UNCOUPLING PROTEIN 1 DECREASES SUPEROXIDE PRODUCTION IN BROWN ADIPOSE TISSUE MITOCHONDRIA

Rebecca Oelkrug, Maria Kutschke, Carola W. Meyer, Gerhard Heldmaier, Martin Jastroch
Department of Animal Physiology, Faculty of Biology, Philipps-Universität Marburg, Karl-von-Frisch-Straße 8, 35043 Marburg, Germany
present address: Buck Institute for Age Research, 8001 Redwood Blvd, Novato, CA 94945, USA
Address correspondence to: Martin Jastroch, Buck Institute for Age Research, 8001 Redwood Blvd, Novato, CA 94945, USA. Fax: 1-415-209-2232; E-mail: mjastroch@buckinstitute.org

In thermogenic brown adipose tissue uncoupling protein 1 (UCP1) potently catalyzes the dissipation of mitochondrial proton motive force as heat. In a cellular environment of high oxidative capacity such as BAT, mitochondrial uncoupling could also reduce deleterious reactive oxygen species (ROS), but the specific involvement of UCP1 in this process is disputed. By comparing brown adipose tissue mitochondria of wildtype mice and UCP1-ablated litter mates, we show that UCP1 potently reduces mitochondrial superoxide production after cold acclimation and during fatty acid oxidation. We address the sites of superoxide production and suggest diminished probability of “reverse electron transport” facilitated by uncoupled respiration as the underlying mechanism of ROS suppression in BAT. Furthermore, ablation of UCP1 represses the cold-stimulated increase of substrate oxidation normally seen in active BAT, resulting in lower superoxide production, presumably avoiding deleterious oxidative damage. We conclude that UCP1 allows high oxidative capacity without promoting oxidative damage by simultaneously lowering superoxide production.

The physiological role of endogenous heat production by non-shivering thermogenesis in brown adipose tissue (BAT) of mammals, in particular small mammals, hibernators and neonates, is the maintenance of a constant high body temperature (T_b) independent of daily and seasonal temperature fluctuations (1). Heat production by BAT is, however, also activated by high calorie diets (2) and since presence of BAT has been recently shown in adult humans, it is in the focus as a potential therapeutic target for the treatment of human body weight disorders (3-5).

BAT is characterized by an exceptionally high presence of mitochondria which possess only a minor amount of ATP synthase to convert nutritional into cellular energy equivalents (6). Instead, brown adipose tissue mitochondria are specialized by a high content of mitochondrial uncoupling protein 1 (UCP1), up to 8% of total mitochondrial protein (7), which generates a proton leak in the mitochondrial inner membrane dissipating proton motive force as heat (8). UCP1 can be inhibited by physiological concentrations of purine nucleoside di- and triphosphates before free fatty acids can overcome inhibition (8).

Cold exposure stimulates lipolysis in BAT therefore activating UCP1. Long-term cold acclimation further increases the thermogenic capacity of BAT by enhanced UCP1 expression and recruitment of oxidative capacity (9). The importance of UCP1 for brown adipose tissue derived heat production has been confirmed in UCP1-ablated mice which are unable to defend their body temperature when acutely exposed to the cold (4°C) (10). Successive acclimation improves cold tolerance of UCP1-ablated mice, suggesting that other sources of heat partially compensate for the loss of functional BAT, namely muscle shivering (11, 12) and accelerated metabolic flux in white adipose tissue (13, 14). These alternative thermogenic mechanisms appear to be less efficient and are disadvantageous as the median survival rate of UCP1-ablated mice decreases in the cold from >24 weeks to ~13 weeks as compared to wildtype mice, raising the question about the advantages of UCP1-mediated thermogenesis.
The cold-recruitment of oxidative capacity increases the chance of oxidative damage. According to the free-radical theory of aging, an increase of radical production could potentially decrease lifespan (15, 16). Superoxide and reactive oxygen species (ROS) can be potently caused by mitochondria: escaping electrons from respiratory chain complexes are the major source of cellular superoxide and its derived reactive oxygen species (17, 18). Complex I (NADH:ubiquinone oxidoreductase; 19) and complex III (ubiquinol:cytochrome c oxidoreductase; 20) are postulated as the major producers of cellular superoxide, but also the mitochondrial glycerophosphate dehydrogenase (G3PDH), which is highly abundant in BAT (21, 22), appears to produce superoxide (23-25).

Functional studies on UCPs (UCP1, 2 and 3) showed activation of uncoupling activity by superoxide and peroxidation metabolites like 4-hydroxy-2-nonenal (26, 27), associating UCPs with a potential role in the prevention of superoxide production. These observations were highly disputed (28-30) and specifically the involvement of UCP1 in cellular reactive oxygen species regulation is questioned (29, 30). Studies comparing BAT from wildtype and UCP1-ablated mice found no activation of UCP1 with 4-hydroxy-2-nonenal (29, 30). Together with unchanged superoxide scavenging systems (29), these results led to the conclusion that conditions of high oxygen tension and high substrate supply in brown adipose tissue do not necessarily lead to increased oxidative stress.

