

Mitochondrial energy utilization maintains young status in the trophocytes and oenocytes of old queen honeybees

Chin-Yuan Hsu^{1,2}, Cheng-Yen Lu²

¹Department of Biomedical Sciences, College of Medicine, Chang Gung University, 259, Wen-Hwa 1st Road Kwei-Shan, 333, Tao-Yuan, Taiwan

²Graduate Institute of Biomedical Sciences, College of Medicine, Chang Gung University, Tao-Yuan, Taiwan

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Abstract – The lifespans of queen honeybees (*Apis mellifera*) are much longer than those of worker bees. Mitochondrial energy utilization in the trophocytes and oenocytes of worker bees during aging has been determined, but it remains unknown in queen bees. In this study, mitochondrial energy utilization was assayed in the trophocytes and oenocytes of young and old queen bees. The mitochondrial density and mitochondrial membrane potential ($\Delta\psi_m$); nicotinamide adenine dinucleotide (NAD^+), nicotinamide adenine dinucleotide reduced form (NADH), and adenosine triphosphate (ATP) levels; NAD^+/NADH ratio; and relative expression of NADH dehydrogenase 1 (ND1) and ATP synthase normalized against mitochondrial density were not significantly different between young and old queen bees. These results indicate that mitochondrial energy utilization maintains a young status in the trophocytes and oenocytes of old queen bees and that trophocytes and oenocytes have longevity-promoting mechanisms that can be investigated to clarify the secret of longevity in queen bees.

aging / mitochondria / energy utilization / honeybee

Scientific abbreviations

$\Delta\psi_m$	Mitochondrial membrane potential
NAD^+	Nicotinamide adenine dinucleotide oxidized form
NADH	Nicotinamide adenine dinucleotide reduced form
ATP	Adenosine triphosphate
ND1	NADH dehydrogenase 1

1. INTRODUCTION

Understanding the mechanisms of aging and longevity could improve the health and extend the lifespan of many organisms, including humans. Honeybees (*Apis mellifera*) provide a unique model for studying aging and longevity because, although worker bees and queen bees develop from the same genome, queen bees have much

longer lifespans. The lifespans of worker bees are 15–38 days in the summer and 150–200 days in the winter, while queen bees live 1–2 years with a maximum of up to 8 years (Page and Peng 2001; Omholt and Amdam 2004; Rueppell et al. 2007; Remolina and Hughes 2008).

Large and irregularly shaped trophocytes attach to small and spherical oenocytes to form a single layer of cells around each segment of honeybee abdomen (Kuterbach and Walcott 1986; Martins et al. 2011). Lack of cell division during adulthood, ease of isolation from the abdomen, and convenient manipulation make trophocytes and oenocytes suitable for use in studies of cellular senescence and longevity (Hsieh and Hsu 2011a, b; Hsu and Chan 2013; Chan et al. 2011; Chuang and Hsu 2013; Hsu and Chuang 2014; Hsieh and Hsu 2013; Hsu and Hsieh 2014; Hsu and Hu 2014).

Studies of age-related molecules have shown that the trophocytes and oenocytes of old queen bees exhibit higher levels of senescence-

Corresponding author: C. Hsu, hsu@mail.cgu.edu.tw
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associated beta-galactosidase, more lipofuscin granules, and higher lipid peroxidation and protein oxidation than those of young queen bees (Hsieh and Hsu 2011b). These results are similar to those reported in worker bees (Hsieh and Hsu 2011a). However, queen bees have much longer lifespans than worker bees, suggesting that the trophocytes and oenocytes of queen bees have longevity-promoting mechanisms.

Studies of oxidative stress and antioxidant enzyme activity have shown that the trophocytes and oenocytes of old queen bees have higher levels of reactive oxygen species (ROS) and higher activity of superoxide dismutase, catalase, and glutathione peroxidase than those of young queen bees (Hsieh and Hsu 2013). These studies suggest that increased oxidative stress and consequent stress defense mechanisms are associated with the longevity of queen bees. In contrast, in worker bees, oxidative stress decreases in trophocytes and oenocytes with age (Hsu and Hsieh 2014).

The trophocytes and oenocytes of old queen bees have similar levels of AMPK activity; FoxO expression; adenosine triphosphate (ATP), ADP, AMP, and cAMP concentrations; SirT1 expression and activity; and PPAR- α expression as those of young queen bees (Hsu and Hu 2014). These results suggest that energy metabolism in trophocytes and oenocytes is similar between young and old queen bees and is different from that reported in worker bees (Hsu and Chuang 2014). These results are consistent with previous studies showing that FoxO overexpression extends the lifespan of *Drosophila* (Giannakou et al. 2004, Hwangbo et al. 2004). These results also suggest that the trophocytes and oenocytes of queen bees have longevity-promoting mechanisms because queen bees have longer lifespans than worker bees and old queen bees have FoxO expression similar to young queen bees.

