

1 **Mitochondria are physiologically maintained at close to 50 °C**

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28

29 **Abstract**

30 In warm-blooded species, heat released as a by-product of metabolism ensures stable internal  
31 temperature throughout the organism, despite varying environmental conditions.  
32 Mitochondria are major actors in this thermogenic process. The energy released by the  
33 oxidation of respiratory substrates drives ATP synthesis and metabolite transport, with a  
34 variable proportion released as heat. Using a temperature-sensitive fluorescent probe targeted  
35 to mitochondria, we measured mitochondrial temperature in situ under different physiological  
36 conditions. At a constant external temperature of 38 °C, mitochondria were more than 10 °C  
37 warmer when the respiratory chain was fully functional, both in HEK cells and primary skin  
38 fibroblasts. This differential was abolished by respiratory inhibitors or in cells lacking  
39 mitochondrial DNA, but enhanced by expressing thermogenic enzymes such as the alternative  
40 oxidase or the uncoupling protein UCP1. The activity of various RC enzymes was maximal  
41 at, or slightly above, 50 °C. Our study prompts a re-examination of the literature on  
42 mitochondria, taking account of the inferred high temperature.

43

44 **Main Text**

45 Mitochondrial targeting of the recently developed, temperature-sensitive fluorescent probe  
46 (Fig. S1A), Mito-Thermo-Yellow (MTY)(*I*) was confirmed in both HEK293 cells and  
47 primary skin fibroblasts, based on colocalization with the well-characterized MitoTracker  
48 green (MTG; Fig. 1A). Initial mitochondrial capture was dependent on the maintenance of a  
49 minimal membrane potential(*I*). Fluorescence remained stable over 2 hours in HEK293 cells,  
50 but mitochondrial MTY retention varied between cell lines (Fig. S2A). The exact sub-  
51 mitochondrial location of the probe is yet to be established, although it has been postulated to  
52 reside within the inner membrane. No toxicity of MTY (100 nM in culture medium) could be  
53 detected after 3 days.

54 The fluorescence of MTY (measured in solution at 562 nm) was progressively and reversibly  
55 decreased by as the medium temperature increased (about 50% from 25 to 45°C; Fig. 1B, a,  
56 b). Using a thermostated, magnetically stirred, closable 750 µl quartz-cuvette fitted with an  
57 oxygen sensitive optode device(2) we simultaneously studied oxygen consumption or tension  
58 and changes in MTY fluorescence (Fig. 1C). Adherent cells were harvested and pre-loaded  
59 for 20 min with 100 nM MTY, harvested and washed, then kept as a concentrated pellet at  
60 38°C for 10 min, reaching anaerobiosis in < 1 min. When cells were added to the oxygen-rich  
61 medium they immediately started to consume oxygen (red trace), accompanied by a  
62 progressive decrease of MTY fluorescence (blue trace; phase I; Fig. 1C). In the absence of  
63 any inhibitor, the fluorescence gradually reached a stable minimum (phase II). Once all the  
64 oxygen in the cuvette was exhausted (red trace) the shift of MTY fluorescence reversed  
65 (phase III), returning gradually almost to the starting value (phase IV). To calibrate the  
66 fluorescence signal, the temperature of the extra-cellular medium was increased stepwise  
67 (green trace). MTY fluorescence returned to the prior value when the medium was cooled to  
68 38 °C (Fig. 1C, phase V). Based on this calibration, we estimate the rise in mitochondrial  
69 temperature due to the activation of respiration as ~10 °C (n=10, range 7-12 °C). At the  
70 lowest (phase II) and highest fluorescence value (38°C, imposed by the water bath; phase IV),  
71 the signal was proportional to the amount of added cells (Fig. 1D, a). Cell number did not  
72 affect the maximal rate of fluorescence decrease (phase I), but, once anaerobic conditions had  
73 been reached (phase III, initial), it was inversely related to the initial rate of fluorescence  
74 increase (Fig. 1D, b). To confirm that the observed fluorescence changes were due to  
75 mitochondrial respiration and not some other cellular process, we depleted HEK cells of their  
76 mtDNA with ethidium bromide (EtBr) to a point where cytochrome c oxidase activity was  
77 less than 2% of that in control cells (Fig. 1E, a; Fig. S1C). In EtBr-treated cells, no MTY  
78 fluorescence changes were observed under aerobiosis and cyanide treatment had no effect

79 (Fig. 1E, b). Because MTY is derived from the membrane potential-sensitive dye rhodamine,  
80 whose fluorescence is essentially unaffected by temperature (Figure 1B, b), we investigated  
81 whether the observed changes MTY fluorescence could be influenced by altered membrane  
82 potential, or by an associated parameter, *e.g.* pH. We therefore compared the response of  
83 MTY fluorescence to cyanide or oligomycin (Fig. 2A, B; Fig. S1D) exerting opposite effects  
84 on membrane potential and pH gradient (Fig. 2C). To avoid the possibly confounding effect  
85 of anaerobiosis, the quartz cuvette was kept uncapped in this experiment, with the oxygen  
86 tension rather than the rates of oxygen uptake being recorded (red traces). Once MTY  
87 fluorescence was stabilized (maximal mitochondrial heating), and the medium re-oxygenated,  
88 cyanide was added, causing a progressive increase in MTY fluorescence to the starting value  
89 (Fig. 2A). Note that, when cyanide was present from the start of the reaction, fluorescence  
90 changes and oxygen uptake were both abolished (Fig. 2A, dotted lines). Adding oligomycin in  
91 lieu of cyanide decreased oxygen consumption and, as observed with cyanide, brought about  
92 an increase in MTY fluorescence (Fig. 2B). Added first, oligomycin strongly decreased  
93 oxygen uptake, abolishing MTY fluorescence decrease (Fig. 2B, dotted lines). Taken  
94 together, these experiments imply that electron flow through the respiratory chain (RC) rather  
95 than membrane potential controls mitochondrial temperature. This conclusion is supported by  
96 the similar effects observed with two other respiratory inhibitors (Fig. S1D), affecting either  
97 RC complex I (CI; rotenone, Fig. 2D) or III (CIII; antimycin, Fig. 2E). Despite their different  
98 effects on the redox state of the various RC electron carriers, these inhibitors blocked oxygen  
99 uptake and again triggered an increase in MTY fluorescence. Taking advantage of cyanide  
100 removal from cytochrome oxidase to form cyanohydrin in the presence of  $\alpha$ -ketoacids under  
101 aerobiosis(3), we confirmed that the blockade of the respiratory chain did not result in MTY  
102 leakage from the mitochondria since pyruvate addition resulted in oxygen uptake resuming  
103 and MTY fluorescence decrease, both being inhibited by a further addition of antimycin (Fig.

104 2F). Noticeably the fluorescence of an endoplasmic reticulum (ER)-targeted version of the  
105 probe in HEK cells and skin fibroblasts was essentially unaffected by the activity of the  
106 mitochondria when modulated by cyanide, pyruvate or antimycin (Fig. S3).

107 We next studied MTY probe behavior in HEK cells made partially deficient for CI by varying  
108 amounts of added rotenone. The rate of MTY fluorescence decrease was proportional to the  
109 residual respiratory electron flux (Fig. 3A) while the maximal temperature as judged from  
110 MTY fluorescence (phase II) was essentially unchanged (Fig. 3A; inset). We next tested the  
111 effect of expressing the cyanide-insensitive non-proton motive alternative oxidase from *Ciona*  
112 *intestinalis* (AOX; Fig. 3B, S1E), whose activity is unmasked in the presence of cyanide(4).  
113 Before cyanide addition, the decrease in MTY fluorescence in AOX-expressing cells was  
114 similar to control cells, consistent with previous inferences that the enzyme does not  
115 significantly participate in uninhibited cell respiration. However upon cyanide addition,  
116 AOX-endowed cells maintained low MTY fluorescence (Fig. 3B, blue trace), despite oxygen  
117 consumption decreasing by more than 50% (red trace). The increased ratio of heat generated  
118 to respiration is consistent with the predicted thermogenic properties of AOX. Subsequent  
119 addition of 0.1 mM propylgallate, which inhibits AOX, almost completely abolished the  
120 residual respiration and brought MTY fluorescence back to its starting value (corresponding  
121 to 38°C).

