

# UQCRC1 downregulation impairs cognitive function in mice via AMPK inactivation (#114201)

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# UQCRC1 downregulation impairs cognitive function in mice via AMPK inactivation

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**Background:** Ubiquinol-cytochrome c reductase core protein 1 (UQCRC1) is a subunit of complex III of the mitochondrial respiratory chain. Although earlier studies have indicated that UQCRC1 down-regulation causes cognitive impairment, the precise processes by which this happens are yet unknown.

**Methods:** In order to investigate its pathophysiological effects, we developed a mouse model with downregulated UQCRC1 expression. We evaluated hippocampal-dependent cognitive performance using behavioral paradigms. Then we quantified changes in bioenergetic state by  level measurements and oxidative stress utilizing reactive oxygen species (ROS) detection.  P-activated protein kinase (AMPK) signaling dynamics and autophagic flux changes were assessed by molecular studies. Intervention strategies involving AMPK activation and lysosomal function potentiation were subsequently employed to elucidate mechanistic pathways.

**Results:** Our results show that UQCRC1 loss causes notable hippocampal-dependent cognitive deficits together with expected mitochondrial bioenergetics (lower ATP synthesis) and higher oxidative stress (more ROS buildup). Mechanistically, this phenotypic expression was linked to reduced AMPK activation and impaired autophagic flux. In UQCRC1 defective mice, pharmacological stimulation of AMPK or therapeutic potentiation of lysosomal activity essentially corrected cognitive deficits and restored mitochondrial redox equilibrium.

**Conclusions:** This work mechanistically defines AMPK as a fundamental metabolic orchestrator of mitochondrial-lysosomal functional crosstalk and reveals its non-canonical function in maintaining neuronal homeostasis via coordinated control of autophagic flux and redox balance. Our identification of AMPK-driven interorganelle communication as a modifiable treatment target creates a fresh paradigm for tackling cognitive decline originating in bioenergetic dysregulation.

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## Abstract

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## Introduction

In mammalian cells, mitochondria are the sites of energy metabolism and signal transmission. They are thus crucial for cellular proliferation, autophagy, apoptosis, and differentiation (Zheng et al., 2021; Zhou et al., 2018). The respiratory chain, a key structure responsible for  $ATP$  generation, is composed of two electron carriers (ubiquinone and cytochrome  $c$ ) and four complexes (CI-CIV). In neurons, mitochondrial malfunction may cause an excessive buildup of reactive oxygen species (ROS) and cytochrome  $c$  release into the cytoplasm. This subsequently reduces  $ATP$  generation, modulates the activity of respiratory chain complexes I, II, and III, and finally causes neuronal apoptosis and cognitive impairments (Fernandez-Vizarra & Zeviani, 2018a). Comprising two monomers, each with eleven subunits, Complex III of the respiratory chain is a symmetric dimer. UQCRC1 is the fundamental protein required for the formation of Complex III (Fernandez-Vizarra & Zeviani, 2018b). Previous research has shown that downregulation of UQCRC1 expression might cause cognitive impairment (Shan et al., 2019). Nonetheless, the underlying processes are still unknown.

Autophagy is a conserved process in eukaryotic evolution, which digests damaged organelles or proteins and recycles them (Mizushima & Komatsu, 2011). Autophagy flux mainly includes formation of autophagosomes, the fusion of autophagosomes with lysosomes, and the degradation of autolysosome contents. Impairment of either step will result in impaired autophagy flux (Levine & Kroemer, 2019; Liu et al., 2023). Mitophagy is the main mechanism controlling the quantity and quality of mitochondria, which is essential for many life activities such as homeostasis, proliferation, aging, apoptosis, etc., and is closely related to the occurrence and development of neurodegenerative diseases, metabolic diseases, tumors, and other diseases (Wang et al., 2023; Picca et al., 2023). After mitochondrial dystrophy or damage caused by undesirable stimuli, it is labeled through various pathways, and LC3-II on the autophagosome membrane recognizes and binds to the labeled mitochondria. Subsequently, autophagosomes wrap the recognized mitochondria and fuse with lysosomes to degrade them. Abnormal mitophagy in neurons can lead to a range of problems, such as metabolic disturbances, oxidative stress, synaptic dysfunction, and calcium homeostasis imbalance (Katayama et al., 2020; Kerr et al., 2017).