Old queen bees have high accumulations of age-related molecules and high oxidative stress (Hsieh and Hsu 2011b; Hsieh and Hsu 2013). However, energy-regulated molecule expression is similar between young and old queen bees. To clarify these phenomena, we studied mitochondrial energy utilization in the trophocytes and

oenocytes of young and old queen bees. In the present study, mitochondrial density, $\Delta\Psi_m$, nicotinamide adenine dinucleotide (NAD^+) concentration, nicotinamide adenine dinucleotide reduced form (NADH) concentration, ATP concentration, NAD^+/NADH ratio, and NADH dehydrogenase 1 (ND1) and ATP synthase expression were evaluated in the trophocytes and oenocytes of young and old queen bees to understand mitochondrial energy utilization in queen bees.

2. MATERIALS AND METHODS

2.1. Queen honeybees

Queen bees were purchased from a single commercial breeder (Hsinchu, Taiwan). As described in our previous studies (Hsieh and Hsu 2011b; Hsieh and Hsu 2013; Hsu and Hu 2014), young (2-month-old) and old (16-month-old) queen bees were collected from different hives on the same dates for use in the following experiments. The young and old queen bees were mated with drones and were able to lay eggs. Individual queens of each group were dissected with scissors, and their abdominal trophocytes and oenocytes were detached from the cuticle using a knife in honeybee saline (Hsu et al. 2007). Trophocytes and oenocytes were used for the following experiments.

2.2. Assay of mitochondrial density

The mitochondrial density was measured as previously described using transmission electron microscopy (TEM) (Hsu and Chan 2011; Chuang and Hsu 2013). Eight microscopy images of the trophocytes near the nucleus from each young and each old queen were analyzed using Photoshop (CS6). This experiment was performed with five biological replicates using a total of five young and five old queen bees.

2.3. Assay of $\Delta\Psi_m$

$\Delta\Psi_m$ assays in the trophocytes and oenocytes were performed as previously described using the specific dye tetramethylrhodamine methyl ester (TMRM) (T668; Life Technologies, Grand Island, NY, USA) (Ward et al. 2000; Chuang and Hsu 2013). Briefly, trophocytes and oenocytes were stained with TMRM, washed with PBS, and visualized under a confocal

microscope (Leica TCS SP2; Leica, Wetzlar, Germany). $\Delta\Psi_m$ was expressed as the red fluorescence intensity/cell area ratio. Five confocal microscopy images of the trophocytes and oenocytes from each young and each old queen were analyzed using QWin image processing and analysis software (version 2.5, Leica, Wetzlar, Germany) and Photoshop (CS6). This experiment was performed with five biological replicates using a total of five young and five old queen bees.

2.4. Assay of NAD^+ and NADH concentration

NAD^+ and NADH were measured in the trophocytes and oenocytes as previously described using an NAD^+ /NADH Quantification Kit (K337-100; BioVision, Mountain View, CA, USA) (Chuang and Hsu 2013). Briefly, the trophocytes and oenocytes from one young or old queen bee were extracted with NADH/NAD extraction buffer; the extracted samples were heated for NADH measurement. After background luminescence was recorded, the extracted samples and the extracted and heated samples were measured. The resultant luminescence was normalized to background. Dilutions of an NADH standard solution (0, 20, 40, 60, 80, and 10 $\text{pmol } \mu\text{L}^{-1}$) were processed concurrently to generate a standard curve, which was used to obtain NAD^+ and NADH values. Protein concentration was determined using a commercial protein assay reagent (500–0006; Bio-Rad Laboratories, Hercules, CA, USA). The NAD^+ and NADH concentrations are expressed as micromoles per milligram protein and were used to determine the NAD^+ /NADH ratio. This experiment was performed with five biological replicates using a total of five young and five old queen bees.

2.5. Assay of ATP concentration

The ATP concentration in the trophocytes and oenocytes was determined as previously described using an ATP Determination Kit (A22066; Invitrogen, Carlsbad, CA, USA) (Chuang and Hsu 2013). Briefly, the trophocytes and oenocytes from one young or old queen bee were homogenized in phosphate buffer containing protease inhibitors and centrifuged to obtain the resulting supernatant. Protein concentration was determined using a commercial protein assay reagent (500–0006; Bio-Rad Laboratories). After recording

background luminescence, diluted ATP standard solution (1, 10, 100, 1000, and 5000 nM) or supernatant was added to the standard reaction solution to measure values. The ATP concentration is expressed as picomoles per milligram protein. This experiment was performed with five biological replicates using a total of five young and five old queen bees.

2.6. Western blotting

The trophocytes and oenocytes from one young or old queen bee were homogenized with a blue pestle and sonicator in 100 μL of radioimmunoprecipitation buffer containing protease inhibitors (11697498001; Roche Applied Science, Indianapolis, IN, USA), and then centrifuged at 5000g for 10 min at 4 °C to obtain the supernatant. Protein concentration was determined using a commercial protein assay reagent (500–0006; Bio-Rad Laboratories). Western blotting was performed as previously described (Hsu et al. 2014; Hsu and Chuang 2014; Hsu and Hu 2014) using primary antibodies against ATP synthase (1:1000; produced in-house), ND1 (h00004535-A01, 1:1000; Abnova, Neihu, Taipei, Taiwan), and tubulin (ab6046, 1:10,000; Abcam, Cambridge, MA, USA) and using the horseradish peroxidase-conjugated secondary antibody (1:10,000). This experiment was performed with five biological replicates using a total of five young and five old queen bees.