122 So as to circumvent the fact that we were not able to use chemical uncouplers with this  
123 probe(1), we used HEK cells engineered to express the uncoupling protein 1 (UCP1; Fig. 3C,  
124 S1F). As expected, UCP1 conferred an increased rate of respiration, which was only partially  
125 inhibited by oligomycin (red trace), and accompanied by an even greater drop in MTY  
126 fluorescence, equivalent to a temperature at least 15 °C above the cellular environment.  
127 The surprisingly high inferred mitochondrial temperatures prompted us to check the  
128 dependence on assay medium temperature of RC enzyme activities measured under  $V_{\max}$

129 conditions (Fig. 4A, B) in crude extracts, where mitochondrial membrane integrity is  
130 maintained. Antimycin-sensitive CIII, malonate-sensitive succinate:cytochrome c reductase  
131 (CII+CIII) and cyanide-sensitive cytochrome c oxidase (CIV) activities all showed  
132 temperature optima at or slightly above 50°C, whilst these activities tended gradually to  
133 decrease again as the temperatures were raised further (Fig. 4A). This was not so for those  
134 enzymes whose activities can be measured in vitro only after osmotic disruption of both outer  
135 and inner mitochondrial membranes (Fig. 4B). Oligomycin-sensitive ATPase (CV) activity  
136 was optimal around 46 °C, whereas rotenone-sensitive NADH quinone-reductase (CI) activity  
137 declined sharply at temperatures above 38 °C. Under the even more disruptive conditions of  
138 clear-native electrophoresis, which requires detergent-dispersal of the membrane-bound  
139 complexes, in-gel activity even of CII or CIV was impaired at temperatures above 42 °C and  
140 46 °C, respectively (Fig. 4C). This strongly suggests a vital role for the lipid components of  
141 the inner mitochondrial membrane in the stabilization of the RC complexes at high  
142 temperature. We next analyzed the temperature profile of RC activity of primary skin  
143 fibroblasts. For CII+CIII, CIII and CIV (Fig. 4D), as well as CV (Fig. 4E) similar temperature  
144 optima were observed as in HEK cells, whilst MTY fluorescence (Fig. 4F) also indicated  
145 mitochondria being maintained at least 6-10 °C above environmental temperature.  
146 Our findings raise numerous questions concerning the biochemistry, physiology and  
147 pathology of mitochondria. The physical, chemical and electrical properties of the inner  
148 mitochondrial membrane and of mitochondria in general, will need to be re-evaluated, given  
149 that almost all previous literature reflects experiments conducted far from physiological  
150 temperature. Traditional views of the lipid component of the respiratory membrane as a lake  
151 in which the RC complexes are floating resulting in a random-diffusion model of electron  
152 transfer or, more recently, as a sealant occupying the space between tightly packed  
153 proteins(5), need to be revised in favour of one that considers it more as a glue that maintains

154 the integrity of the respiratory complexes. The effects of respiratory dysfunction need to be  
155 reconsidered, to include those attributable to temperature changes, such as effects on  
156 membrane fluidity, electrical conductance and transport. RC organization into  
157 supercomplexes(6, 7) should be re-examined at more realistic temperatures, using methods  
158 other than CNE, which does not maintain functional integrity even of the RC complexes  
159 individually. Finally, whilst the subcellular distribution of mitochondria (e.g. perinuclear, or  
160 synaptic) has previously been considered to reflect ATP demand, mitochondria should also be  
161 considered as a source of heat, potentially relevant in specific cellular or physiological  
162 contexts, not just in specifically thermogenic tissues like brown fat.

163

164 **Legends to Figures**

165 **Figure 1** Determination of mitochondrial temperature in intact human cells.

166 A: The temperature-sensitive probe MitoThermoYellow (MTY: a, b) co-localizes with  
167 MitoTrackerGreen (MTG: c, d, merge: e, f) in HEK293 cells and in primary skin fibroblasts  
168 (e, f). B: a, Fluorescence excitation (green) and emission (red) spectra of MTY (1 mM) in  
169 solution in 2 ml PBS at 25°C and 45°C; b, Response of MTY (blue and red) and rhodamine  
170 (green) fluorescence (1 mM) in solution in 2 ml PBS to temperature (34 to 64°C). Note that  
171 the linear decrease of MTY fluorescence to increasing temperature (blue) is essentially  
172 reversed upon cooling (red) of the medium to the initial temperature. c, Linear increase of  
173 fluorescence of HEK cells (preloaded 10 min before trypsinization with 100  $\mu$ M MTY)  
174 according to cell number (using cell protein concentration as surrogate parameter). C, The  
175 definition of the various phases of MTY-preloaded HEK cell fluorescence adopted in this  
176 study. phase I: cell respiration (red trace) after cells are exposed to aerobic conditions in PBS,  
177 resulting in decreased MTY fluorescence (blue trace) as mitochondria heat up; phase II: cell  
178 respiration under aerobic conditions, which has reached a steady-state of MTY fluorescence  
179 (maximal warming of mitochondria). Cells were initially maintained for 10 min at 38°C under  
180 anaerobic conditions, before being added to the cuvette; phase III: cell respiration that has  
181 arrested due to oxygen exhaustion - MTY fluorescence progressively increases to the starting  
182 value, as mitochondria cool down; phase IV: respiration remains stalled due to anaerobiosis;  
183 after reaching steady-state, MTY fluorescence is dictated only by the water bath temperature;  
184 phase V, respiration remains stalled due to anaerobiosis; temperature of the cell suspension  
185 medium (green trace) shifted by stepwise adjustments to water-bath temperature, followed by  
186 return to 38°C. Measurements were carried out in a quartz chamber closed except for a 0.6  
187 mm addition hole in the hand-made cap. D, Maximal rate of decrease of MTY fluorescence  
188 (% , blue circles; corresponding with mitochondrial warming) is not significantly affected by



189 cell number, whereas initial fluorescence increase in presence of cyanide (% , green circles,  
190 corresponding with initial rate of mitochondrial cooling) is modulated by cell number. D: a,  
191 HEK cells were made severely deficient for cytochrome oxidase by culture (10 days) in the  
192 presence of ethidium bromide (EtBr; 1  $\mu\text{g}/\text{ml}$ ). Cytochrome c oxidase activity (blue circles)  
193 declined to a few percent of the activity measured at  $t=0$ , whilst citrate synthase activity  
194 (green circles) was little changed; b, The fluorescence of EtBr-treated HEK cells (10 days of  
195 EtBr treatment) pre-loaded with MTY (blue continuous line) does not decrease following  
196 suspension in oxygenated medium, whilst that of control HEK cells (blue dotted lines)  
197 follows the profile documented in Fig. 1C; in contrast to control cells (red dotted line), EtBr-  
198 treated HEK cells do not consume oxygen (red continuous line).

199

200 **Figure 2.** The rate of respiratory electron flow determines the temperature of mitochondria in  
201 intact HEK cells.

202 A: The effect of 1 mM cyanide on MTY fluorescence (blue lines) and cell respiration (red  
203 lines), when added under aerobic conditions (continuous lines), or when present from the start  
204 of the experiment (dotted lines). Changes in the temperature of the cell suspension medium  
205 (green line), imposed by water bath adjustment, were used to calibrate the MTY fluorescence  
206 changes. B: The effect of 12.5  $\mu\text{M}$  oligomycin on MTY fluorescence (blue lines) and oxygen  
207 tension (affected by cell respiration balanced by medium stirring) in the uncapped quartz-  
208 cuvette (red lines), when added to free respiring cells (continuous lines), or when present from  
209 the start of the experiment (dotted lines). C: a, effects of different inhibitors on rhodamine  
210 fluorescence, in digitonin (0.001%)-permeabilized HEK293 cells supplied with 10 mM  
211 succinate and 0.1 mM ADP as indicated. Under state 3 conditions 12.5  $\mu\text{M}$  oligomycin and  
212 0.8 mM KCN have qualitatively opposite effects on rhodamine fluorescence, used as an  
213 indicator of membrane potential ( $\Delta\Psi$ ). The effects of 0.6  $\mu\text{M}$  antimycin (D) and 2  $\mu\text{M}$

214 rotenone (E) on MTY fluorescence and oxygen tension, plotted as for oligomycin in (B, b). F:  
215 The effect of adding pyruvate on MTY fluorescence (blue line) and oxygen uptake (red line)  
216 by KCN-inhibited HEK cells. Temperature calibration (green line) of MTY fluorescence as in  
217 A.