The AMP-activated protein kinase (AMPK) is an essential regulator and sensor of cellular metabolism and stress responses. Triggers include energy stress, changes in cytoplasmic  $Ca^{2+}$  levels, and the presence of reactive oxygen species (ROS) (Trefts & Shaw, 2021). In several

ways, AMPK is involved in cellular homeostasis maintenance. It affects mitochondrial fusion and fission, therefore regulating mitochondrial biogenesis and hence affecting mitochondrial function (Virga et al., 2024; Herzog & Shaw, 2018). Moreover, AMPK plays a multifaceted role in lysosomal biogenesis and function, hence supporting cellular homeostasis (Paquette et al., 2021). AMPK therefore may be a fundamental protein linking mitochondrial malfunction to the autophagy mechanism (Hu et al., 2021). Although earlier studies on mitophagy mostly concentrate on how autophagy controls mitochondria, new investigations indicate that mitochondria may potentially reciprocally control autophagy.

This study investigated the mechanistic role of UQCRC1 in cognitive regulation. Our findings demonstrate that UQCRC1<sup>+/-</sup> mice showed hippocampal-dependent cognitive impairment along with lower ATP generation, higher ROS levels, poor autophagy, and more hippocampal cell apoptosis. Especially, our results underlined the important roles lysosomal and AMPK activation play in these abnormalities.

## Materials & Methods

### Animals

C57BL/6 wild-type (WT) mice were provided by Charles River Laboratories (Chengdu, China) and UQCRC1<sup>+/-</sup> heterozygous mice were generated by Professor Zhiyi Zuo (Shan et al., 2019), with all animals housed under specific pathogen-free (SPF) conditions at Army Medical University's animal facility. Mice (8-12 weeks old, 20-30 g) were maintained in standard cages (330 × 210 × 170 mm; 5 mice/cage) with ad libitum access to food and water, under controlled environmental parameters: 12-h light/dark cycle (08:00-20:00), 20-23 °C ambient temperature, and 50-60% relative humidity.

Complemented by 16 female animals (8 UQCRC1<sup>+/-</sup> and 8 WT) for behavioral assessments, this study included 36 male WT mice and 63 male UQCRC1<sup>+/-</sup> mice. After pre-treatment behavioral testing, 8 WT and 8 UQCRC1<sup>+/-</sup> males were paired with 10 more genotype-matched males (total n=18 per genotype) for hippocampus tissue collection at baseline. Six specimens per genotype were assigned to ROS/ATP/caspase assays, six to Western blot analysis, and six to transmission electron microscopy (TEM). The therapeutic evaluation phase consisted of 45 UQCRC1<sup>+/-</sup> males equally split into three treatment cohorts (solvent vehicle, A-769662, and LH2-051). Eight mice per group received post-treatment behavioral assessment, with the addition of 7 more mice per group (total n=15/group) for parallel tissue studies (six specimens per group were assigned to ROS/ATP/caspase assays, three to Western blot analysis, and six to TEM). For a comparison study, a separate cohort of 36 WT males followed identical experimental schedules and tissue allocation guidelines.

Animals were euthanized prior to the planned endpoint only if they met predefined humane endpoints. These criteria included (but were not limited to): severe weight loss (>20% of baseline body weight) or failure to thrive; signs of irreversible distress or pain (e.g., labored breathing, prolonged immobility, inability to access food/water); unexpected complications directly related to the experimental intervention (e.g., neurological deficits). No animals required



early euthanasia in this study, as all subjects maintained stable health metrics within predefined thresholds throughout the experimental timeline. No animals were retained beyond the study period due to the terminal nature of the experimental design. Male subjects received 1% sodium pentobarbital (50 mg/kg) intraperitoneally, while all female cohorts were euthanized following behavioral assessment using gradual  $O_2$  asphyxiation (30%–99%, 15 min) with secondary death confirmation. Randomization using random number tables ensured objective group assignments, with sample sizes calculated by power analysis incorporating preliminary data and literature benchmarks (Shan et al., 2019; Lin et al., 2020; Fernandez-Mosquera et al., 2019a; Chen et al., 2022; Kim et al., 2021). The experimental protocol was developed prior to study commencement and conducted in compliance with guidelines approved by the Army Medical University's Laboratory Animal Welfare and Ethics Committee (Approval No.: AMUWEC20245280; Approval date: 10/1/2024).