2.7. Statistical analysis

The differences in the mean values between the two age groups were examined using a two-sample *t* test. $P < 0.05$ was considered statistically significant.

3. RESULTS

3.1. Mitochondrial density

To examine mitochondrial energy utilization in queen bees, we assayed the mitochondrial density in the trophocytes of young and old queen bees. The mitochondrial size in these trophocytes, determined using TEM, was small in young queen

bees and large in old queen bees (Figure 1a). The mitochondrial area in these trophocytes was significantly lower in young queen bees than in old queen bees ($n=137$, $P<0.01$; Figure 1b). However, the mitochondrial density, presented as the mitochondrial area/cell area, was not significantly different between young and old queen bees ($n=40$, $P>0.05$; Figure 1c).

3.2. $\Delta\Psi_m$

To further evaluate mitochondrial energy utilization in queen bees, we assayed $\Delta\Psi_m$ in the trophocytes and oenocytes of young and old queen bees. The trophocytes and oenocytes were stained with TMRM and showed red fluorescence (Figure 2a). The $\Delta\Psi_m$ was presented as the ratio

of red fluorescence/cell area, and it was not significantly different between young and old queen bees ($n=25$, $P>0.05$; Figure 2b).

3.3. NAD^+ , $NADH$, and ATP concentration

To examine the relationship between $\Delta\Psi_m$ and levels of NAD^+ , $NADH$, and ATP, we assayed the concentrations of NAD^+ , $NADH$, and ATP in the trophocytes and oenocytes of young and old queen bees. The mean values for NAD^+ concentration in the trophocytes and oenocytes were 1.05 ± 0.19 and $1.28 \pm 0.12 \mu\text{mol mg}^{-1}$ of protein in young and old queen bees, respectively ($n=5$, $P>0.05$; Figure 3a). The mean values for $NADH$ concentration in trophocytes and oenocytes were 0.23 ± 0.01 and $0.25 \pm 0.01 \mu\text{mol mg}^{-1}$ of protein

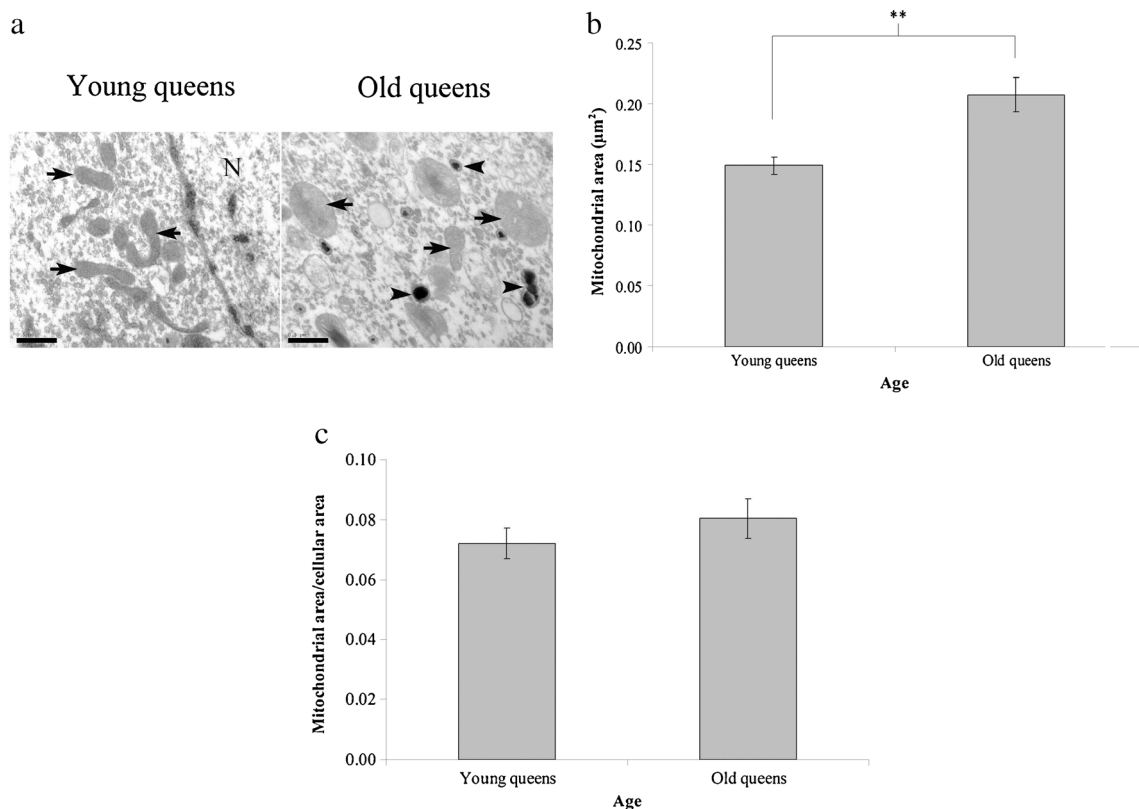


Figure 1. Mitochondrial density in the trophocytes of queen bees. **a** Mitochondria in trophocytes were examined using TEM. Scale bar, 1 μm . Arrows point to mitochondria. Arrowheads point to iron granules. **b** Mitochondrial area, presented as the mean \pm standard error of the mean (SEM), was calculated from the TEM images. **c** Mitochondrial density is presented as the mitochondrial area/cellular area ratio. Bars are presented as the mean \pm SEM. Asterisks indicate statistically significant difference (** $P<0.01$; two-sample t test).