218

219 **Figure 3.** Effects on mitochondrial temperature of respiratory inhibitors, uncouplers and  
220 expression of heterologous mitochondrial proteins.

221 A: Effect of variable rotenone addition to control HEK cells on the rates of oxygen uptake and  
222 fluorescence decrease of MTY. Rotenone was added at  $t=4$  min, rates calculated from 4 to 7  
223 min expressed as a percent of initial rate. Inset: Maximal warming up of HEK cell  
224 mitochondria in the absence or presence of 20  $\mu$ M rotenone. B, C: Changes in MTY  
225 fluorescence (blue lines), cell respiration (B, C) (red lines) and temperature of cell suspension  
226 medium (green line), with additions of inhibitors as shown (KCN, n-propyl gallate and/or  
227 oligomycin), alongside Western blots confirming expression or knockdown of the indicated  
228 genes *Ciona intestinalis* alternate oxidase, AOX (B), UCP1 (C), alongside loading controls as  
229 indicated. Traces for control cells are shown by dotted lines.

230

231 **Figure 4.** Effects of assay medium temperature on RC activities. Temperature profile of (A,  
232 D) cytochrome c oxidase (CIV), malonate-sensitive succinate-cytochrome c reductase  
233 (CII+CIII) and antimycin-sensitive decylubiquinol:cytochrome c reductase (CIII) activity in  
234 (A) HEK293 cells, and (D) primary skin fibroblasts, after two freeze-thaw cycles, and (B, E)  
235 oligomycin-sensitive ATPase (CV) and rotenone-sensitive NADH:decylubiquinone (CI)  
236 activity, after disruption of inner mitochondrial membrane in frozen (B) HEK293 cells and  
237 (E) primary skin fibroblasts, by osmolysis with water(8). Colours denote optimal (green),  
238 minimal (red) and degrees of intermediate (pale greens, yellow) activity. Grey bars indicate

239 optimal temperature range. C: CNE in-gel activities of CI, CIV and CII extracted from  
240 mitochondria and incubated for 10 min at (a) 4 °C , (b) 37°C, (c) 42 °C, (d) 46 °C and (e) 55  
241 °C, also plotted graphically (lower panel). F: Changes in MTY fluorescence (blue), cell  
242 respiration (red) and temperature of cell suspension medium (green), for primary skin  
243 fibroblasts, as denoted in Fig. 2A, with addition of KCN as shown.

244

## 245 **Material and Methods**

246

### 247 **Cell culture**

248 Human cells derived from Embryonic Kidney (HEK293), Hepatoma Tissue Culture (HTC-  
249 116) and from large cell lung cancer (NCI-H460) cells (American Type Culture Collection,  
250 Manassas, VA 20108 USA) were cultured in DMEM medium containing 5 g/l glucose  
251 supplemented by 2 mM glutamine (as Glutamax™), 10% foetal calf serum, 1 mM pyruvate,  
252 100 µg/ml penicillin/streptomycin each. AOX-(9) or UCP-endowed(10, 11) HEK293 cells  
253 were obtained as previously described.

254 Primary skin fibroblasts were derived from healthy individuals and grown under standard  
255 condition in DMEM glucose (4.5 g/l), 6 mM glutamine, 10% FCS, 200 µM uridine,  
256 penicillin/streptomycin (100 U/ml) plus 10 mM pyruvate.

257

### 258 **Immunoblot analyses and *in gel* enzyme activity assays**

259 For the Western blot analysis, mitochondrial proteins (50 µg) were separated by SDS–PAGE  
260 on a 12% polyacrylamide gel, transferred to a nitrocellulose membrane and probed overnight  
261 at 4°C with antibodies against the protein of interest, AOX 1:10,000<sup>10</sup> , UCP1 1:10,000<sup>11</sup> .  
262 Membranes were then washed in TBST and incubated with mouse or rabbit peroxidase-  
263 conjugated secondary antibodies for 2 h at room temperature. The antibody complexes were

264 visualized with the Western Lightning Ultra Chemiluminescent substrate kit (Perkin Elmer).  
265 For the analysis of respiratory chain complexes, mitochondrial proteins (100 µg) were  
266 extracted with 6% digitonin and separated by hrCN-PAGE, on a 3.5–12% polyacrylamide gel.  
267 Gels were stained by in gel activity assay (IGA) detecting CI, CII and CIV activity as  
268 described<sup>12</sup>.

269

### 270 **Staining procedures and life cell imaging**

271 Cells (HEK293, MTC, NIC, primary skin fibroblasts) were seeded on glass coverslips and  
272 grown inside wells of a 12 well-plate dish for 48 h in standard growth media at 37 °C, 5%  
273 CO<sub>2</sub>. The culture medium was replaced with pre-warmed medium containing fluorescent  
274 dyes, 100 nmol MitoTracker Green (Invitrogen M7514) and 100 nmol MitoThermo  
275 Yellow(1) or 100 nmol ER Thermo Yellow(12). After 10 min the staining medium was  
276 replaced with fresh pre-warmed medium or PBS buffer and cells were observed immediately  
277 by Leica TCS SP8 confocal laser microscopy.

278

### 279 **Assay of mitochondrial respiratory chain activity**

280 The measurement of RC activities was carried out using a Cary 50 spectrophotometer (Varian  
281 Australia, Victoria, Australia), as described(13). Protein was estimated using the Bradford  
282 assay.

283

### 284 **Simultaneous spectrofluorometric, temperature, oxygen uptake assay**

285 Detached sub-confluent HEK (NCI, HTS) cells (25 cm<sup>2</sup> flask) or trypsinized sub-confluent  
286 skin fibroblasts (75 cm<sup>2</sup> flask) were treated for 20 min with 100 µM MTY in 10 ml DMEM,  
287 and spun down (1,500 g x 5 min). The pellet is resuspended in 1 ml PBS, cells being next  
288 spun down (1,500 g x 5 min) and kept as a concentrated pellet. After anaerobiosis (checked

289 by inserting an optic fiber equipped with an oxygen-sensitive fluorescent terminal sensor  
290 (Optode device; FireSting O<sub>2</sub>, Bionef, Paris, France) was established (10 min incubation of  
291 the pellet at 38°C;), cells (1 mg prot) were added to 750 µl of 38°C-thermostated medium.  
292 The fluorescence (excitation 542 nm, emission 562 nm for MTY; excitation 559 nm, emission  
293 581 nm for ERTY), the temperature of the medium in the cuvette and the respiration of the  
294 intact cell suspension were simultaneously measured in a magnetically-stirred, 38°C-  
295 thermostated 1 ml-quartz cell in 750 µl of PBS using the Xenius XC spectrofluorometer  
296 (SAFAS, Monaco). Oxygen uptake was measured with an optode device fitted to a handmade  
297 cap, ensuring either closing of the quartz-cell yet allowing micro-injections (hole with 0.6 mm  
298 diameter) or leaving the quartz-cell open to allow for constant oxygen replenishment.  
299 Alternatively, untreated HEK293 cells (250 µg protein) were added to 750 µl of medium  
300 consisting of 0.25 M sucrose, 15 mM KCl, 30 mM KH<sub>2</sub>PO<sub>4</sub> (pH 7.4), 5 mM MgCl<sub>2</sub>, EGTA 1  
301 mM, followed by addition of rhodamine to 100 nM and digitonin to 0.01 % w/v. The  
302 permeabilized cells were successively given a mitochondrial substrate (10 mM succinate) and  
303 ADP (0.1 mM) to ensure state 3 (phosphorylating) conditions, under which either 6.5 µM  
304 oligomycin or 1 mM cyanide was added.