### Behavioral testing

One week before the trials began, all the mice were adjusted to their surroundings. Every behavioral test took place beginning at 10 AM. Two hours before the start of every trial, mice were allowed to become acquainted with the testing room. Different behavioral assessments were kept one day apart. Following every test, the equipment was carefully cleaned with 75% alcohol before the next one. Examiners left the room during testing to reduce outside influence. EthoVision XT 11.5 (Noldus Inc., Netherlands) was used for data capture and analysis.

#### Novel Object Recognition test (NOR)

The ability of short-term memory in the hippocampus of mice was evaluated using the Novel Object Recognition (NOR) test (Bevins & Besheer, 2006). The experimental setup consisted of an acrylic rectangular enclosure (40 x 40 x 40 cm). The test comprised three separate phases. Mice in the first step, the habituation phase, were allowed to explore freely for ten minutes in the apparatus. Conducted 24 hours after habituation, in the second step, two identical objects (Familiar Object, F) were symmetrically placed within the apparatus. Reintroduced into the center area, mice had ten minutes to wander freely. Two hours after the second stage, in the third stage of the novel-object test, one of the familiar items was substituted with a new object of the same size but different form (Novel Object, N). Reintroduced into the center area, the mice were allowed to search for another ten minutes. The time spent exploring the familiar object (tF) and the novel object (tN) was recorded, and the recognition rate (RR) was calculated as follows:  $RR = tN / (tF + tN)$ .

#### Nest Building test (NBT)

We used NBT to evaluate the general effect of UQCRC1 knockdown on hippocampally reliant cognition. Experimental mice were individually housed in conventional cages with a single 2.5 g, 5 cm<sup>2</sup> tearable cotton pad and a tiny quantity of wood shavings on the day of the test at 6:00 PM. Enough food and drink were offered. Photographs were taken the next day to record the nesting activity, paying special attention to the degree to which the cotton pads were ripped and used in nest building (Fig. 1C). After that, nesting quality was assessed and scored according to criteria



defined in earlier research (Deacon, 2006). Data from mice whose cotton pads were moist were excluded from the analysis.

# **Barnes Maze**

The Barnes maze test was a low-stress and effective way to evaluate spatial learning and memory in mice. The apparatus consisted of a circular platform with 20 evenly spaced holes, one of which was selected as the target hole. Under this hole was a detachable black box that functioned as a sanctuary, enabling the mice to escape from lights and aural stimulation (Fig. 1E). Two stages comprised the test: the training phase and the testing phase. Mice positioned in the middle of the labyrinth during the training period were watched for behavior. The test was stopped when the mouse found and entered the black box or when the test lasted for three minutes. Should a mouse fail to locate the box during this period, it was softly directed to the black box and allowed to stay for one minute. Each mouse underwent three daily training sessions for four consecutive days. The main performance indicator was the latency to enter the black box. Conducted 24 hours after the last training session, the testing phase proceeded exactly like the training phase. Performance measures for this phase were the latency to reach the black box and the number of tries to find the box.

# **Transmission Electron Microscope (TEM)**

Mice were initially perfused with ice-cold PBS. The hippocampus was carefully isolated and sectioned into small fragments. The tissue samples were then fixed, dehydrated, infiltrated, and immersed as required. Ultra-thin sections were prepared and stained with 2% uranyl acetate. Quantitative analysis of the images was performed using ImageJ software (NIH, <https://imagej.nih.gov/ij/>, version 1.54).