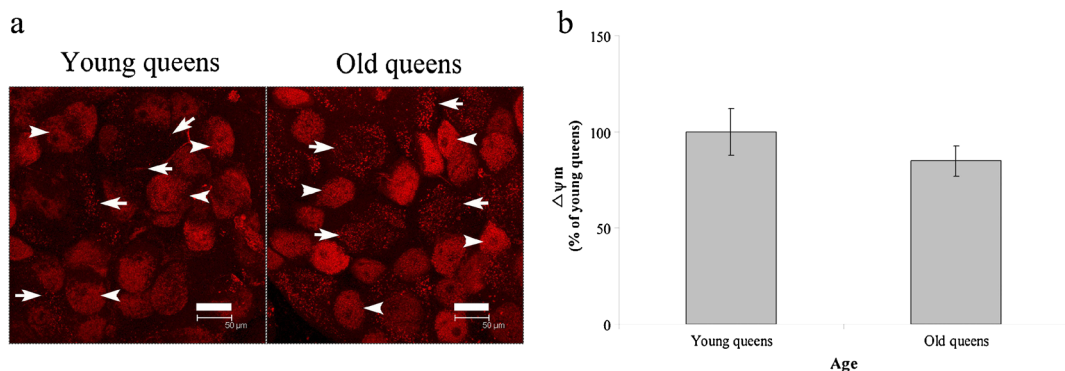


Figure 2. $\Delta\psi_m$ in the trophocytes and oenocytes of queen bees. **a** Representative confocal images of the stained TMRM. Scale bar, 50 μm . Arrows point to trophocytes. Arrowheads point to oenocytes. **b** $\Delta\psi_m$ is presented as the red fluorescence/cellular area ratio. The red fluorescence/cellular area ratio was normalized to that in young worker bees. The results are presented as the mean \pm SEM and are expressed as percentages.

in young and old queen bees, respectively ($n=5$, $P>0.05$; Figure 3b). The mean values for NAD^+ /NADH ratio in trophocytes and oenocytes were

4.47 \pm 0.67 and 5.25 \pm 0.59 in young and old queen bees, respectively ($n=5$, $P>0.05$; Figure 3c). The mean values for ATP concentration in trophocytes