Using mitochondria sampled from wildtype and UCP1-ablated mice, we here show that the presence of UCP1 potently lowers mitochondrial superoxide production during cold acclimation and fatty acid oxidation. Our results therefore demonstrate that uncoupling is a physiologically relevant mechanism to reduce oxidative stress in a mammalian organ of high oxidative capacity.

**Experimental procedures**

**Animals** - Wildtype and UCP1-ablated mice were provided by Dr. Leslie Kozak (Pennington Medical Research Center) and littermates (genetic background C57Bl/6J) were derived from heterozygous breeding pairs in the animal facility of the Philipps-Universität of Marburg. In these mice UCP1 was inactivated by homologous recombination with a deletion vector in which exon 2 and parts of exon 3 had been replaced with a neomycin resistance gene. In brown adipose tissue of these mice, no UCP1 could be detected with polyclonal antibodies (10). For experimental procedures, only homozygotes for UCP1 (wildtype; UCP1+/+) and UCP1-ablated (UCP1−/) mice were used. Mice were fed ad libitum (Sniff 1534) with free access to water and kept on a 12:12h light:dark cycle. For experiments, animals were kept in single cages. Warm acclimated mice (WA) were maintained at 30°C at least 3 weeks prior to sacrifice, while wildtype and UCP1-ablated adult mice were cold acclimated by first transferring them to 18°C for 3 weeks followed by 3 weeks at 5°C (11). The final body mass of the mice did not differ between the experimental groups (WA: UCP1+/+: 23.5±1.0g, UCP1−/+: 24.5±1.0g; CA: UCP1+/+: 24.7±0.3g, UCP1−/+: 24.4±0.4g). All experimental procedures were approved by the German Animal Welfare Authorities.

**Genotyping** - Mice were genotyped by amplifying a 201 bp (UCP1+/+) and 409 bp (UCP1−/) fragment from the UCP1 gene, using the primers 8265-5F: GGT AGT ATG CAA GAG AGG TGT and E2Rev: CCT AAT GGT ACT GGA AGC CTG and NeoRev: CCT ACC CGC TTG CAT TGC TCA, according to a protocol kindly provided by L. Kozak. Additionally the presence or absence of UCP1 protein was validated post mortem by immunological detection in brown adipose tissue mitochondria (as published previously 31). Membranes were probed with a rabbit anti-UCP1 polyclonal antibody (1:30,000 dilution; 3046, CHEMICON), followed by the relevant peroxidase-conjugated secondary antibody (goat-anti-rabbit-IgG at 1:10,000 dilution; Dako). Antigens were visualised on X-Ray Film (Super RX, Fuji) using an ECL Plus Western Blotting Detection system (SRX -101A, Konika Minolta).

**Mitochondria isolation** - Mitochondria were prepared by homogenisation and differential centrifugation as described previously (32). Mitochondria from one wildtype and one UCP1-ablated mouse were always isolated in parallel to control for possible day-by-day variability of mitochondrial quality. All available brown
adipose tissue depots were removed (interscapular, dorsal cervical, axillar, suprasternal, subcostal) and immediately placed in ice-cold isolation medium. Protein concentration was determined photometrically using the Bradford method (33).

Measurement of oxygen consumption
Mitochondrial oxygen consumption was measured using a Clark-type oxygen electrode (Rank Brothers Ltd, United Kingdom) maintained at 37°C and calibrated with air-saturated medium (50 mM KCl, 5 mM TES, 2 mM MgCl₂ x 6H₂O, 4 mM KH₂PO₄, 1 mM EGTA, BSA 0.4% (w/v), pH 7.2 at RT) which was assumed to contain 406 nmol O*ml⁻¹ (34). Mitochondrial respiration was measured in 500 µl medium at a concentration of 0.1 mg*ml⁻¹ incubated with 4.8 µM rotenone (inhibiting complex I of the respiratory chain) and 2 µM oligomycin (inhibiting the ATP synthase). Mitochondria were energized by addition of glycerol-3-phosphate (G3P, titrating up to 16 mM) or 4 mM succinate (state 4 respiration). Subsequently GDP was titrated to a final concentration of 5 mM. The ability of GDP to establish respiratory control in wildtype mice also confirmed the integrity of the mitochondria after the isolation process. At the end of each respiration measurement, the artificial uncoupler FCCP was titrated in ~1µM steps to estimate the maximal substrate oxidation rate. State 4 respiration in some cases was directly translated into proton leak rate assuming that 6 protons were transported (and leaked back to the matrix) per atomic oxygen when respiring on succinate or glycerol-3-phosphate.