305

### 306 **Statistics**

307 Data are presented as mean ± SD statistical significance was calculated by standard unpaired  
308 one-way ANOVA with Bonferroni post-test correction; a  $p < 0.05$  was considered statistically  
309 significant (GraphPad Prism).

310

### 311 **Data availability statement.**

312 The authors declare that all data supporting the findings of this study are  
313 available within the paper and its supplementary information files.

314

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331

## 332 **End notes**

333

## 334 **Abbreviations**

335 AOX, Alternative oxidase; CNE, Clearnative Electrophoresis; ERTY, Endoplasmic  
336 Reticulum Thermo Yellow; EtBr, Ethidium bromide; MTG, MitoTracker Green; MTY,  
337 MitoThermo Yellow; RC, Respiratory chain; UCP1 Uncouplin protein 1

338

## 339 **Acknowledgements**

340 This work was supported by French (ANR MITOXDRUGS to DC, PB, MR and PR) and  
341 European (E-rare Genomit to DC, PB, MR and PR) institutions and patient's associations to  
342 PB and PR Association d'Aide aux Jeunes Infirmes (AAJI), Association contre les Maladies  
343 Mitochondriales (AMMi), Association Française contre l'Ataxie de Friedreich (AFAF), and  
344 Ouvrir Les Yeux (OLY).

345

## 346 **Authors contribution**

347 M.R., HT.J. and P.R. conceived the project. HT.J., M.J., YT. C. and P.R. wrote the  
348 manuscript. D.C., P.B., R. EK., HH.H., S.K., M.R. and P.R. conducted research. All authors  
349 contributed to data analysis and manuscript preparation.

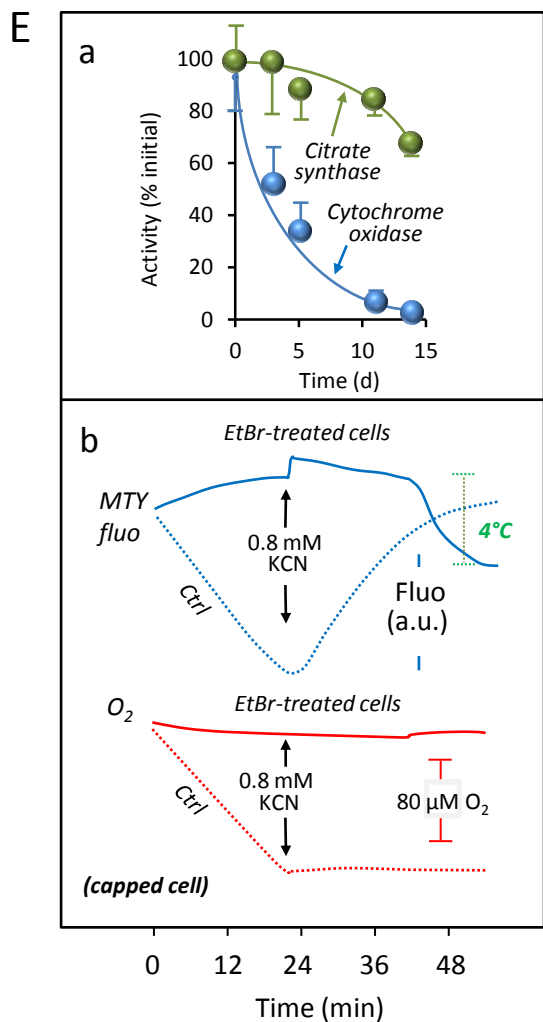
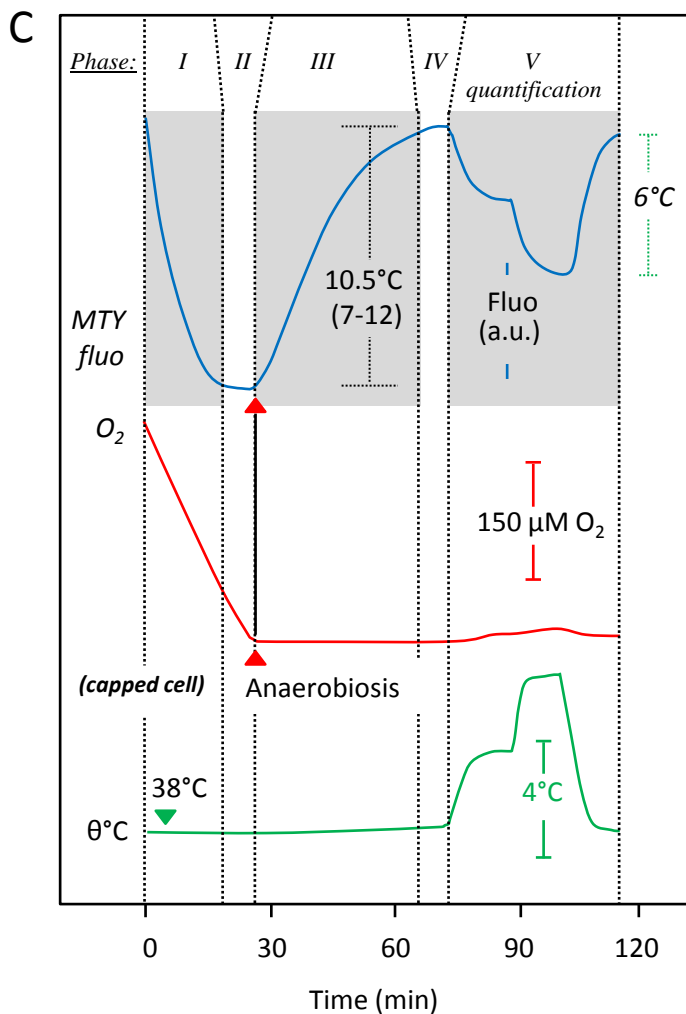
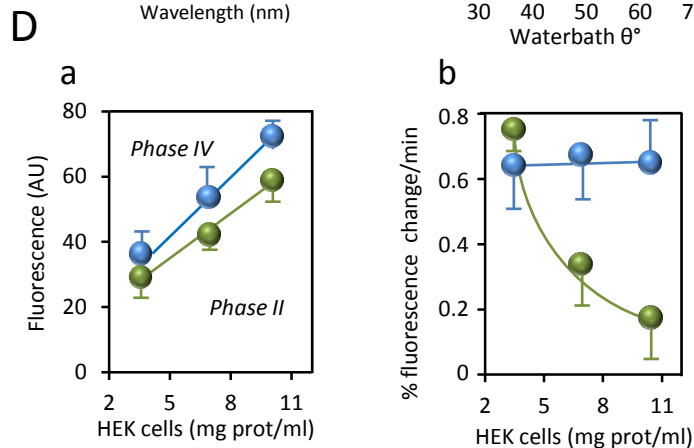
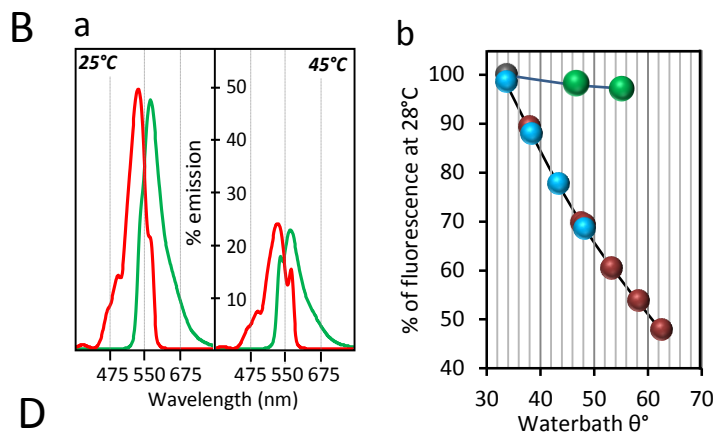
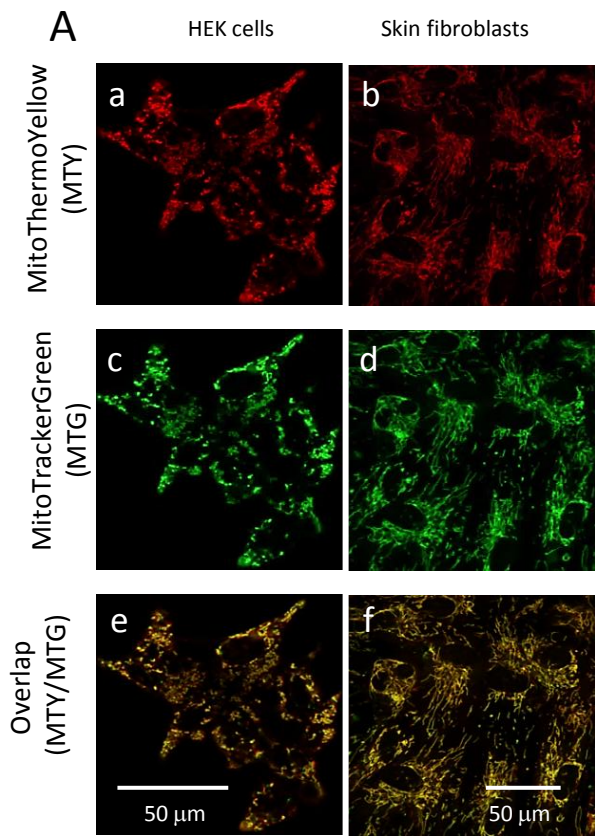
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351 **Author information**