# **Western Blotting**

The hippocampus was harvested and homogenized, and 20 µg of protein was loaded onto the 4–20% gels (ACE Biotechnology, cat# ET15420LGel) and then transferred to a PVDF membrane using the Bio-Rad system. Membranes were blocked with a rapid blocking buffer (MedChemExpress, cat# HY-K1027) for 10 minutes. The primary antibodies were incubated overnight at 4°C. The main antibodies used were rabbit polyclonal anti-IL-1β (1:1000 dilution, Abcam, cat# ab48394), rabbit polyclonal anti-AMPKα (1:1000 dilution, Cell Signaling Technology, cat# 2532), and rabbit monoclonal anti-phospho-AMPKα (Thr172) (1:1000 dilution, Cell Signaling Technology, cat# 2535). After being diluted 1:3000 (ZSGB-BIO, cat# ZB-2301), the HRP-conjugated goat anti-rabbit IgG was incubated at room temperature for one hour. Images were processed using the ImageJ program (<https://imagej.nih.gov/ij/>, version 1.54) for densitometric analysis.

# **ATP Assay**

The method offered in the ATP content test kit handbook (Servicebio, cat# G4309-48T) was followed to measure ATP content. Extracted hippocampal tissue was homogenized for lysis, was boiled, cooled to room temperature. The sample was then centrifuged at 10,000 × g for 15 minutes at 4°C. After obtaining the supernatant, 20 µL of supernatant was added to 100 µL of

ATP assay reagent and mixed thoroughly. The bioluminescent intensity was then assessed using a luminometer (Fig. 2A).

### Assessment of reactive oxygen species (ROS)

Reactive oxygen species (ROS) levels were gauged using the ROS Detection Kit (Bestbio, cat# BB-47051). The homogenate was centrifuged at 1000 g for three minutes at 4°C after the hippocampal harvest and homogenization. Two  $\mu$ L of dihydroethidium (DHE) probe was then added to 200  $\mu$ L of supernatant and incubated in the dark at 37°C for thirty minutes. Measuring fluorescence intensity at 510 nm as an excitation wavelength and 610 nm as an emission wavelength. Calculating the ratio of fluorescence intensity to protein concentration to determine the ROS levels in the tissue.

### Assessment of Caspase 3 and Caspase 9

The Caspase 3/Caspase 9 Activity Assay Kit (made by MULTI SCIENCES, cat# APC03/APC09) was used to evaluate the activation of caspase 3 and caspase 9. In a nutshell, homogenized hippocampal tissue was centrifuged at 12,000 rpm for 15 minutes at 4 °C after extraction. Following the directions on the kit, we collected the supernatant and made the reaction mixture. After four hours of incubation at 37 °C, the samples were tested for absorbance at 405 nm.

### Intraperitoneal injection

UQCRC1<sup>+/-</sup> mice were randomly allocated into three groups: UQCRC1<sup>+/-</sup> + A-769662, UQCRC1<sup>+/-</sup> + LH2-051, and UQCRC1<sup>+/-</sup> + solvent. Mice in the UQCRC1<sup>+/-</sup> + A-769662 and UQCRC1<sup>+/-</sup> + LH2-051 cohorts received intraperitoneal injections of A-769662 (30 mg/kg, MCE, cat# HY-50662) or LH2-051 (10 mg/kg, MCE, cat# HY-161723), respectively, administered twice daily for a duration of 30 consecutive days. Mice in the UQCRC1<sup>+/-</sup> + solvent group received an equivalent volume of solvent consisting of 10% I<sub>2</sub>SO, 40% P<sub>300</sub>, 5% Tween-80, and 45% saline.

### Statistical analysis

All experiments and analyses were performed under blinded conditions, with investigators unaware of group allocations during both experimental procedures and data interpretation. All acquired data were retained for statistical analysis without exclusion. The Shapiro-Wilk test was employed to assess normality. Data conforming to a normal distribution were analyzed using parametric tests, while non-parametric tests were applied to data that violated the normality assumption. In the two-way ANOVA analysis, no outliers were detected using the method of evaluating whether studentized residuals exceeded  $\pm 3$ . The Kruskal-Wallis H test, unpaired two-tailed t-tests, Mann-Whitney U test, one-way ANOVA, and two-way ANOVA were used to assess differences between groups. Statistical significance was defined as a p-value of less than 0.05. Data analysis was performed using GraphPad Prism (GraphPad Software, Boston, USA, version 9.5) and SPSS (IBM, New York, USA, version 27.0). Statistical comparisons were represented as \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, and \*\*\*\*P < 0.0001.