Measurements of mitochondrial hydrogen peroxide release - Measurements of hydrogen peroxide production of isolated mitochondria were performed similar to Lambert (35). 10-20µg brown adipose tissue mitochondria were incubated in assay buffer (50 mM KCl, 5 mM TES, 2 mM MgCl₂ x 6H₂O, 4 mM KH₂PO₄, 1 mM EGTA, BSA 0.4% (w/v), pH 7.2 at RT) containing a cocktail of the fluorescent probe Amplex Red (50 µM, Invitrogen), 30 U*ml⁻¹ superoxide dismutase (to convert superoxide to hydrogen peroxide), 6 U*ml⁻¹ horseradish peroxidase (catalyzing the reaction of hydrogen peroxide with Amplex Red resulting in fluorescent resorufin) and 2 µM oligomycin (to inhibit ATP synthase). Amplex Red reacts with H₂O₂ at a 1:1 stoichiometry, while the stoichiometry of conversion from superoxide to H₂O₂ is assumed 1:2.

H₂O₂ formation was initiated by addition of glycerol-3-phosphate (15 mM), succinate (5 mM) or a cocktail of pyruvate (5 mM) and malate (3 mM). Experiments aimed to measure the H₂O₂ formation after palmitate addition were performed according to the protocol/substrates of the mitochondrial respiration measurements. Fluorescence was detected at 37°C in a microplate reader (BMG Labtech, FLUOstar OPTIMA) in 96 well microplates (GREINER 96-Well μClear, F-Bottom, black). The excitation wavelength was set to 560-10 nm, and the fluorescence emission was detected at 590 nm. Fluorescence was calibrated using known amounts of H₂O₂ at each experimental day. Optionally, superoxide production was measured in the presence of rotenone (2 µM, inhibiting complex I-derived ROS production), GDP (5 mM, to inhibit UCP1) and carboxyatractylate (CAT, 2.5 nM) to distinguish from adenine-nucleotide transporter-dependent effects.

Measurement of oxygen consumption and hydrogen peroxide release during beta oxidation of palmitoyl-CoA - To measure the palmitate-dependent mitochondrial respiration and hydrogen peroxide release, we incubated the mitochondria with 5 µM coenzyme A and 2 mM L-carnitine in a measuring buffer without BSA. Mitochondria were energized with 3 mM malate. After 7 minutes we added 1 mM ATP (to allow the activation of residual free fatty acids). After further 15 minutes 20 µM palmitate (equilibrated in a final concentration of 0.02% BSA) was added that activated beta oxidation and UCP1 in parallel (36).

Statistics - In figures and text, data are expressed as means±standard error (SE). In some cases, normal distribution of data could not be assumed. Therefore, in order to reveal significant differences in respiration rates or ROS production rates between genotypes, Student’s t-test or Mann Whitney U-test was used, as indicated. To assess the effect of different substrates or inhibitors within genotypes, paired t-test or Wilcoxon Wilcox Test was used. All calculations were performed using SigmaStat 3.5. The overall level of significance was set to p<0.05. To account for type one error in multiple comparisons effective p-values were adjusted according to Bonferroni.
RESULTS

In the following experiments, we investigated the impact of different substrates on mitochondrial respiration and superoxide production in isolated BAT mitochondria from warm 30°C (WA) and cold 5°C (CA) acclimated wildtype and UCP1-ablated mice. Figure 1 gives an overview on the respiratory chain complexes in the mitochondrial inner membrane, potential sites of substrate entry and electron leak, the latter resulting in superoxide production.

UCP1 reduces superoxide production in BAT mitochondria of warm acclimated mice

Initially, we investigated mitochondria respiring on glycerol-3-phosphate (G3P) as BAT has substantial amounts of glycerol-3-phosphate dehydrogenase (Figure 2a). As the affinity of the G3P dehydrogenase to its substrate is comparably low, we titrated G3P to saturating amounts (Supplemental Fig. 1). State 4 respiration in wildtype and UCP1-ablated mice up to 7 mM G3P was similar. More than 7 mM G3P resulted in a further increase of respiration rates of wildtype but not of UCP1-ablated mice, before saturating at concentrations above 13 mM (Figure 2a). Then, GDP was added to inhibit UCP1 and to induce respiratory control. The concentration of 5 mM GDP, that fully inhibited UCP1, was determined in mitochondria of CA wildtype animals (Supplemental Fig. 1). By addition of GDP and re-coupling of wildtype mitochondria, we also assured mitochondrial integrity (as brown adipose tissue mitochondria possess very little ATP synthase, state 3 respiration is not applicable to show respiratory control and oligomycin had minor or no effect on respiration). Addition of 5mM GDP strongly re-coupled wildtype mitochondria but had only minor effects in UCP1-ablated mitochondria (Figure 2a). At the end of each measurement, FCCP was added in order to induce maximal uncoupled respiration. FCCP-induced respiration was similar between wildtype and UCP1-ablated mitochondria.