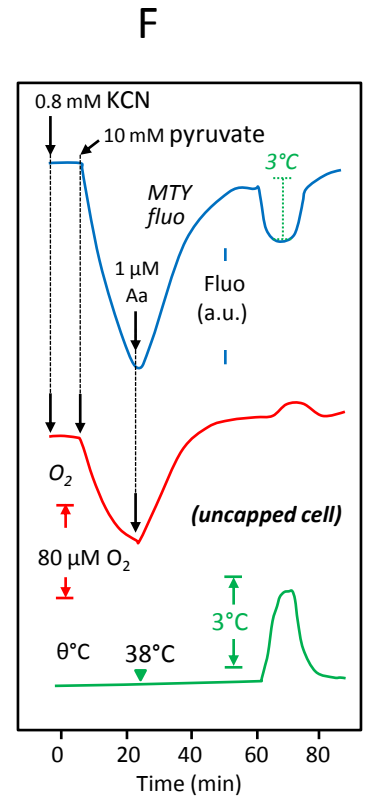
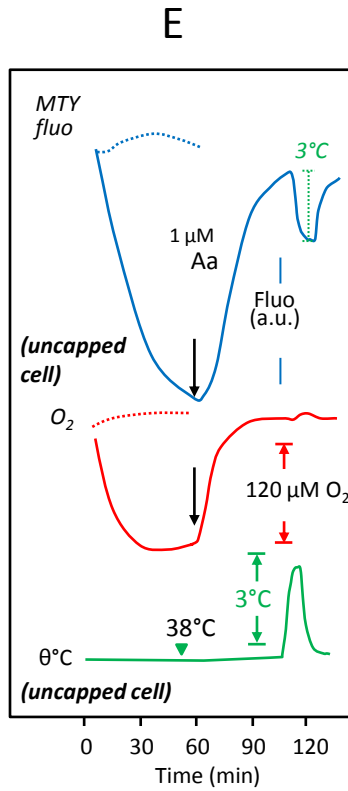
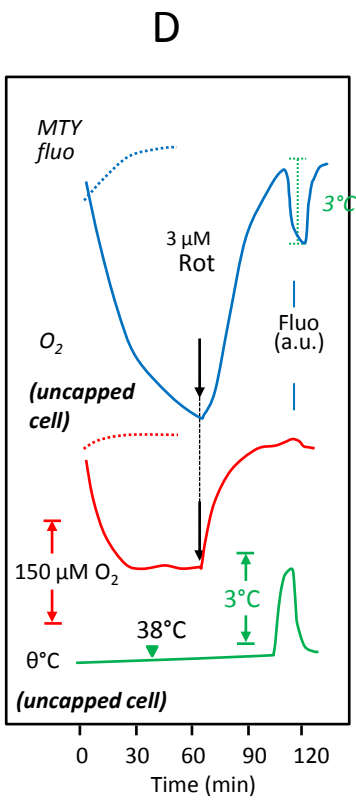
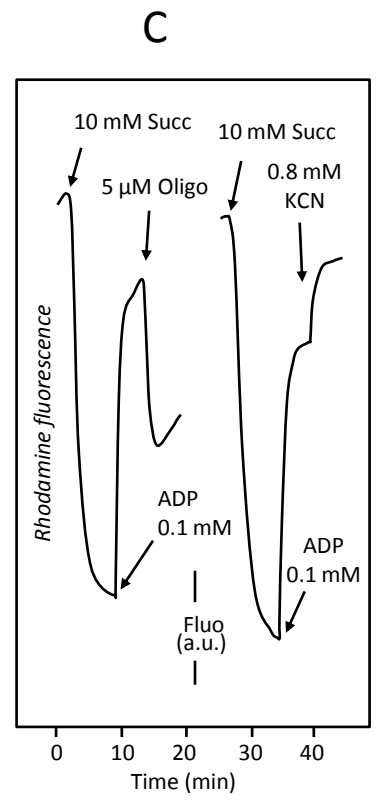
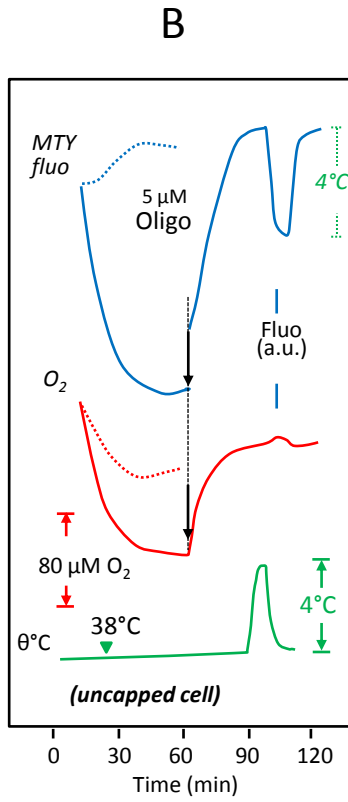
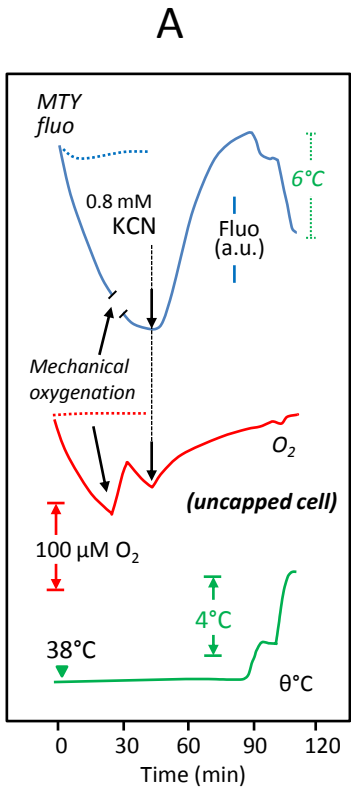
352 Reprints and permissions information is available at [www.nature.com/reprints](http://www.nature.com/reprints).

353 We confirm that there are no known conflicts or competing financial interests of interest  
354 associated with this publication and there has been no significant financial support for this  
355 work that could have influenced its outcome.