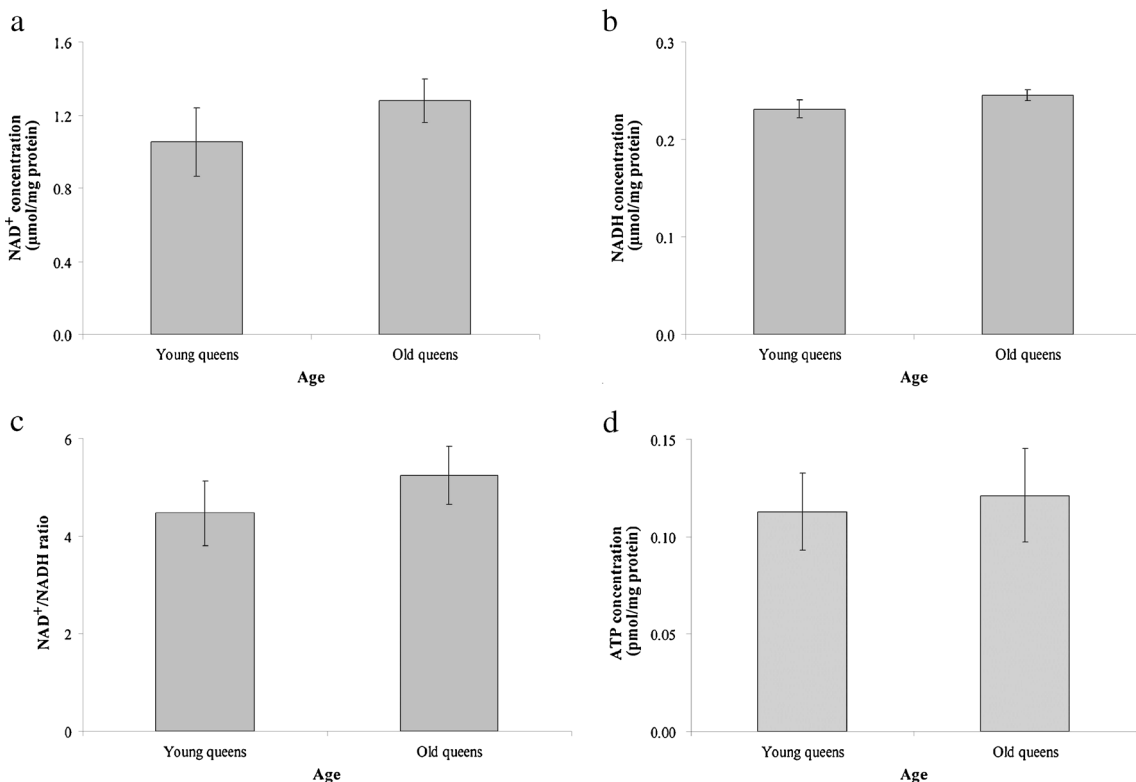


Figure 3. NAD^+ concentration (a), NADH concentration (b), NAD^+ /NADH ratio (c), and ATP concentration (d) in the trophocytes and oenocytes of queen bees. The values denote the mean \pm SEM.

and oenocytes were 0.11 ± 0.02 and 0.12 ± 0.02 pmol mg^{-1} of protein in young and old queen bees, respectively ($n=5$, $P>0.05$; Figure 3d).

3.4. ND1 and ATP synthase expression

To extend our evaluation of NAD^+ and ATP, we assayed the expression of ND1 and ATP synthase in the trophocytes and oenocytes of young and old queen bees. ND1 expression was lower in young queen bees than in old queen bees (Figure 4a), and this difference was statistically significant ($n=5$, $P<0.05$; Figure 4b). Similarly, expression of ATP synthase was also low in young queen bees and high in old queen bees

(Figure 4c), and this difference was statistically significant ($n=5$, $P<0.05$; Figure 4d). Because ND1 and ATP synthase are located in the mitochondria, the expression levels of these enzymes were normalized to the mitochondrial area. The normalized ND1 and ATP synthase levels were not significantly different between young and old queen bees ($n=5$, $P>0.05$; Figure 5a; $n=5$, $P>0.05$; Figure 5b).

4. DISCUSSION

In the present study, we evaluated mitochondrial energy utilization in the trophocytes and oenocytes of young and old queen bees.

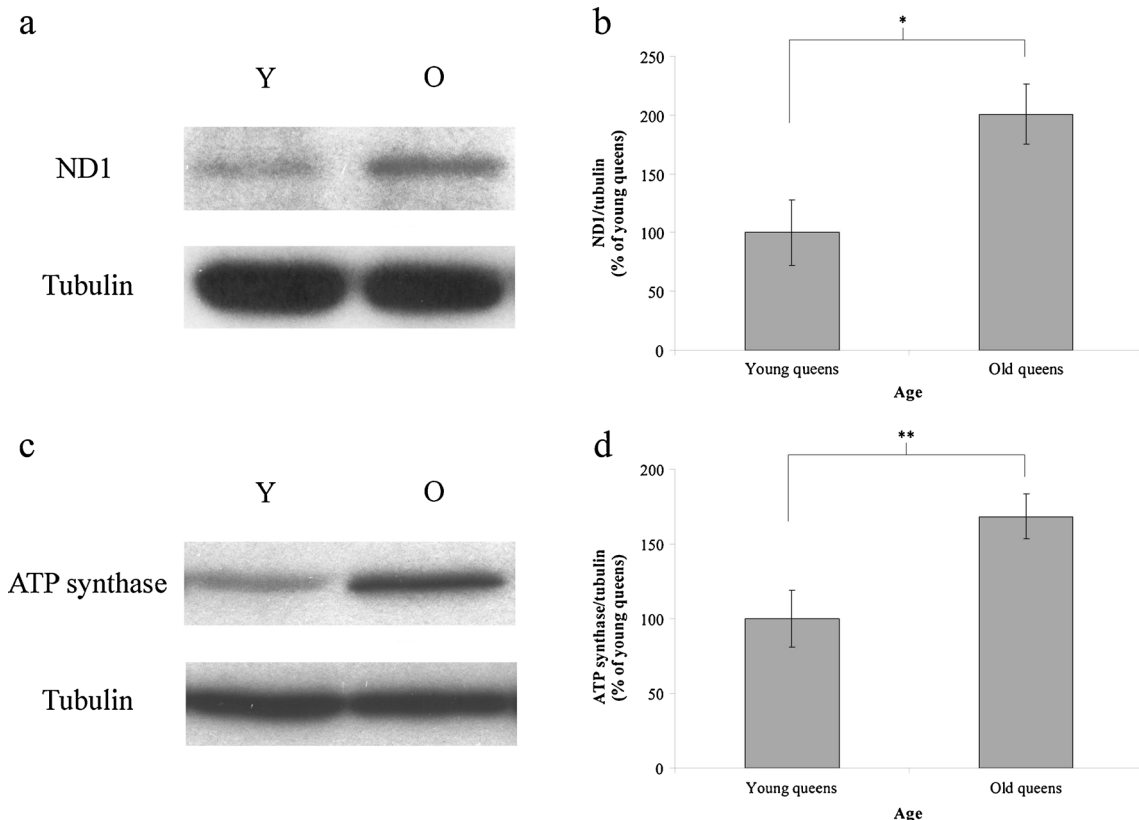


Figure 4. ND1 and ATP synthase expression in the trophocytes and oenocytes of queen bees. **a** ND1 protein concentration was analyzed using Western blotting. Tubulin served as a loading control. **b** ND1 expression was normalized to that in young worker bees. The results are presented as the mean \pm SEM and are expressed as percentages. **c** ATP synthase concentration was analyzed through Western blotting. Tubulin served as a loading control. **d** ATP synthase expression was normalized to that in young worker bees. The results are presented as the mean \pm SEM and are expressed as percentages. Asterisks indicate a statistically significant difference (* $P<0.05$, ** $P<0.01$; two-sample t test). *Y* young queen bees, *O* old queen bees.