We then measured the hydrogen peroxide release rate from isolated brown adipose tissue mitochondria which is established to report mitochondrial superoxide production rates (37, 38). Mitochondrial superoxide production rates were directly calculated from hydrogen peroxide concentrations. Superoxide production rates at state 4 (leak) respiration were lower in wildtype mitochondria than in UCP1-ablated mitochondria (Figure 2b). This reduction was attributable to UCP1 as the addition of the UCP1-inhibitor GDP abolished genotypic differences. The administration of rotenone allowed to determine the amount of superoxide production by the Iₜ site of complex I (Figure 1). G3P-dependent superoxide production was only slightly sensitive to rotenone (Figure 2b), indicating that the major proportion of superoxide was produced independently of complex I. Superoxide production rates and the associated effect of GDP were highly dependent on substrate concentration and only slightly sensitive to rotenone (Supplemental Fig. 1) supporting the idea that the G3P dehydrogenase itself generates superoxide. In particular, when aiming to compare superoxide levels from warm and cold acclimated animals, membrane potential and UCP1-sensitive effects on superoxide production may be overwritten by differences in G3P dehydrogenase content and its self-generated superoxide (25). Therefore, we chose to further investigate mitochondria respiring on succinate (Figure 2c), as the succinate dehydrogenase possesses high substrate affinity and is a weak producer of superoxide.

When respiring on succinate, leak respiration was about 3-fold higher in wildtype than in UCP1-ablated BAT mitochondria (Figure 2c). In wildtype mitochondria, leak respiration could be reduced to UCP1-ablated levels by administration of GDP. FCCP-respiration was indistinguishable between wildtype and UCP1-lacking mitochondria. Superoxide production rates of wildtype mitochondria respiring on succinate were barely detectable but GDP-sensitive, whereas in UCP1-ablated mitochondria, production rate was about 5 fold higher and GDP-insensitive (Figure 2b).

In the presence of GDP, superoxide production increased in wildtypes to levels as high as in UCP1-ablated mice which showed no GDP effect on superoxide production.

While superoxide production with G3P could only be slightly inhibited in the presence of rotenone, the potent inhibition in the presence of succinate demonstrated that the major proportion was produced by complex I (Figure 2d).
The sites of UCP1-dependent superoxide generation

The experimental setup measuring superoxide production with G3P and succinate, as well as the administration of rotenone, allowed for the determination of superoxide producing sites in BAT mitochondria. The rotenone-sensitive proportion of superoxide can be attributed to the I0 site of complex I. Superoxide generated by the G3P dehydrogenase can be calculated by subtracting the rotenone-insensitive superoxide production rate of succinate (residual superoxide) from the rate with G3P energization. Analyzing this (Figure 2e), UCP1 decreases superoxide production at the IQ site by about 40-50%, at the G3P dehydrogenase just about 20% and superoxide production at residual sites such as the Q pool and complex III by 50-70%.

Mitochondrial respiration and superoxide production during cold acclimation

While brown adipose tissue should be inactive in the thermoneutral zone (30°C) of the mouse, cold temperatures should lead to proliferation and activation of brown adipose tissue to produce heat. To examine this physiological condition we acclimated the two genotypes (UCP1+/+; UCP1−/−) to 5°C as described in the Material and Methods section. We then investigated thermoregulatory effects on respiration and superoxide production of isolated brown adipose tissue mitochondria.

Cold acclimation caused an increase in UCP1 protein levels (Figure 3a) and in leak respiration in wildtype mitochondria (compare Figure 2c with Figure 3b) but had no increasing effect in UCP1-ablated mice.

The genotype difference in succinate respiration observed in WA animals was further increased in response to cold acclimation (Figure 3b). Surprisingly, state 4 respiration in the CA UCP1-ablated mice (Figure 3b) was decreased in contrast to WA UCP1-ablated mice (Figure 2c), while in wildtype mitochondria, cold acclimation led to an increase of state 4 respiration (Figure 3b). Addition of 5 mM GDP inhibited UCP1 and decreased state 4 respiration of cold acclimated wildtype mice mitochondria (Figure 3b). Similar to WA animals, GDP had only minor effects on UCP1-lacking mitochondria (Figure 3b). Maximal substrate oxidation rate of CA UCP1-ablated mice determined with FCCP was not increased, suggesting that their mitochondria were unable to recruit oxidative capacity in the cold. The state 4 superoxide production rate of BAT mitochondria from CA animals was about 7.5 fold lower in wildtype than in UCP1-ablated mitochondria (Figure 3d). Addition of GDP increased superoxide production of wildtype mitochondria by 30-fold, exceeding values found for UCP1-ablated mice.