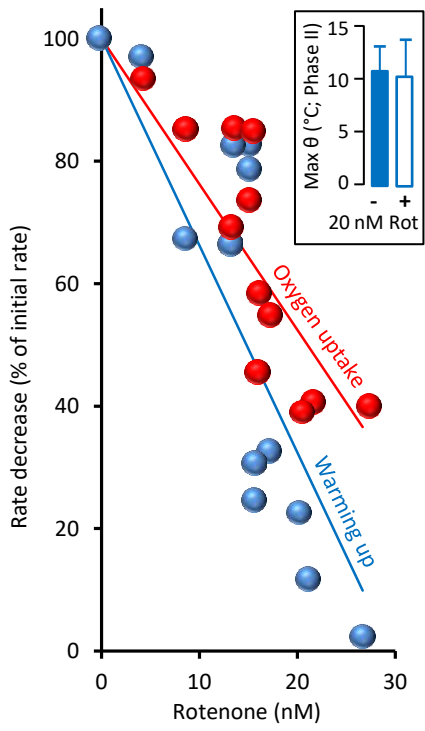
356 Correspondence and requests for materials should be addressed to Pierre Rustin,  
357 [pierre.rustin@inserm.fr](mailto:pierre.rustin@inserm.fr), or Malgorzata Rak, [malgorzata.rak@inserm.fr](mailto:malgorzata.rak@inserm.fr).



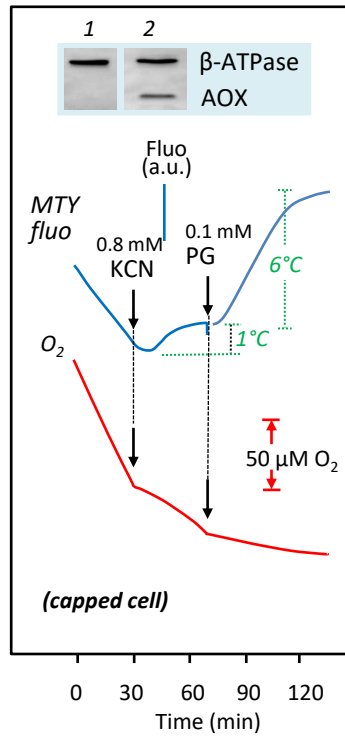




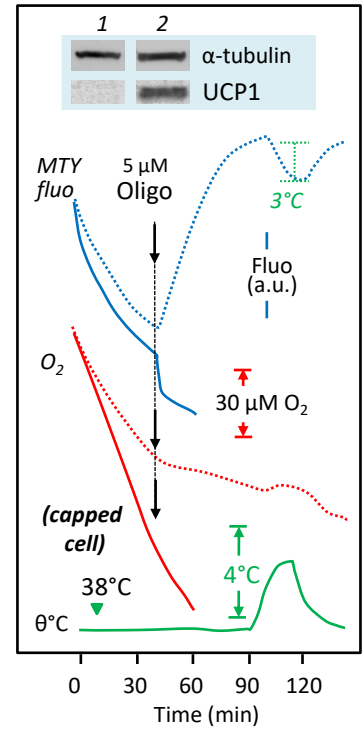
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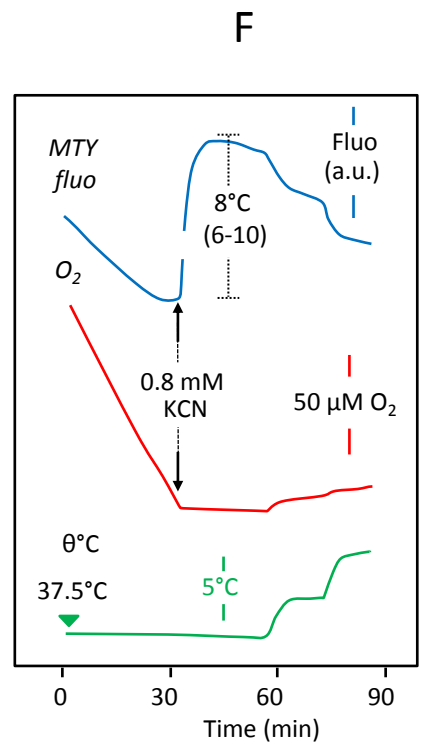
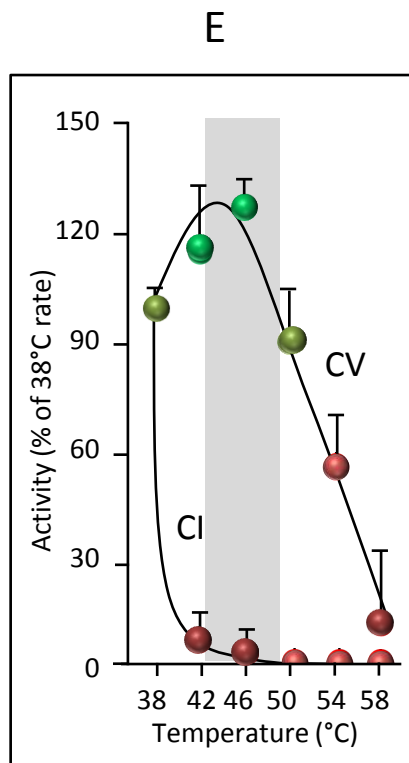
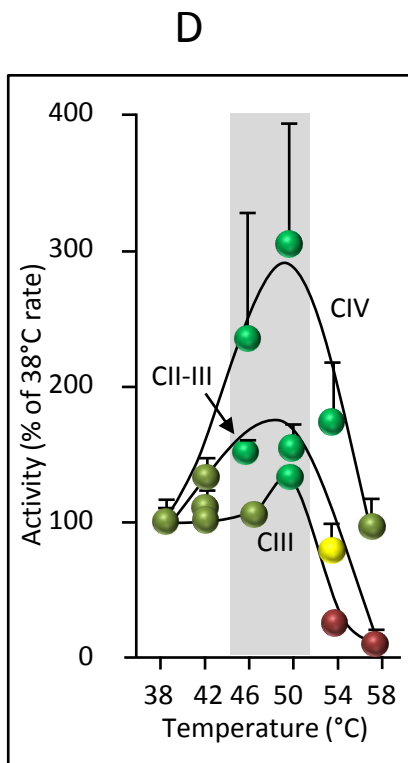
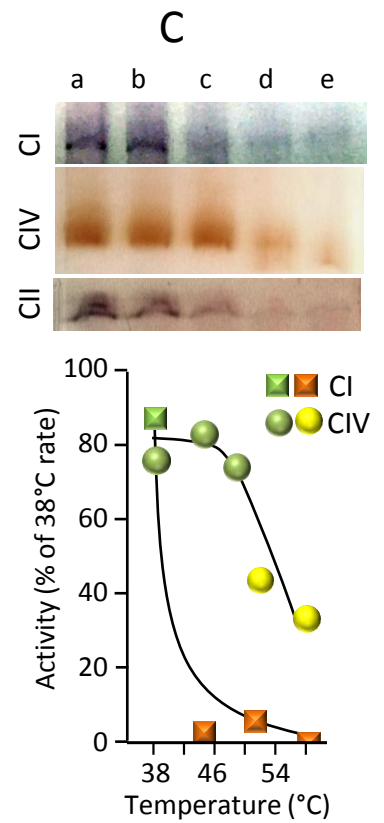
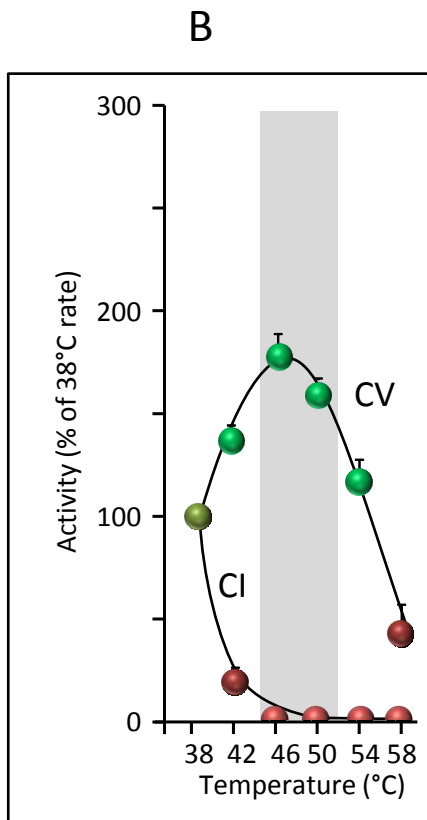
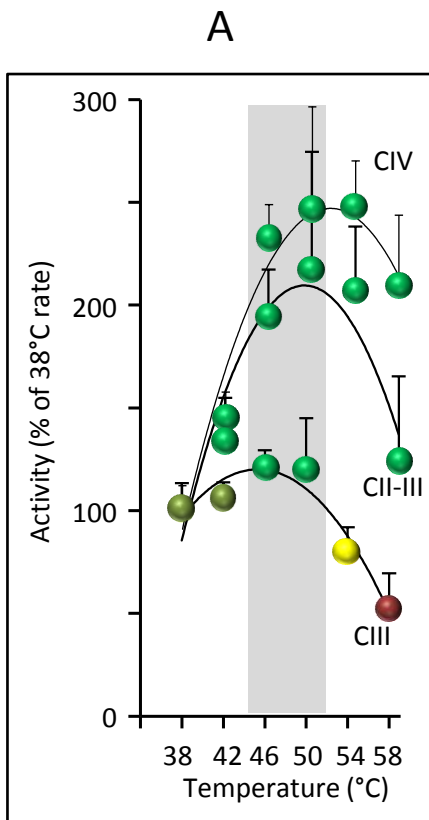