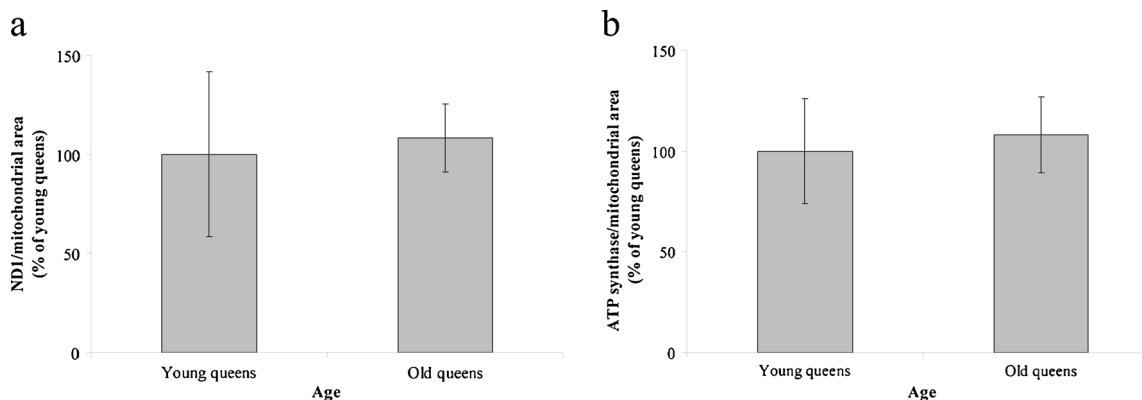


Figure 5. ND1 and ATP synthase expression normalized to mitochondrial density. **a** ND1/mitochondrial density ratio. **b** ATP synthase/mitochondrial density ratio. Ratios are expressed relative to those in young worker bees. The results are presented as the mean \pm SEM and are expressed as percentages.

Young and old queen bees had similar mitochondrial density; $\Delta\psi_m$; NAD^+ , $NADH$, and ATP concentrations; and area-normalized ND1 and ATP synthase levels. These results suggest that mitochondrial energy utilization is unchanged in the trophocytes and oenocytes of young and old queen bees, and these cells seem to have longevity-promoting mechanisms that can be used to investigate the longevity of queen bees.

4.1. Mitochondrial density

An increase in mitochondrial density with age has been reported in many contexts, such as in senescent human diploid fibroblasts (Martinez et al. 1991; Xu and Finkel 2002); the lung and brain of aging humans (Lee et al. 1998; Barrientos et al. 1997; Lee et al. 2000); the liver, heart, and brain of old rats (Gadaleta et al. 1992); the liver of old mice, rats, guinea pigs, rabbits, sheep, cows, hamsters, and humans (Passos et al. 2007); and shorter-lived mammals (Passos et al. 2007). Similarly, the mitochondrial density increases with age in the trophocytes and oenocytes of 50-day-old worker bees (Chuang and Hsu 2013). In the present study, however, the mitochondrial density in the trophocytes and oenocytes of young and old queen bees was similar, although the old queen bees were 16 months old. These results most likely reflect delayed senescence mechanisms in

the trophocytes and oenocytes of queen bees, which maintain these cells in a young status.

Although mitochondrial density did not appear to change, a change in mitochondrial morphology was observed. The mitochondrial area (and thus presumably the mitochondrial volume) of old queen bees was larger than that of young queen bees, and the mitochondrial morphology of old queen bees was more ovoid than that of young queen bees. In addition, because mitochondrial density did not appear to change, we infer that the size of the trophocytes of old queen bees was also increased. The enlargement of the mitochondria and trophocytes in old queen bees may be associated with cellular functions in old queen bees. Further research is required to clarify this point.

4.2. $\Delta\Psi_m$

$\Delta\Psi_m$ controls respiration, ATP synthesis, and ROS production. In turn, $\Delta\Psi_m$ is controlled through electron transport and proton leakage (Nicholls 2004). $\Delta\Psi_m$ decreases with age in the hepatocytes of rats and mice (Hagen et al. 1997; Kokoszka et al. 2001), the lymphocytes of mice (Wikowski and Micklem 1985; Rottenberg and Wu 1997), and the heart of rats (Savitha and Panneerselvam 2006). Similarly, $\Delta\Psi_m$ decreases with age in the trophocytes and oenocytes of worker bees (Chuang and Hsu 2013). These

phenomena reflect mitochondrial dysfunction during the aging process (Ames et al. 1995; Trifunovic and Larsson 2008; Artal-Sanz and Tavernarakis 2008; López-Lluch et al. 2008; Chanséaume and Morio 2009). In this study, however, the $\Delta\Psi_m$ in the trophocytes and oenocytes of young queen bees was similar to that in old queen bees, potentially reflecting delayed senescence mechanisms in these cells that maintain them in a young status.

