =$ 194 0.33; $F_{(1,8)} = 6.38$, p < 0.05); the variation in the basal state failed to reach significance (J_{H2O2} 195 = 4786 × mass^{-0.073}; r^2 = 0.50; $F_{(1,8)}$ = 4.61, p = 0.06). The electron leak (%ROS/O) was not 196 correlated with body mass in either the basal or active states (% $H_2O_2/O = 3.06 \times mass^{0.048}$; r² 197 = 0.31; $F_{(1,8)} = 0.11$; p = 0.74 and % H₂O₂/O = 0.09 × mass^{0.007}; $r^2 = 0.003$; $F_{(1,8)} = 0.92$; p =198 0.37, in basal and active states respectively). 199

200

201 3.2. Relationships between H_2O_2 generation and oxygen consumption

202 Mitochondrial H_2O_2 release and oxygen consumption at different mitochondrial 203 phosphorylating steady-state rates are presented in Fig. 2. H_2O_2 generation and electron leak 204 ($\%H_2O_2/O$) decreased non-linearly as mitochondrial oxidative phosphorylation activity 205 increased. From the maximum oxidative phosphorylation state to the basal state (from right to 206 left in Fig 2), the rate of H_2O_2 release and corresponding % H_2O_2/O increased slowly at high

oxygen consumption, up to a phosphorylating steady-state rate below which a sharp increase 207 in these parameters occurred. The rate of oxygen consumption at which H₂O₂ release start to 208 increase sharply (i.e. the breaking point of the non-linear curve shown in Fig. 2) was 209 determined for each individual of the ten mammal groups (see "Materials and Methods" 210 section for more details). Fig. 3 shows that the oxygen consumption rate at the breakpoint is 211 negatively correlated with body mass ($F_{(1,8)} = 30.87$; p < 0.05), but the correlation was not 212 quite significant when data were corrected for phylogenetic relatedness of species ($F_{(1,8)}$ = 213 214 4.84; p = 0.059).

215

216 *3.3. Threshold values of mitochondrial ROS generation*

217 The above result indicates that small mammals trigger a sharp H_2O_2 release at higher oxygen consumption rate than larger species. However, the respiratory activities of mitochondria from 218 219 small mammals were also significantly higher than in larger species (Table 1). To test whether the higher mitochondrial oxidative activity of small mammals was responsible for the shift, 220 221 we expressed the oxygen consumption rate at the inflexion point as a percentage of maximal oxidative rate. This fractional value defines a threshold in mitochondrial H₂O₂ release for 222 223 each of the ten mammal species, i.e. the percentage maximal oxidative rate at which H_2O_2 release starts to increase sharply (Roussel et al., 2019). Fig. 3 indicates that threshold values 224 were independent of body mass ($F_{(1,8)} = 0.0002$, p = 0.99). Regardless of the species studied, 225 the oxidative phosphorylation activity had to decrease by approximately 74% before a major 226 increase in H₂O₂ release occurred in skeletal muscle mitochondria. 227

228

229 **4. Discussion**

We followed the entire non-linear dynamic relationship between the intensity of oxidative 230 phosphorylation (i.e. from basal non-phosphorylating to maximal phosphorylating rates) and 231 mitochondrial H₂O₂ release or electron leak (%H₂O₂ /O) in ten mammal species. The present 232 results show a clear threshold for mitochondrial H₂O₂ release and electron leak (%H₂O₂ /O), 233 234 which remain low until a low level of oxidative phosphorylation activity is reached. Hence, the activity of mitochondrial oxidative phosphorylation must reach 26% of the maximal 235 respiration before an H₂O₂ release burst occurs in skeletal muscle of mammals. The threshold 236 values for H₂O₂ release in mammals are slightly lower than those recently published in birds 237 (Roussel et al., 2019). Compared with the value of 26% in skeletal muscle mitochondria from 238 mammals (present study), the threshold values for H₂O₂ release in skeletal muscle 239 mitochondria from ducklings average 35% of the maximal oxidative phosphorylation activity, 240