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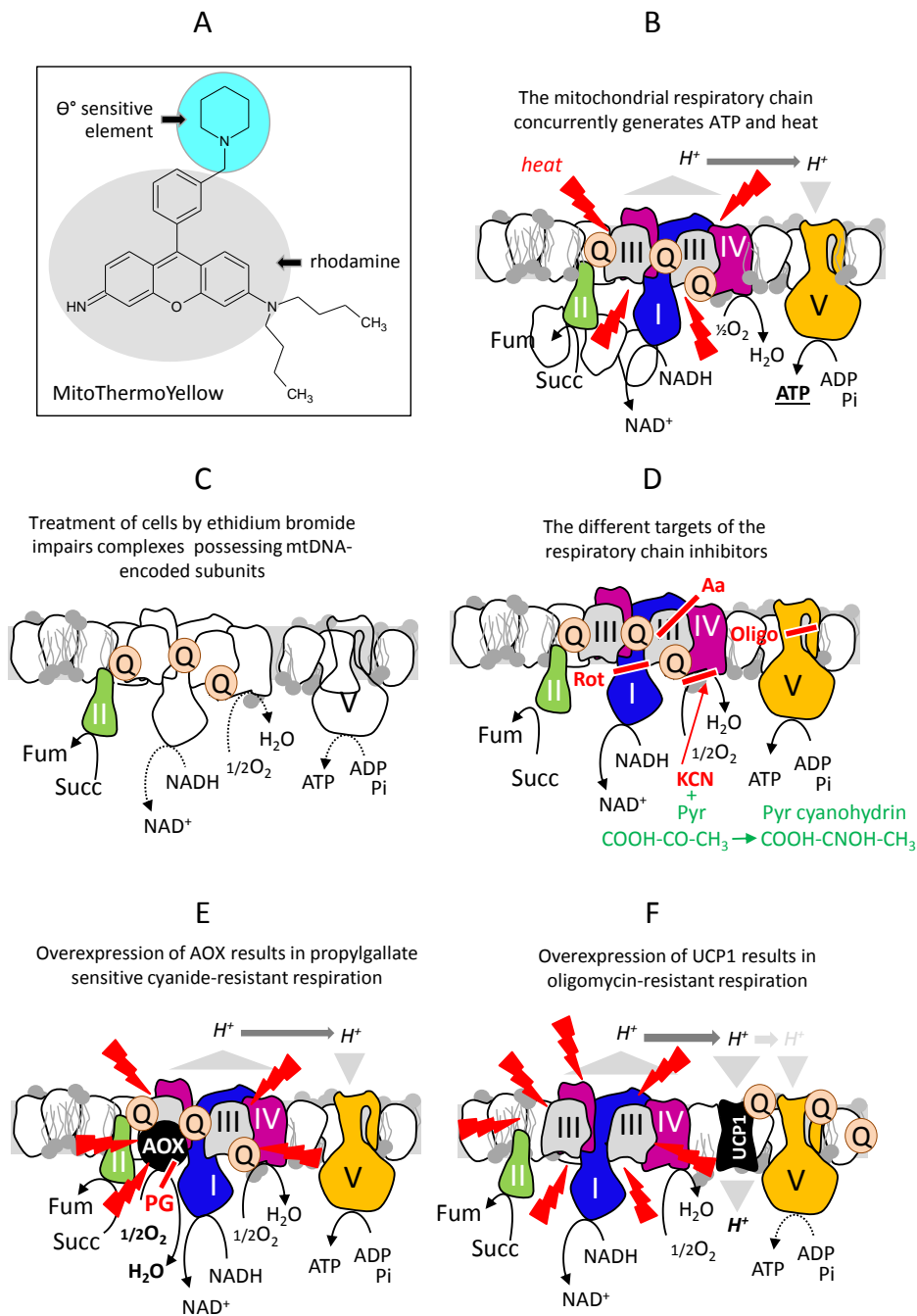
C





1 **Supplemental Material**

3 *Figures and their legends*



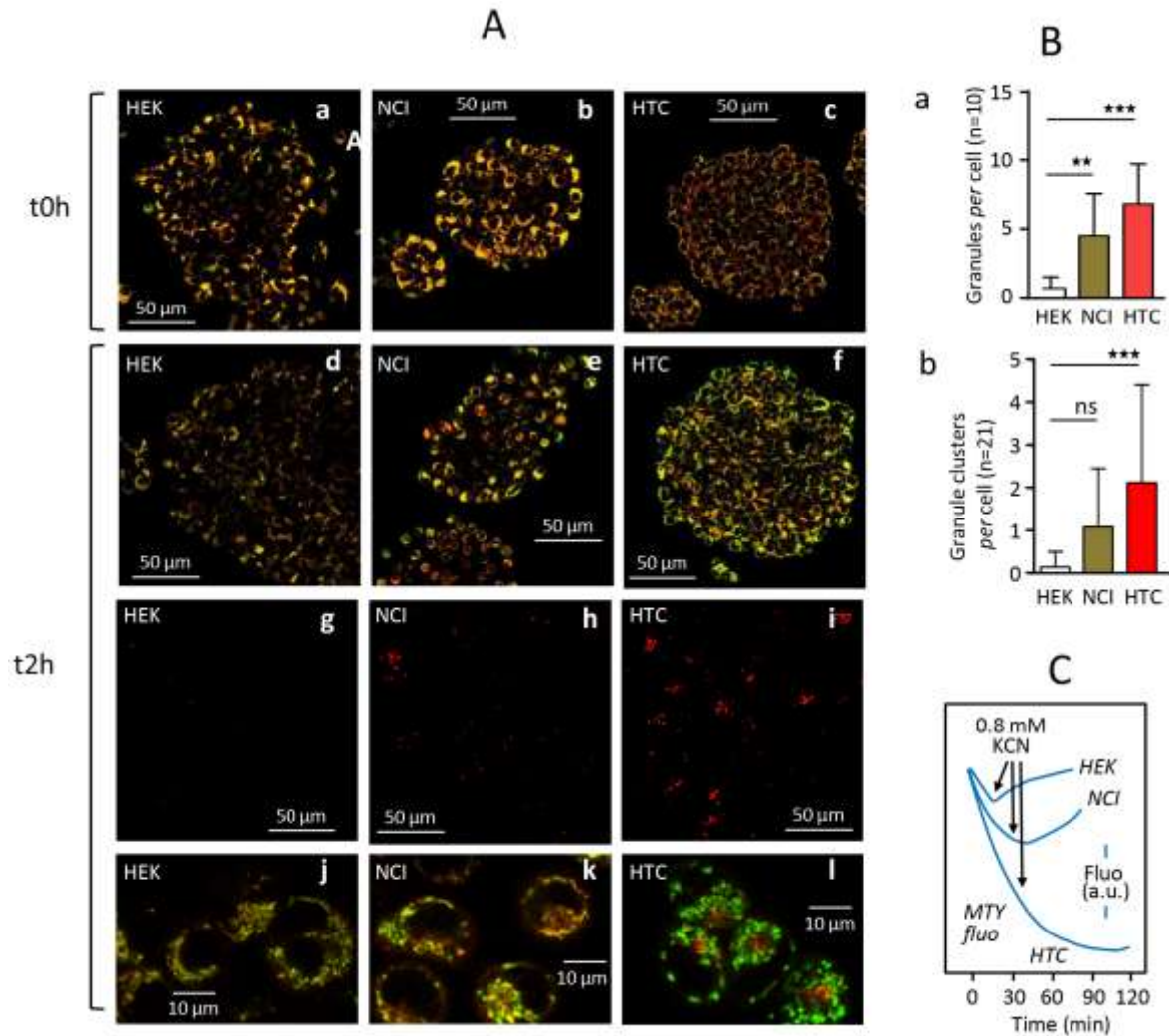
2

3 **Figure S1.** The structure of the MitoThermo Yellow probe and a series of schematized views  
 4 of the various conditions used for testing warming up of mitochondria *in situ* in cells.