UCP1-sensitive changes in proton leak rate

Mitochondrial oxygen consumption in state 4 can be directly translated to proton leak rate assuming that with complex II substrates, six protons are transported out of the matrix (and leak back) per reduced oxygen atom. Calculated from Figure 2c, GDP led to a decrease of proton leak rate by 1.00 µmol protons *min⁻¹*mg⁻¹ protein in WA wildtype, but only by 0.18 µmol protons *min⁻¹*mg⁻¹ mitochondrial protein in UCP1-ablated mice (UCP1-independent effects). The UCP1-dependent proportion of proton leak rate of mitochondria respiring on succinate in state 4 can therefore be given as 0.82 µmol protons*min⁻¹*mg⁻¹ (Figure 3c). To investigate the physiological relevance of UCP1-mediated proton leak during cold acclimation, we also determined the UCP1-dependent proportion of proton leak from CA wildtype brown adipose tissue mitochondria (Figure 3b), which was about 2.74 µmol protons*min⁻¹*mg⁻¹ and about 0.30 µmol protons*min⁻¹*mg⁻¹ for UCP1-lacking mitochondria. Consequently 2.44 µmol protons*min⁻¹*mg⁻¹ of the proton leak was GDP-dependent. Taken together, the UCP1-dependent proton turnover rate in response to cold was increased by about 1.6 µmol protons*min⁻¹*mg⁻¹ or 300% (Figure 3c).

UCP1-dependent changes in superoxide production

What is the contribution of UCP1 in reducing mitochondrial superoxide? Inhibiting UCP1 with GDP should give a differential value of superoxide production rate that reflects the production rate depending on UCP1. This value can be further corrected with the effect of GDP in the UCP1-ablated mice. The calculation of the UCP1-dependent superoxide production (Figure 3d) showed that under warm conditions UCP1 blunts 83% of potential mitochondrial superoxide production. During cold stress, when respiration rate and the flow of electrons through the respiratory chain are elevated, the protective
function of UCP1 increased and mitigated more than 96% of potential superoxide production (calculated as difference of values ±GDP).

Other potential sources of uncoupling affecting mitochondrial superoxide production

The adenine nucleotide translocase (ANT) contributes significantly to basal proton leak (39) and can also regulate uncoupling (40-42), albeit less than UCP(s).

In BAT mitochondria of our experiments, the ANT, inhibited with carboxyatractylate (CAT), did not affect superoxide production rates (Supplemental Fig. S2), although it has been shown previously that isoforms may contribute to uncoupling in BAT mitochondria (29).

Superoxide production with the complex I substrates pyruvate/malate

Using pyruvate and malate as substrate of BAT mitochondria from both CA genotypes, we found barely detectable superoxide production rates (Figure 4) which were not increased by GDP. Substantial superoxide production could only be provoked by adding rotenone (produced at the I_f site, see Figure 1) and was higher in mitochondria of CA wildtype compared to UCP1-ablated mice (Figure 4).

UCP1 mitigates mitochondrial superoxide during fatty acid oxidation

During adaptive non-shivering thermogenesis, triglycerides are broken down by lipolytic activity and fatty acids are delivered to brown adipose tissue mitochondria. Free fatty acids activate UCP1, but in the presence of coenzyme A (CoA) and ATP, they are subsequently converted to acyl-CoA clearing this physiological UCP1 activator. To highlight the relevance of UCP1 in preventing superoxide production in a physiological scenario by mimicking cold stimulated lipolysis, we incubated brown adipose tissue mitochondria from cold acclimated mice with compounds of the carnitine cycle (L-carnitine, CoA), let them respire on malate to provide C4-bodies for beta oxidation, and then added ATP to activate and oxidize residual free fatty acids as described previously (43, Figure 5). Addition of ATP increased mitochondrial respiration in all mice (Figure 5a/b), clearly demonstrating that fatty acids were oxidized despite absence or partial purine-nucleotide related inhibition of UCP1.

After subsequent addition of a 20 µM pulse of equilibrated palmitate (containing 512 nM free fatty acids according to 44), the initial respiratory burst appeared to be larger in wildtype mice. This increase of respiration is likely caused by a combination of palmitoyl-CoA oxidation and uncoupled respiration, and was, however, depressed in UCP1-ablated mitochondria.

While superoxide production rates of mitochondria from both genotypes energized with malate were similar, the activation of endogenous free fatty acids induced by adding 1 mM ATP resulted in more pronounced increase of superoxide production of UCP1-ablated mitochondria as compared to wildtype mitochondria (Figure 5 c-f). Although the following palmitate pulse induced a stronger respiration rate of CA wildtype mitochondria, the superoxide production rate remained on a constant level, while it further increased in UCP1-ablated mitochondria.

DISCUSSION

Cold stress causes the activation of brown adipose tissue mitochondria, which quantitatively increases the influx of electrons, the reduction state of the respiratory chain and the amount of respiratory complexes. This is seen in our experiments by an increase of UCP1-dependent proton leak rate (Figure 3c). The probability of electrons to leak from the respiratory chain to form superoxide should increase, but our results show that this increase is blunted by UCP1. Major sites of superoxide production that are affected by the presence of UCP1 are the I_f site (about 40-50% lower) and other sites in the respiratory chain like the Q pool or complex III (about 50-70% lower).