## Results

### The downregulation of UQCRC1 impaired cognitive function.

In the NOR, both male (Fig. 1B,  $p < 0.0001$ ) and female (Fig. 1B,  $p < 0.05$ ) UQCRC1<sup>+/-</sup> mice demonstrated a markedly decreased recognition rate relative to WT controls. In the NBT, the male UQCRC1<sup>+/-</sup> mice exhibited reduced performance (Fig. 1D,  $p < 0.01$ ). From day 3 of the Barnes maze training phase, the male UQCRC1<sup>+/-</sup> mice exhibited substantially longer escape latencies (Fig. 1F,  $p < 0.05$  on day 3,  $p < 0.01$  on day 4). During the testing phase, male UQCRC1<sup>+/-</sup> mice exhibited prolonged escape latencies (Fig. 1G,  $p < 0.05$ ) and significantly increased error rates (Fig. 1H,  $p < 0.05$ ), whereas female UQCRC1<sup>+/-</sup> mice required more attempts to find the target hole (Fig. 1H,  $p < 0.05$ ). These results suggest that mitochondrial dysfunction caused by the downregulation of UQCRC1 was strongly associated with cognitive decline in male mice, while its effects on female mice were more variable. Thus, we focused on male mice in the following research.

### **The downregulation of UQCRC1 led to autophagy impairment.**

Our findings so far showed that the downregulation of UQCRC1 seriously reduced hippocampally reliant cognitive capacities. We investigated this phenomenon further and discovered a significant drop in ATP levels (Fig. 2B,  $p < 0.01$ ) and a rise in ROS levels (Fig. 2C,  $p < 0.05$ ) in UQCRC1<sup>+/-</sup> mice. Autolysosomes (Fig. 2D, E,  $p < 0.05$ ) were shown to be accumulating along with rising LC3B II expression (Fig. 2F, G,  $p < 0.01$ ). These results implied that autophagy failure in mice followed from UQCRC1 downregulation. Previous research has revealed that lysosomal malfunction in HeLa cells resulted from prolonged mitochondrial respiratory chain failure reducing AMPK activation (Fernandez-Mosquera et al., 2019a). In accordance with these results, our work showed that UQCRC1 downregulation resulted in a drop in AMPK levels (Fig. 2F, H,  $p < 0.00001$ ) and a lower pAMPK/AMPK ratio (Fig. 2F, J,  $p < 0.00001$ ), while total AMPK levels remained unchanged (Fig. 2F, I). These findings implied that UQCRC1 downregulation reduced hippocampal AMPK activation and impaired autolysosome degradation. Notably, we observed an increase in the activation of caspase 3 (Fig. 2K,  $p < 0.0001$ ) and caspase 9 (Fig. 2K,  $p < 0.001$ ), which suggested that increased apoptosis in mice resulted from UQCRC1 downregulation.

### **Enhancing AMPK activity or improving lysosomal function ameliorated autophagy in UQCRC1<sup>+/-</sup> mice.**

Our findings so far showed that the downregulation of UQCRC1 seriously reduced hippocampally reliant cognitive capacities. We investigated this phenomenon further and discovered a significant drop in ATP levels (Fig. 3B,  $p < 0.01$ ) and a rise in ROS levels (Fig. 3C,  $p < 0.05$ ) in UQCRC1<sup>+/-</sup> mice. Autolysosomes (Fig. 3D, E,  $p < 0.05$ ) were shown to be accumulating along with rising LC3B II expression (Fig. 3F, G,  $p < 0.01$ ). Furthermore, a rise in the activation of caspase 3 (Fig. 3K,  $p < 0.0001$ ) and caspase 9 (Fig. 3K,  $p < 0.001$ ) was noted. These results implied that autophagy failure and increased apoptosis in mice followed from UQCRC1 downregulation. Previous research had revealed that lysosomal malfunction in HeLa cells resulted from prolonged mitochondrial respiratory chain failure, reducing AMPK activation. In accordance with these results, our work showed that UQCRC1 downregulation resulted in a drop in pAMPK levels (Fig. 3F, H,  $p < 0.00001$ ) and a lower pAMPK/AMPK ratio