ranging from 25% to 45% depending on nutritional status and muscle phenotypes (Roussel et 241 242 al., 2019). It has been clearly shown that the ROS generation is non-linearly sensitive to the redox state of respiratory chain intermediates, which remain the main driver of ROS 243 generation (Quinlan et al., 2013). Thus, the kinetics of ROS release could be well explained 244 by the rate of electron leak to generate ROS, which decreases non-linearly during the 245 246 transition from basal to maximal phosphorylating activity due to an enhanced oxidized state of electron transport chain intermediates (Quinlan et al., 2013; Starkov and Fiskum, 2003; 247 Cortassa et al., 2014). These studies further suggest that the non-linear relationships between 248 249 H₂O₂ release and mitochondrial activity might be a general feature of mammal and avian mitochondrial functioning, i.e. a low release of H₂O₂ followed by a sharp increase once a 250 251 threshold of oxidative phosphorylation inactivity is reached.

252 Previous studies have reported a negative relationship between body mass and 253 mitochondrial ROS production in several mammalian tissues, including kidney, heart and liver (Sohal et al., 1990; Ku et al., 1993; Lambert et al., 2007). The present data also show 254 255 that the rates of H₂O₂ release during basal non-phosphorylating respiration and maximal skeletal muscle mitochondria 256 phosphorylating respiration in respiring on 257 pyruvate/malate/succinate were on the whole negatively correlated with body mass of mammals. Yet, mitochondrial respiration and H_2O_2 emission may vary according to the 258 muscle phenotypes, being respectively high and low in slow twitch/oxidative muscle and 259 conversely in fast twitch/glycolytic muscle (Anderson and Neufer, 2006; Picard et al., 2008). 260 In the present study, we assumed that collected muscles were predominantly constituted of 261 mixed glycolytic/oxidative muscle fibers and representative of the whole musculature across 262 the ten species. Nevertheless, it has been reported that for a given skeletal muscle, the 263 activities of oxidative and glycolytic enzymes negatively and positively correlate to body 264 mass in mammals, respectively (Emmett and Hochachka, 1981). If the metabolic 265 (oxidative/glycolytic) and contractile (slow/fast twitch) phenotypes are related each other 266 across mammalian species, then we should expected the mitochondrial oxidative activity and 267 268 H_2O_2 emission in small mammals to be respectively greater (which was the case in the present study) and lower (which was not the case in the present study) than in large mammals. 269 Although, we cannot exclude differences in the composition of skeletal muscle fiber types 270 between mammalian species, we suggest that this characteristic was not the main driver of the 271 negative relationship between body mass and mitochondrial H₂O₂ release reported here. 272

Further insight, it is important to keep in mind that we did not directly measure mitochondrial ROS production, but the rate of H_2O_2 release from mitochondria, which is a

function of ROS producing and consuming processes within mitochondria. Allometry studies 275 276 on mitochondrial antioxidant systems are scarce, but there is some data suggesting no strong relationship between antioxidant enzyme activities of different tissues and body mass in 277 mammals (Tolmasoff et al., 1980; Sohal et al., 1990; Stuart et al., 2013), including 278 279 mitochondrial superoxide dismutase (Page et al., 2010). Another study on four species (camel, 280 goat, rat and mouse) suggests that the activity of mitochondrial glutathione peroxidase might be negatively related to body mass (Al-Otaiba et al., 2010). Although this result has to be 281 confirm with a larger number of species, it would be associated with a higher H₂O₂ release 282 283 from mitochondria of large mammals, which is not the case (present study) (Sohal et al., 1990; Ku et al., 1993; Lambert et al., 2007). If mitochondrial antioxidant systems were the 284 285 main determinant, the rate of H₂O₂ release from mitochondria respiring on pyruvate/malate should be as significantly correlated with body mass as in the presence of succinate, which 286 287 was not the case (present study). Even though, we cannot completely rule out the possible influence of interspecific differences in the matrix H₂O₂ consuming processes, these processes 288 289 were not the main driver of the negative allometric relation reported in muscle mitochondria respiring on pyruvate/malate/succinate (Table 1). 290