4.3. NAD^+ , NADH, and ATP concentrations

Protons from NADH are pumped out of the mitochondrial matrix, creating a proton gradient that forms the $\Delta\Psi_m$. A decline in this activity results in a decrease in $\Delta\Psi_m$ and a subsequent decrease in NAD^+ concentration. NAD^+ concentration decreases with age in the pancreatic β cells and neurons of mice (Ramsey et al. 2008; Imai 2009) and in the liver, heart, kidney, and lung of rats (Braidy et al. 2011). Similarly, the NAD^+ concentration and $NAD^+/NADH$ ratio decreases with age in the trophocytes and oenocytes of worker bees (Chuang and Hsu 2013). In this study, however, the NAD^+ concentration and $NAD^+/NADH$ ratio in trophocytes and oenocytes were similar in young and old queen bees.

ATP concentration decreases with age in the gastric mucosa and muscle of humans (Kawano et al. 1991; Gurd et al. 2008), the blood and brain of mice (Jayachandran et al. 2005; Joo et al. 1999), the erythrocytes of rabbits (Subasinghe and Spence 2008) and cows (Bartosz et al. 1982), and the heart of rats (Guerrieri et al. 1996). Similarly, we have found that ATP concentration decreases with age in the trophocytes and oenocytes of worker bees (Hsu and Chan 2013; Chuang and Hsu 2013). In this study, however, the ATP concentration in trophocytes and oenocytes was similar in young and old queen bees. This finding is consistent with the results obtained for $\Delta\Psi_m$, NAD^+ concentration, and $NAD^+/NADH$ ratio. A possible explanation for these results is that the trophocytes and oenocytes of queen bees have delayed senescence mechanisms that maintain these cells in a young status.

4.4. ND1 and ATP synthase expression

ND1 expression increased with age in the trophocytes and oenocytes of queen bees. This finding is consistent with the previous finding that NADH dehydrogenase expression increases with age in the muscle of rats (Chang et al. 2007), the lung fibroblasts of humans (Allen et al. 1999), and the trophocytes and oenocytes of worker bees (Chuang and Hsu 2013). The expression of ATP synthase also increased with age in the trophocytes and oenocytes of queen bees. Previous studies have reported similar findings, showing that the expression of ATP synthase increases with age in the brain and muscle of rats (Nicoletti et al. 1995; Chang et al. 2007; Doran et al. 2008; Donoghue et al. 2010), the muscle of humans (Gelfi et al. 2006), and the trophocytes and oenocytes of worker bees (Chuang and Hsu 2013).

Although ND1 and ATP synthase expression, measured in absolute terms, both increased with age in the trophocytes and oenocytes of queen bees, the levels of ND1 and ATP synthase normalized against the mitochondrial density were not significantly different between young and old queens. This observation is consistent with the results of a previous study of worker bees (Chuang and Hsu 2013). In worker bees, ND1 and ATP synthase levels normalized against the mitochondrial density were not significantly different between young and old worker bees, but NAD^+ and ATP concentrations decreased in old worker bees. These phenomena suggest that old worker bees have reduced the efficiency of ND1 and ATP synthase (Chuang and Hsu 2013). NAD^+ and ATP concentrations were also not significantly different between young and old queen bees, indicating ND1 and ATP synthase do not experience age-related declines in efficiency.

Taken together, the results of the present study indicate that young and old queen bees had similar mitochondrial density; $\Delta\Psi_m$; NAD^+ , NADH, and ATP concentration; and ND1 and ATP synthase expression. These findings suggest that mitochondrial energy utilization does not decline in the trophocytes and oenocytes of old queen bees. These results are consistent with a recent study showing that energy-regulated molecules

maintain a young status in the trophocytes and oenocytes of old queen bees (Hsu and Hu 2014). In addition, this study demonstrates that the trophocytes and oenocytes of queen bees have longevity-promoting mechanisms. This speculation is consistent with previous studies (Hsieh and Hsu 2011b; 2013; Hsu and Hu 2014).