(Fig. 3F, I,  $p < 0.00001$ ), while total AMPK levels remained unchanged (Fig. 3F, J). These findings implied that UQCRC1 downregulation reduced hippocampal AMPK activation and impaired autolysosome degradation.

### **Increasing AMPK activity or enhancing lysosomal function could both rescue cognitive deficits in UQCRC1<sup>+/-</sup> mice.**

We performed behavioral evaluations after giving rescue therapy to investigate the potential causal relationship between autophagy dysfunction and the cognitive abnormalities observed in UQCRC1<sup>+/-</sup> mice. The findings indicated that augmenting AMPK activity or improving lysosomal function markedly enhanced the performance of UQCRC1<sup>+/-</sup> mice in the new object recognition test (Fig. 4A, B), nest-building test (Fig. 4C, D), and Barnes maze (Fig. 4E, F, G, H, I). The results indicated that AMPK-mediated autophagy failure was pivotal in the cognitive impairments linked to mitochondrial malfunction.

## **Discussion**

Since UQCRC1<sup>-/-</sup> mice displayed embryonic lethality, UQCRC1<sup>+/-</sup> mice were therefore chosen as the experimental subjects. The brain, being a highly oxygen-consuming and energy-demanding organ, requires a significant quantity of mitochondria to maintain its standard functions. A multitude of studies have established that mitochondria can impact cognitive abilities by modulating the functions of hippocampal neurons (Khacho et al., 2019; He et al., 2022). Furthermore, extensive research has underscored the strong correlation between mitochondrial activity and neurodegenerative disorders, including Alzheimer's disease (Wang et al., 2020; Bishop et al., 2010). In this study, our findings showed that the downregulation of UQCRC1 expression resulted in lower ATP levels and increased oxidative stress in the hippocampal tissue. Moreover, this downregulation hampered autophagy flux by reducing AMPK activity, which in turn resulted in increased neuronal apoptosis and contributed to hippocampus-dependent cognitive dysfunction. This suggests that increasing AMPK activity or improving autophagic flux may be a potential effective approach for treating cognitive decline caused by mitochondrial dysfunction.

In the context of cognitive disorders associated with mitochondrial dysfunction, existing research has highlighted a significant gender disparity. Before menopause, females appear to be better equipped to combat oxidative stress due to their higher antioxidant defenses, which may be attributed to the protective effects of estrogen (Viña & Borrás, 2010; Mandal et al., 2012; Grimm et al., 2016). Consistent with these findings, our study also demonstrated that the downregulation of UQCRC1 expression had a more pronounced and stable impact on males. Therefore, in the mechanistic research section, we only used male mice to eliminate potential confounding factors.

Autophagy has a complex and key role in neurons. It can keep the balance in neurons and help neural functions work normally, but it might also cause harm to neurons in some situations (Nixon & Rubinshtein, 2024). Produced during the start of autophagy, LC3II is a structural protein linked to the autophagosome membrane that finally breaks down with the contents in the autolysosome (Iriondo et al., 2022). Its expression degree exactly matches the count of autophagosomes and autolysosomes. In this work, we found higher LC3II expression in UQCRC1<sup>+/-</sup> mouse hippocampal tissue. Confirming a significant increase in the number of autophagosomes/autolysosomes in this area, transmission electron microscopy also indicated

poor breakdown and recycling of autolysosomes in the hippocampal tissue of UQCRC1<sup>+/-</sup> mice. Researchers have shown that LH2-051 significantly improves learning, memory, and cognitive capacity in a mouse model of Alzheimer's disease (AD) via increasing lysosomal degradation capability (Yin et al., 2023; Yin et al., 2023). This study found that LH2-051 greatly reduced cognitive deficits in UQCRC1<sup>+/-</sup> mice, indicating that lysosomal dysfunction was a key component of mitochondrial damage-induced cognitive impairments, and improving the function of lysosomes was a targeted therapy.