291 Of note, these interspecific correlations between body mass and mitochondrial ROS 292 release are essentially reported when mitochondria are respiring on succinate (Lambert et al., 2007). Succinate oxidation generates ROS production both with a reverse electron transfer at 293 the quinol site of complex I and with a forward reaction at the flavin site of complex II 294 (Quinlan et al., 2012b). Interestingly, these two sites account for half of the total rate of H_2O_2 295 production in skeletal muscle mitochondria during rest (Goncalves et al., 2015). In the present 296 study, the addition of succinate in mitochondria respiring on pyruvate/malate resulted in a 297 drastic increase in H₂O₂ release, which was highly sensitive to the addition of ADP (Table 1). 298 Such high dependence of H₂O₂ release upon mitochondrial activity (basal non-299 300 phosphorylating state versus active phosphorylating state) suggests that the reverse electron transfer through complex I would be involved in the differences in the H₂O₂ release between 301 302 species (Zoccaroto et al., 2009; Goncalves et al., 2015). Indeed, the endergonic nature of the reversed electron transport makes ROS generation by this process highly sensitive to proton 303 motive force (Korshunov et al., 1997; Votyakova and Reynolds, 2001; Starkov and Fiskum, 304 2003), and so to proton motive force consuming pathways such as ATP synthesis. These 305 observations thus suggest that the low level of H₂O₂ production in the larger mammals might 306 result from a low content of complex I (Lambert et al., 2010). The allometry of ROS 307 production in mammals might also be explained by changes in complex II level and/or 308

activity. This is suggested by an earlier study reporting a negative relationship between the 309 activity of succinate oxidase and body mass in several tissues of mammals (Fried and Tipton, 310 1953). Nevertheless, the ROS generation from complex II reaches a maximum at a low 311 concentration of succinate (<400 µM), becoming very low at high concentrations (Quinlan et 312 al., 2012b), such as the 5 mM used in the present study. We thus postulate that the low level 313 of H₂O₂ production in mitochondria from the larger mammals would result mainly from a 314 lower activity of the reversed electron transport through complex I, a hypothesis that remains 315 316 to be investigated thoroughly.

317 The quinol site in complex I, which is active during reverse electron transport, produces a significant rate of ROS in resting skeletal muscle, heart and cell lines from several 318 other tissues (Goncalves et al., 2015; Brand et al., 2016; Wong et al., 2019). This ROS 319 generation by the reverse electron transport in complex I is exacerbated with increasing 320 321 concentrations of succinate (Zoccaroto et al., 2009). Hence, this mechanism appears to be of pathological and physiological importance in situations promoting succinate accumulation 322 323 within mitochondria, such as ischemia or hypoxia (Hochachka et al., 1975; Starkov, 2008; Ralph et al., 2011). Succinate accumulates in several tissues of mammals, including heart, 324 325 liver, kidney and brain during ischemia (Chouchani et al., 2013), or during breath-hold dives 326 (Hochachka et al., 1975). During reperfusion, its rapid oxidation leads to a burst of ROS production and ischemia-reperfusion injury (Dröse, 2013; Chouchani et al., 2016). In turn, the 327 ROS release from mitochondria could also serve as a signal, which may regulate cell 328 functions (Guzy and Schumaker, 2006; Stowe and Camara, 2009). For instance, ROS play a 329 vital role in the regulation of muscle metabolism and adaptation towards exercise (Morales-330 Alamo and Calbet, 2016). ROS signaling may also be involved in the up-regulation of 331 antioxidant pathways and repair mechanisms of cellular damage reported in diving 332 vertebrates, limiting the oxidative stress associated with repeated prolonged apnea and re-333 oxygenation cycles (Filho et al., 2002; Rey et al., 2016). Another interesting perspective is the 334 differential oxygen sensitivity of these two mitochondrial processes, i.e. the oxygen 335 respiration and the H₂O₂ release (Hoffman et al., 2007; Treberg et al., 2018). This oxygen 336 sensitivity of mitochondrial ROS generation varies between the respiratory substrates and the 337 ROS-generating sites within mitochondria (Hoffman and Brookes, 2009). For instance, it has 338 been shown that complex I remains the main source of ROS generation at low concentration 339 of oxygen (Hoffman and Brookes, 2009). Whether the oxygen sensitivity differs between 340 tissues and across mammalian species remain an open question. 341

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- **Competing interests**. The authors declare that they have no competing financial interests.

Data Sharing. Data available on request from the corresponding authors.