Parallel increases in age-related molecule accumulation and oxidative stress in the trophocytes and oenocytes of old queen bees indicate that high oxidative stress results in high expression of age-related molecules (Hsieh and Hsu 2011b; Hsieh and Hsu 2013). Higher energy metabolism increases oxidative stress (Kowaltowski et al. 2009; Hsu and Hsieh 2014), and ROS are generated as by-products of energy metabolism (Halliwell and Gutteridge 1999). Therefore, the trophocytes and oenocytes of old queen bees are inferred to have high energy metabolism. However, our recent study shows that energy-regulated molecule expression is similar between young and old queen bees. These results are consistent with the current finding that mitochondrial energy utilization is similar between young and old queen bees. Thus, the patterns of age-related molecule accumulation and oxidative stress are inconsistent with those of energy-regulated molecule expression and mitochondrial energy utilization. The most likely reason is that high ROS levels in old

queen bees do not derive from high-energy metabolism but perhaps from other superoxide generators, such as reduced nicotinamide adenine dinucleotide phosphate (NADPH) oxidase. This inference is supported by our previous study showing that old queen bees have high NADPH oxidase activity in trophocytes and oenocytes (Hsieh and Hsu 2013). In addition, high ROS may derive from hydrogen peroxide, which has important roles as a signaling molecule in the regulation of biological processes (Veal et al. 2007; Giorgio et al. 2007). This inference is supported by our study of worker bees (Hsu and Hsieh 2014). The above phenomena are consistent with previous studies showing that increased oxidative stress promotes the longevity and metabolic health of organisms (Ristow and Zarse 2010; Yang and Hekimi 2010; Ristow and Schmeisser 2011).

Furthermore, if high ROS levels derive from high metabolism, the superoxide most likely derives from the conversion of NADH to NAD⁺ (Camello-Almaraz et al. 2006). High superoxide levels then induce UCP2 expression and activate UCP2-mediated proton leak (Krauss et al. 2003). In queen bees, NAD⁺ concentration, UCP2 expression ($n=5$, $P>0.05$; Figure 6a), and $\Delta\Psi_m$ were similar between young and old queen bees, showing that they have similar energy metabolism. In contrast, young worker bees had high

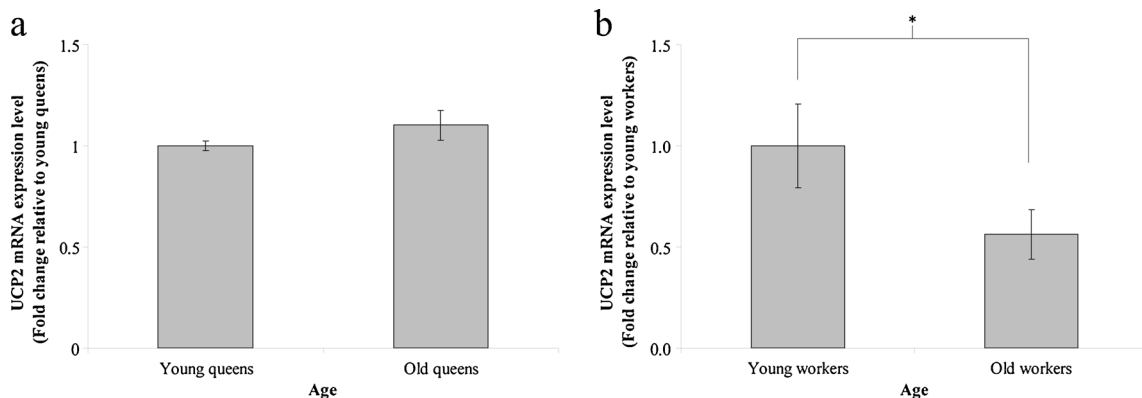


Figure 6. The mRNA expression of UCP2 in trophocytes and oenocytes was measured by quantitative PCR. **a** The mRNA expression of UCP2 in queen bees. The results are normalized to young queen bees, are shown as fold changes, and represent the mean \pm SEM. **b** The mRNA expression of UCP2 in worker bees. The results are normalized to that of young worker bees, are shown as fold changes, and represent the mean \pm SEM. Asterisks indicate a statistically significant difference ($*P<0.05$; two-sample t test).

energy metabolism and simultaneously had high NAD^+ concentration, high UCP2 expression ($n = 10$, $P < 0.05$; Figure 6b), and high $\Delta\Psi_m$ (Chuang and Hsu 2013). Therefore, high ROS levels in old queen bees do not derive from high energy metabolism.

Oxidative stress decreases in the trophocytes and oenocytes of worker bees because young worker bees have high ROS levels from high mitochondrial energy utilization (Hsu and Hsieh 2014). These ROS damage lipids and proteins over time, resulting in the accumulation of lipid peroxidation and protein oxidation. Although old queen bees have high ROS levels, ROS are also present in young queen bees. ROS in young queen bees also damage lipids and proteins resulting in the accumulation of lipid peroxidation and protein oxidation. In older queen bees, lipid peroxidation and protein oxidation may be more obvious and mitochondrial energy utilization and the expression of energy-regulated molecules may show significant differences. Further research is required to clarify these questions.

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L'utilisation de l'énergie mitochondriale maintient les trophocytes et les oenocytes à l'état jeune chez les vieilles reines d'abeilles

Vieillesse / mitochondrie / utilisation de l'énergie / *Apis mellifera*

Die Nutzungsweise der mitochondrialen Energieproduktion hält die Trophozyten und Oenozysten alter Königinnen der Honigbiene auf einem jungen Niveau

Alterung / Mitochondrien / Energienutzung / *Apis mellifera*

Conflict of interest statement We have no conflicts of interest or disclosures.

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