While autophagy typically works to maintain cell survival by removing damaged components, apoptosis is triggered when the damage is beyond repair. Apoptosis is a controlled and physiological type of cell apoptosis that happens under certain physiological or pathological conditions. It is controlled by genetic pathways (Renehan et al., 2001; Bredesen, 1995). Mitochondria play a crucial role in the apoptotic pathway, and dysfunctional mitochondria can trigger apoptosis via multiple pathways. The main events that initiate the cascade include the release of cytochrome c and the apoptosis-inducing factor (AIF) and the excessive opening of the mitochondrial permeability transition pore (mPTP). Reduced ATP content and elevated reactive oxygen species (ROS) can make neurons more sensitive to apoptosis-related proteins, which speed up the process (Nguyen et al., 2023; Bock & Tait, 2020). In this work, we found out that more neural apoptosis in the hippocampal tissue made the cognitive ability worse in UQCRC1<sup>+/-</sup> mice.

AMPK is a highly conserved protein kinase and a major responder to mitochondrial stress, which is very sensitive to changes in the ratio of AMP to ATP. Taking out ndufs4, which is a subunit of respiratory chain complex I, has been shown to lower AMPK activation in the brain (Fernandez-Mosquera et al., 2019b). Previous studies have also shown that AMPK was involved in lysosome formation in vitro and in living organisms (Cheng et al., 2021; Alers et al., 2012; Patra et al., 2019). In this study, we also demonstrated that via AMPK-mediated processes, mitochondrial malfunction caused autolysosomes accumulation. Together, AMPK might be an important mediator between mitochondrial dysfunction and an altered autophagy flux.

There are various limits to this research that need thought. First, our studies indicated sex-dependent different effects of UQCRCR1 downregulation on murine cognitive function, while the underlying mechanisms are still unknown. Second, while behavioral assessments preliminarily identified the hippocampus as the primary affected region, given the widespread neuronal network underpinning cognitive processes it is impossible to completely eliminate possible involvement from extra-hippocampal locations. Thirdly, although we demonstrated that UQCRCR1-mediated mitochondrial dysfunction impairs cognition through AMPK-dependent autophagic dysregulation, two critical mechanistic gaps persist: 1) the precise signaling cascades linking AMPK activation to autophagic modulation, and 2) the exact neurobiological mechanisms through which autophagy perturbations mediate cognitive deficits.



## Conclusions

The brain depends primarily on oxygen and energy, hence it needs significant mitochondrial reserves to maintain normal physiological operations. Because of its increased metabolic needs, limited vascular reserve, and unique synaptic plasticity properties, the hippocampus region shows especially sensitivity to hypoxic-ischemic damage and is thus a target of great importance for neuroprotective treatments. This work shows that in mouse models UQCRC1 deficiency causes hippocampus dependent cognitive impairment. UQCRC1 downregulation has been shown in subsequent studies to cause a cascade of pathological changes in the hippocampal cells including lowered ATP synthesis capacity, reduced AMPK activation ratio, along with elevated ROS accumulation, decreased autophagolysosome density, and caspase-3 and caspase-9 enhanced activation. By means of pharmacological therapies in UQCRC1<sup>+/-</sup> mice, we found that lysosomal enhancement by LH2-051 therapy and AMPK activation by intraperitoneal A-769662 essentially corrected cognitive abnormalities. Especially, AMPK potentiation greatly reduced autophagolysosome accumulation, but lysosomal modification had little effect on AMPK activity in turn. These results define a molecular route wherein UQCRC1 deficiency-mediated mitochondrial dysfunction aggravates neuronal apoptosis via lysosomal impairment in the hippocampal region with AMPK acting as a central regulating hub. This cascade offers fresh treatment options for reducing cognitive impairment linked with mitochondrial diseases.