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		Respiration rate (nmol O.min ⁻¹ .mg protein ⁻¹)		H ₂ O ₂ release (pmol H ₂ O ₂ .min ⁻¹ .mg protein ⁻¹)		Electron leak (%H ₂ O ₂ /O)	
Species (n)	Body weight (g)	Basal state	Active state	Basal state	Active state	Basal state	Active state
Pygmy mouse (10) (<i>M. minutoides</i>)	7.2 ± 0.4	141 ± 8	832 ± 49	5,062 ± 357	$1,035 \pm 76$	3.69 ± 0.32	0.13 ± 0.02
House mouse (5) (<i>M. musculus</i>)	18.8 ± 0.6	98 ± 4	518 ± 16	$4,598 \pm 728$	643 ± 178	4.75 ± 0.86	0.12 ± 0.03
Golden hamster (8) (<i>M. auratus</i>)	105.5 ± 2.6	95 ± 6	418 ± 14	$4,067 \pm 290$	483 ± 24	4.29 ± 0.17	0.12 ± 0.0
Black rat (8) (<i>R. rattus</i>)	194.4 ± 7.2	88 ± 10	487 ± 50	$2,452 \pm 212$	216 ± 34	2.92 ± 0.27	0.04 ± 0.0
Muskrat (1) (<i>O. zibethicus</i>)	1,600	85	443	2,091	206	2.5	0.05
Rabbit (6) (<i>O. cuniculus</i>)	$3,200 \pm 301$	43 ± 3	218 ± 9	$2,041 \pm 190$	209 ± 27	4.84 ± 0.49	0.10 ± 0.0
Nutria (3) (<i>M. coypus</i>)	$5,150 \pm 760$	51 ± 2	222 ± 25	$2,254 \pm 418$	181 ± 30	4.42 ± 0.83	0.08 ± 0.0
Sheep (6) (<i>O. aries</i>)	34,217 ± 1,633	42 ± 7	334 ± 60	$1,834 \pm 173$	311 ± 92	4.97 ± 0.70	0.08 ± 0.0
Boar (5) (S. scrofa)	$79,800 \pm 4,420$	52 ± 7	211 ± 18	$3,422 \pm 357$	460 ± 36	6.94 ± 0.80	0.23 ± 0.0
Bovine (4) (<i>B. Taurus</i>)	486,300 ± 25,367	36 ± 5	260 ± 39	2,112 ± 226	242 ± 63	6.21 ± 0.85	0.09 ± 0.0

Table 1: Properties of muscle mitochondria isolated from mammals of different mass.

- 519 Values are means ± SEM [number of independent mitochondrial preparations (n) in parentheses alongside the name of animals]. Basal state of
- respiration was initiated by the addition of pyruvate/malate/succinate (5/2.5/5 mM) as respiratory substrates. Active state of respiration was
- 521 measured following the addition of $500 \,\mu M \,ADP$

522 Figure legends

Figure 1: Body mass dependence of basal respiration rate (J_0) and H_2O_2 release (J_{H2O2}) in skeletal muscle mitochondria respiring on pyruvate/malate. Skeletal muscle mitochondria were isolated from ten mammal species. Red symbols are two species taken to cover the whole range of body mass (bovine with the highest mass and pygmy mice with the lowest mass). Values are mean \pm s.d. from (n) independent mitochondrial preparations; (n) is given alongside the name of the species

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Figure 2: Relationships between oxidative phosphorylation activity and H_2O_2 release (A) or electron leak (B). Mitochondria were isolated from skeletal muscle of ten mammal species and were respiring on pyruvate/malate/succinate. Red symbols illustrate the relationships for two species taken to cover the whole range of body mass (bovine with the highest mass and pygmy mice with the lowest mass). Values are mean \pm s.d. from (n) independent mitochondrial preparations; (n) is given alongside the name of the species

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Figure 3: Body mass dependence of threshold H₂O₂ production values. Data are expressed as 537 538 the oxygen consumption rate at the breaking point of H₂O₂ release or as a percentage of maximal oxidative phosphorylation activity (see text for more details) in skeletal muscle 539 540 mitochondria isolated from ten mammal species. A mixture of pyruvate/malate/succinate was used as respiratory substrates. Red symbols are two species taken to cover the whole range of 541 542 body mass (bovine with the highest mass and pygmy mice with the lowest mass). Values are 543 mean \pm s.d. from (n) independent mitochondrial preparations; (n) is given in Fig. 2 alongside the name of the species. 544

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