In contrast, a previous study (29) concluded no effect of UCP1 on superoxide production, but measured rates in the presence of rotenone (inhibiting complex I) and antimycin A (inhibiting complex III). Under these conditions, however, there is reduced electron flow and membrane potential is negligible (see Figure 1). Therefore, no effect of UCP1 under those conditions is expected.

We show that the G3P dehydrogenase, which is highly abundant in brown adipose tissue, produces a major proportion of superoxide which can only be mildly reduced (about 20%) by uncoupling. Previous reports confirm that the
G3P dehydrogenase per se generates high levels of superoxide under in vitro conditions (24). The oxidative capacity of BAT mitochondria from warm acclimated animals, as measured by maximal uncoupling with FCCP, shows no genotype differences. However, during cold acclimation, it appears that only BAT of wildtype mice can derive the typical cold-induced oxidative capacity requiring the presence of UCP1.

In fact, superoxide production in UCP1-ablated mice is lower than expected from wildtype mitochondria with inhibited UCP1. The unexpectedly low superoxide production rates in cold acclimated UCP1-ablated mitochondria can be explained by compensatory lower respiratory chain activity. There is evidence for lower complex I content in UCP1-ablated mitochondria (seen in Figure 4), as the electron leak provoked by rotenone at the complex I flavin site is lower in UCP1-ablated than in wildtype mitochondria. Other downstream components of the respiratory chain are also affected as maximal succinate respiration in the presence of FCCP is also reduced in CA UCP1-ablated mitochondria (Figure 2b). The failure to increase oxidative capacity in UCP1-ablated mitochondria (as shown with pyruvate/malate) most of the superoxide during beta-oxidation in UCP1-ablated mice can be addressed to reverse electron transport. Generation of superoxide by ETF per se may be of minor significance as UCP1-ablated mitochondria have less respiration/oxidative capacity in the cold, but substantially produce more superoxide (Figure 5).

Our data support physiological significance of UCP1 in blunting enhanced superoxide production during beta-oxidation, the major pathway of substrate oxidation in brown adipose tissue during cold-induced thermogenesis.

Pertaining to superoxide and the lack of cold induced substrate oxidation in CA UCP1-ablated mice, superoxide may also act as a signalling molecule reporting the redox status of mitochondria thereby adjusting mitochondrial biogenesis, as recently hypothesized (45). It has been emphasized that some reactive oxygen species are required to induce mitochondrial biogenesis via PGC-1α (46) but our results suggests that pathologically high ROS concentrations inhibit respiration. Initial superoxide bursts during cold acclimation in UCP1-ablated mice may compromise the function of respiratory chain complexes, as seen for superoxide dismutase 2 (SOD2) knockout mice (47). Further indirect evidence that superoxide negatively controls oxidative capacity was derived from SOD2 overexpression which resulted in enhanced oxidative capacity (30).

Survival rates of UCP1-ablated mice are markedly reduced in the cold (11). Although local oxidative damage in BAT could be responsible for shorter lifespan, the absence of UCP1 requires recruitment of other tissues for heat production with less mitochondrial uncoupling capacity. This might increase systemic radical damage affecting lifespan.
Our results show the interrelationship of brown adipose tissue UCP1, energy dissipation and superoxide production, which may become instrumental when considering human brown adipose tissue for the treatment of obesity. A model proposing activation of UCP1 directly by superoxide as a negative feedback, thereby preventing further superoxide production (27), is highly disputed (28 - 30). Independent of such a mechanism, we here demonstrate that UCP1 regulates mitochondrial superoxide production upon physiological challenge. We comprise our results in a model shown in figure 6. Mammalian UCP1 allows high oxidation rates in the absence of deleterious oxidative stress and due to this dual role in heat production and superoxide reduction, evolution may have selected for the implementation of UCP1 in adaptive non-shivering thermogenesis of mammals. The role of recently discovered ancient UCP1 orthologues in ectotherms (48) may well be prevention of superoxide production prior to evolution of non-shivering thermogenesis.
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FOOTNOTES

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The abbreviations we used are: ANT, adenine nucleotide translocase; CA, cold acclimated; CAT, carboxyatractylate; FCCP, carbonyl cyanide p-trifluoromethoxyphenylhydrazone; UCP, uncoupling protein; WA, warm acclimated
FIGURE LEGENDS

Fig. 1: Sites of substrate entry and superoxide production in the respiratory chain of BAT mitochondria. NADH linked substrates donate their electrons to complex I. Further substrate entry sites, which are investigated in this study, are the mitochondrial glycerol-3-phosphate dehydrogenase (G3PDH), the succinate dehydrogenase (complex II) and the electron transfer flavoprotein (ETF), which is fuelled by beta oxidation respectively. Beta oxidation also generates NADH, which enters complex I in parallel. The electrons from the entry complexes are transferred to ubiquinone (Q) forming QH2. In the normal electron forward transfer, the electrons are donated to complex III, then complex IV, before finally reducing oxygen to water. Electrons may leak from the I_F site and the I_Q site of complex I. These sites can be distinguished by using the complex I inhibitor rotenone. Substantial amount of superoxide production can also derive from the G3PDH and the ETF. The sites of superoxide production are indicated with asterisks in the figure. Proton motive force, generated by complexes I, III and IV, is dissipated by either the ATP synthase (complex V) or UCP1. UCP1 is activated by free fatty acids and inhibited by purine nucleoside di- and triphosphates (here: GDP).