5

6 A, The structure of the rhodamine-derived MTY probe. B, The concurrent synthesis of ATP  
 7 and heat generation by the respiratory chain. C, The defective respiratory chain of EtBr

8 treated cells. D, The sites of action of the several inhibitors used in this study. The respiratory  
 9 chain of AOX- (E) or UCP1- (F) expressing HEK cells. I, II, III, IV, V, the various complexes  
 10 of the respiratory chain; Aa, antimycin A; AOX, Alternative oxidase; Fum, fumarate; Oligo,  
 11 oligomycin; Q, ubiquinone 10 (coenzyme Q); Rot, rotenone; Succ, succinate; UCP1,  
 12 uncoupling protein 1.



13

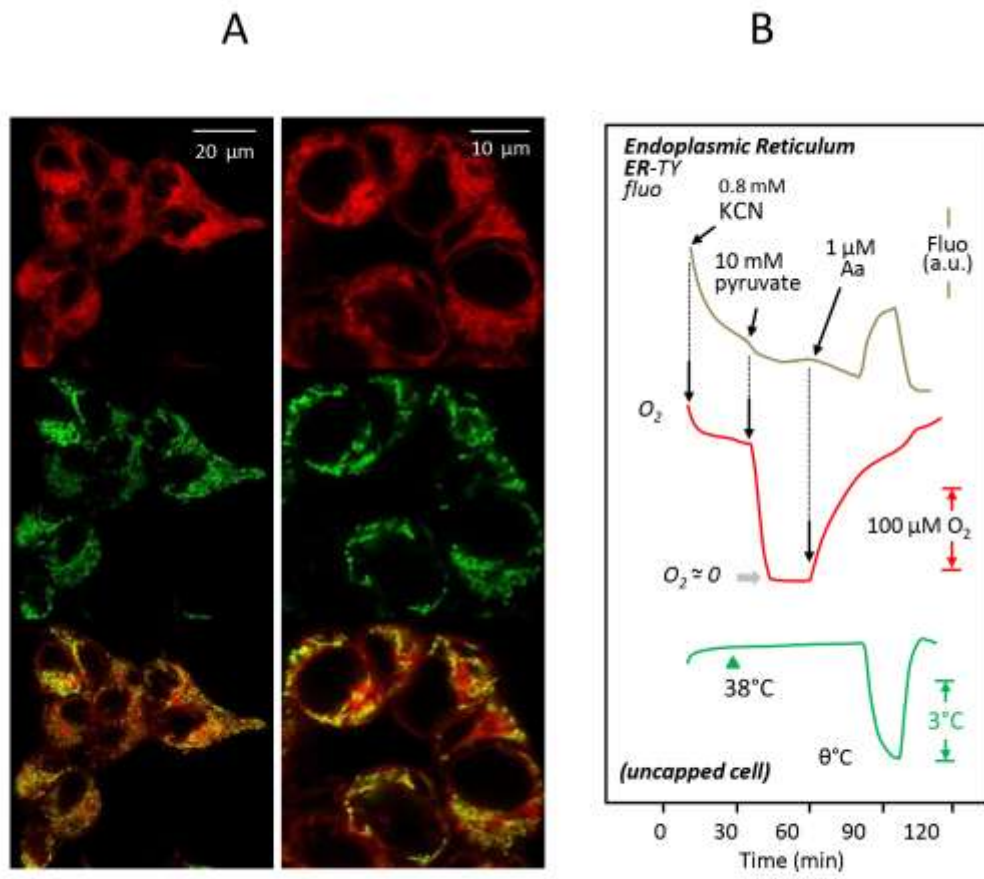
14 **Figure S2.** MTY probe well preserved *in situ* in HEK mitochondria is slowly (NCI cells) or  
 15 rapidly (HTC cells) expelled from mitochondria.

16

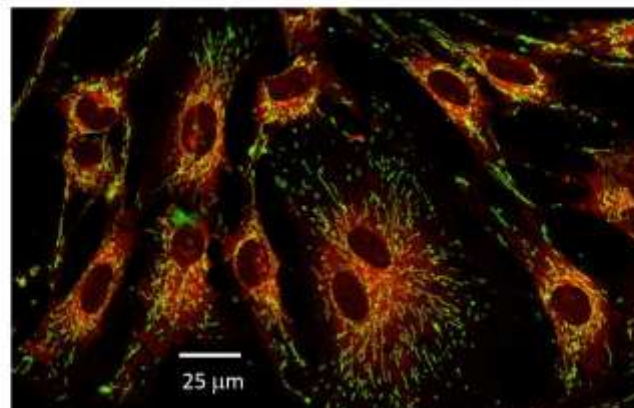
17 A, Initial MTY fluorescence is mostly localized to mitochondria in HEK (a), NCI (b), and  
 18 HTC (c) cell lines as shown by the overlapping staining of MTY and MitoTracker green  
 19 (yellow color). After 2 hours, a significant amount of the probe is excluded from  
 20 mitochondria in NCI cells (e), resulting in a number of cells being green or red colored.

21 Noticeably red (MTY) fluorescence is observed in cytosolic small granules (h). A similar but

22 much more pronounced phenomenon is observed in HTC cells where large granules can be  
23 observed (f, i, j). B, Quantification of MTY-stained (red) granules (a) and clustered-granules  
24 (b) in HEK, NCI and HTC cells. C, MTY-fluorescence changes (as in Fig. 2A) in HEK, NCI  
25 and HTC cells upon shift from anaerobic to aerobic conditions and the effect of a subsequent  
26 addition of cyanide. Noticeably while cyanide restores the initial fluorescence value in HEK  
27 cells it does not in NCI and even less in HTC cells. All together these experiments indicated  
28 that depending on cell types MTY can be either preserved for long (2 hours) in mitochondria  
29 (HEK cells) or more or less rapidly excluded as cytosolic granules (NCI, HTC cells) causing  
30 an irreversible loss of MTY fluorescence as measured in the spectrofluorometer quartz  
31 cuvette (C).  
32



**C**



34

35 **Figure S3.** Modulating respiratory chain activity does not change the fluorescence of an  
36 endoplasmic reticulum-targeted (ER thermos yellow; ERTY) version of MTY.

37

38 A, ERTY- (red) and MitoTracker green (green) fluorescence did not overlap (bottom) in HEK  
39 cells. B, Tested as MTY in the spectrofluorometer (Fig. 1), the fluorescence of ERTY within  
40 HEK cells (brown line) was unaffected by the activity of the mitochondria modulated by  
41 cyanide, pyruvate or antimycin, chemicals that fully controlled oxygen uptake (red line).  
42 Noticeably, fluorescence decrease of ERTY was similar in the initial absence or presence of  
43 cyanide (not shown). C, ERTY- (red) and MitoTracker green (green) fluorescence did not  
44 overlap either in skin fibroblasts (C) at variance with MTY and MitoTracker green which  
45 staining perfectly overlaps in the same cells (Fig. 1Af).

46