Fig. 2: Respiration and superoxide production rates of brown adipose tissue mitochondria from warm acclimated wildtype (UCP1+/+) and UCP1-ablated (UCP1−/−) mice energized with glycerol-3-phosphate (a, b) or with succinate (c, d). The effect of UCP1 on superoxide derived from different sites is illustrated in e. Mitochondrial respiration of 0.1 mg/ml mitochondrial protein was measured in a temperature controlled chamber with a Clark-type electrode at 37°C. 2 µM oligomycin, 4.8 µM rotenone and 16 mM G3P (a) or 4 mM succinate (c) were added to establish state 4 respiration, 5 mM GDP to inhibit UCP1 and re-couple mitochondria and at the end of each run, FCCP to artificially uncouple mitochondria and estimate maximum substrate oxidation. Comparing succinate with G3P respiration, succinate respiration was higher in every treatment. State 4 respiration with G3P tended to be higher in wildtype as compared to UCP1-ablated mitochondria (P=0.1, Student’s t-test, n=8 per group), but was significantly different during succinate respiration. (**P<0.001, Student’s t-test n=7 per group). Superoxide production was calculated from hydrogen peroxide release assayed with Amplex Red in the presence of superoxide dismutase (30 U/ml). Superoxide production was measured in mitochondria incubated with oligomycin (2 µM, State 4), 5 mM GDP was added to inhibit UCP1 and 4.8 µM rotenone to inhibit complex I. Genotype differences in superoxide production were found in state 4 and in the presence of rotenone (inhibiting complex I) (*P<0.05, Student’s t-test; n=7 per group). Superoxide production after addition of succinate (d) was generally lower but rotenone addition resulted in a greater decrease demonstrating that the major proportion of superoxide was produced at complex I (8P<0.05, Mann Whitney U-test, UCP1+/+ n=11 and UCP1−/− n=15 (12 with GDP)).(e) Rotenone inhibits superoxide production from the IQ-site of complex I.
superoxide leaving from the \( I_Q \)-site \((a', a)\) when respiring on glycerol-3-phosphate is about 40 to 50\% lower in brown adipose tissue mitochondria of wildtype mice as compared to \( UCP1 \)-ablated mice. 

Respiring on succinate with rotenone, the residual superoxide production rate \((c', c)\) derives from other sites such as the Q radical and complex III. This residual superoxide production rate is about 50\% lower in wildtypes. Subtracting \( c \) and \( c' \) from the rotenone insensitive rate with glycerol-3-phosphate determines superoxide generated by the glycerol-3-phosphate dehydrogenase \((b, b')\). This rate was about 20\% lower in \( UCP1 \)-wildtype mice. All experiments were performed from brown adipose tissue mitochondria of individual animals kept at 30°C.

**Fig. 3: Cold-induced adaptive changes in mitochondrial physiology.** Succinate-respiration rates/\( UCP1 \)-dependent proton leak rate and superoxide production rates/\( UCP1 \)-dependent superoxide production of brown adipose tissue mitochondria from cold acclimated wildtype (\( UCP1^{+/+} \)) and \( UCP1 \)-ablated (\( UCP1^{-/-} \)) mice. The ablation and cold-induced protein levels of \( UCP1 \) were confirmed by immunological detection of \( UCP1 \) in brown adipose tissue with an anti-hamster \( UCP1 \) antibody \((a)\). Experiments on mitochondrial respiration were performed on 0.1 mg mitochondrial protein per ml from wildtype \((n=7)\) and \( UCP1 \)-ablated mice mitochondria \((n=5)\) incubated in an air-saturated medium at 37°C \((b)\). Respiration was started in state 4 with 2µM oligomycin, 4.8µM rotenone and 4 mM succinate. Addition of GDP (5 mM) acted as an inhibitor of \( UCP1 \) and a final addition of 2 - 4 µM FCCP was made to compare maximum substrate oxidation rates. State 4 respiration of CA wildtype mitochondria was higher than of \( UCP1 \)-lacking mitochondria, and the difference persisted after GDP addition \((b)\) \((^P<0.05 \text{ Mann Whitney U-test})\). The \( UCP1 \)-dependent proton leak rate \((c)\) was calculated based on the assumption that 6 protons are transported per molecule oxygen and increased by 1.6 µmol*min^{-1}*mg^{-1} in response to cold acclimation. Data for WA mice were taken from Figure 1 \((***P<0.001, \text{ Student’s t-test})\). Superoxide production was assessed by measurement of the hydrogen peroxide release rate, determined fluorometrically over the reduction of Amplex Red to resorufin in the presence of superoxide dismutase \((30 \text{ U*ml}^{-1})\) and HRP \((6 \text{ U*ml}^{-1})\). d) In state 4 conditions \((\text{with 2 µM oligomycin})\) superoxide production was higher in \( UCP1 \)-ablated mitochondria \((n=5)\). Administration of GDP increased wildtype \((n=7)\) mitochondrial superoxide production, exceeding levels found for \( UCP1 \)-ablated mitochondria as well as WA wildtype mitochondria \((\text{see Fig. 1d})\) \((^P<0.05 \text{ Mann Whitney U-test})\). e) The \( UCP1 \)-dependent superoxide production rate was calculated as superoxide production after GDP inhibition minus the respective state 4 superoxide production, resulting in a 2.7 fold increase of \( UCP1 \)-dependent superoxide production of wildtype mitochondria in response to cold \((^P<0.05 \text{ Mann Whitney U-test})\).

**Fig. 4: Superoxide production of brown adipose tissue mitochondria from cold acclimated wildtype (\( UCP1^{+/+} \)) and \( UCP1 \)-ablated (\( UCP1^{-/-} \)) mice mitochondria by complex I**
A cocktail of pyruvate (5 mM) and malate (3 mM) was used to provide complex I with NADH. By supplemental addition of rotenone we showed that UCP1 provides superoxide production by reverse electron transfer. Independent experiments were performed with wildtype mitochondria (n=4) and UCP1-ablated mice mitochondria (n=8) ($^eP<0.05$ Mann Whitney U-test).

**Fig. 5:** Effects of UCP1 ablation on superoxide production during fatty acid oxidation in brown adipose tissue mitochondria of cold acclimated animals. a) Palmitoyl-CoA oxygen consumption response curve of a wildtype mouse and b) of a UCP1-ablated mouse. The arrows indicate the time points when substrates were added. c) Superoxide production of the same wildtype mouse as shown above measured as fluorescence and d) of the same UCP1-ablated mouse. At t=0 the mitochondria were respiring on malate (3 mM) in all performed measurements. The lower panels show the mean values±SE of wildtype mice (n=4) and UCP1-ablated mice (n=8) for every treatment (***,P<0.001, One Way RM ANOVA (Bonferroni-adjusted)).

**Fig. 6:** Physiological significance of mitochondrial uncoupling on mitochondrial superoxide production in brown adipose tissue mitochondria. Cold exposure activates brown adipose tissue for heat production which is generally referred as adaptive non-shivering thermogenesis. The heat output of the brown adipocytes is increased by stimulation of oxidative capacity and mitochondrial biogenesis. These factors would usually potently increase the probability of mitochondrial superoxide production. In BAT mitochondria, however, cold acclimation also stimulates UCP1 gene transcription and protein levels, and cold-induced free fatty acid release activates UCP1 which dissipates proton motive force as heat. Based on our study, we put forward a model that proposes an additional physiological role of UCP1 by preventing cellular damage through the decrease of superoxide formation.
Figures:
Figure 1
Figure 3

a) Western blot analysis showing UCP1+/- and UCP1-/- at 40 kDa and 30 kDa at 30°C and 5°C.

b) Mitochondrial respiration (nmol O2/min/mg protein).
   - State 4
   - + GDP
   - + FCCP

b) UCP1-dependent proton leak rate (nmol H+min/mg protein).
   - 30°C
   - 5°C

b) Super oxide production (nmol O2/min/mg protein).
   - State 4
   - + GDP
   - + ROT

b) UCP1-dependent super oxide production (nmol O2/min/mg protein).
   - 30°C
   - 5°C
Figure 4

Comparison of superoxide production in different states and treatments.

- UCP\textsuperscript{+/+}
- UCP\textsuperscript{--/--}

Parameters:
- Superoxide production (nmol O\textsubscript{2}^{-} \cdot min\textsuperscript{-1} \cdot mg protein\textsuperscript{-1})
- State 4
- + GDP
- + Rot

Significance:
- P = 0.09
- # indicates a significant difference
Figure 5

(a) 1mM ATP

(b) 1mM ATP

(c) Palmitate

(d) Palmitate

(e) 1mM ATP

(f) Palmitate

Superoxide production (mmol O$_2$·min$^{-1}$·mg protein$^{-1}$)

Molde  + 1mM ATP  + Palmitate

Superoxide production (mmol O$_2$·min$^{-1}$·mg protein$^{-1}$)

Molde  + 1mM ATP  + Palmitate

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Uncoupling protein 1 decreases superoxide production in brown adipose tissue mitochondria
Rebecca Oelkrug, Maria Kutschke, Carola W. Meyer, Gerhard Heldmaier and Martin Jastroch

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