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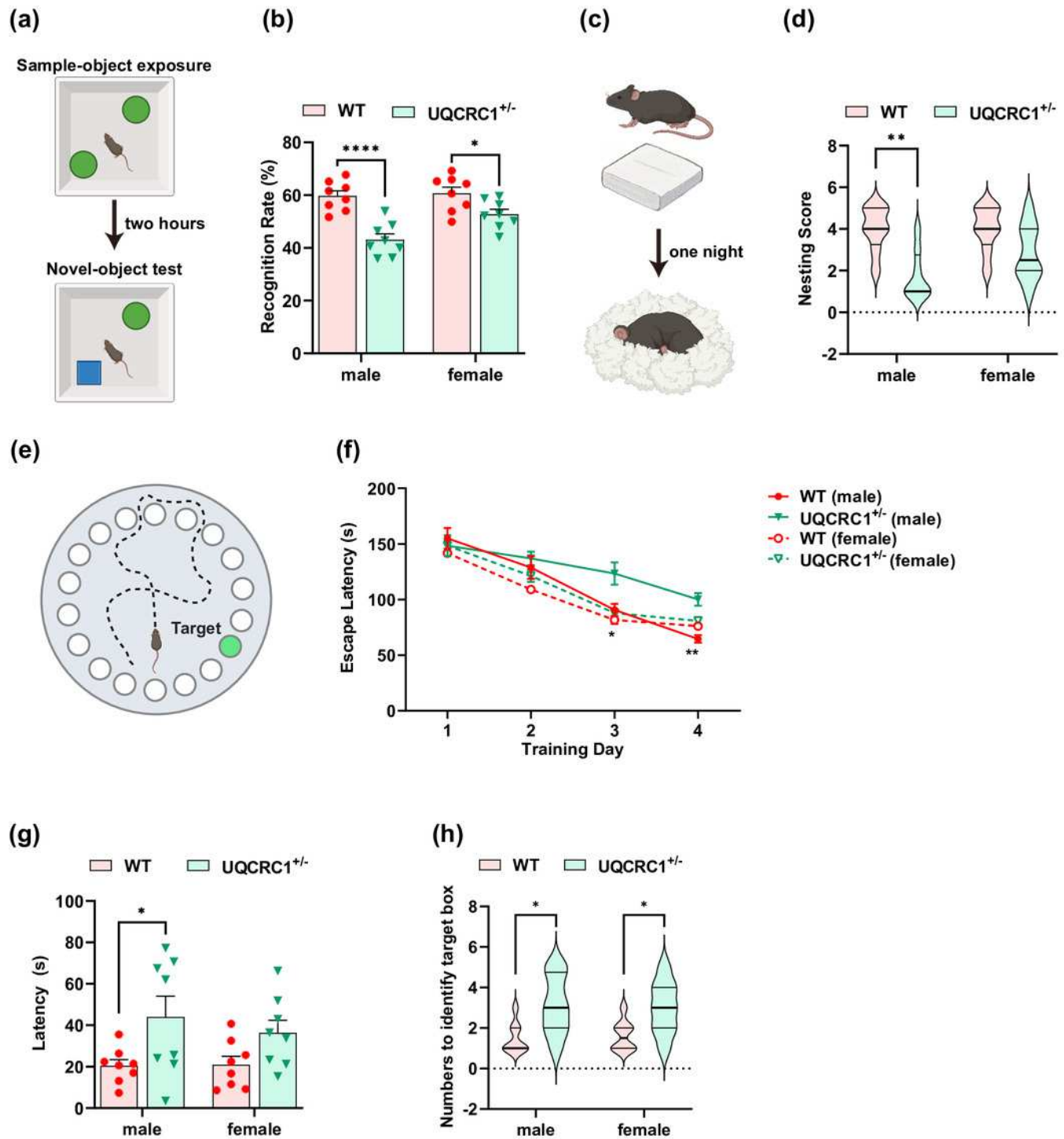
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# Figure 1

The downregulation of *UQCRC1* impaired cognitive function.

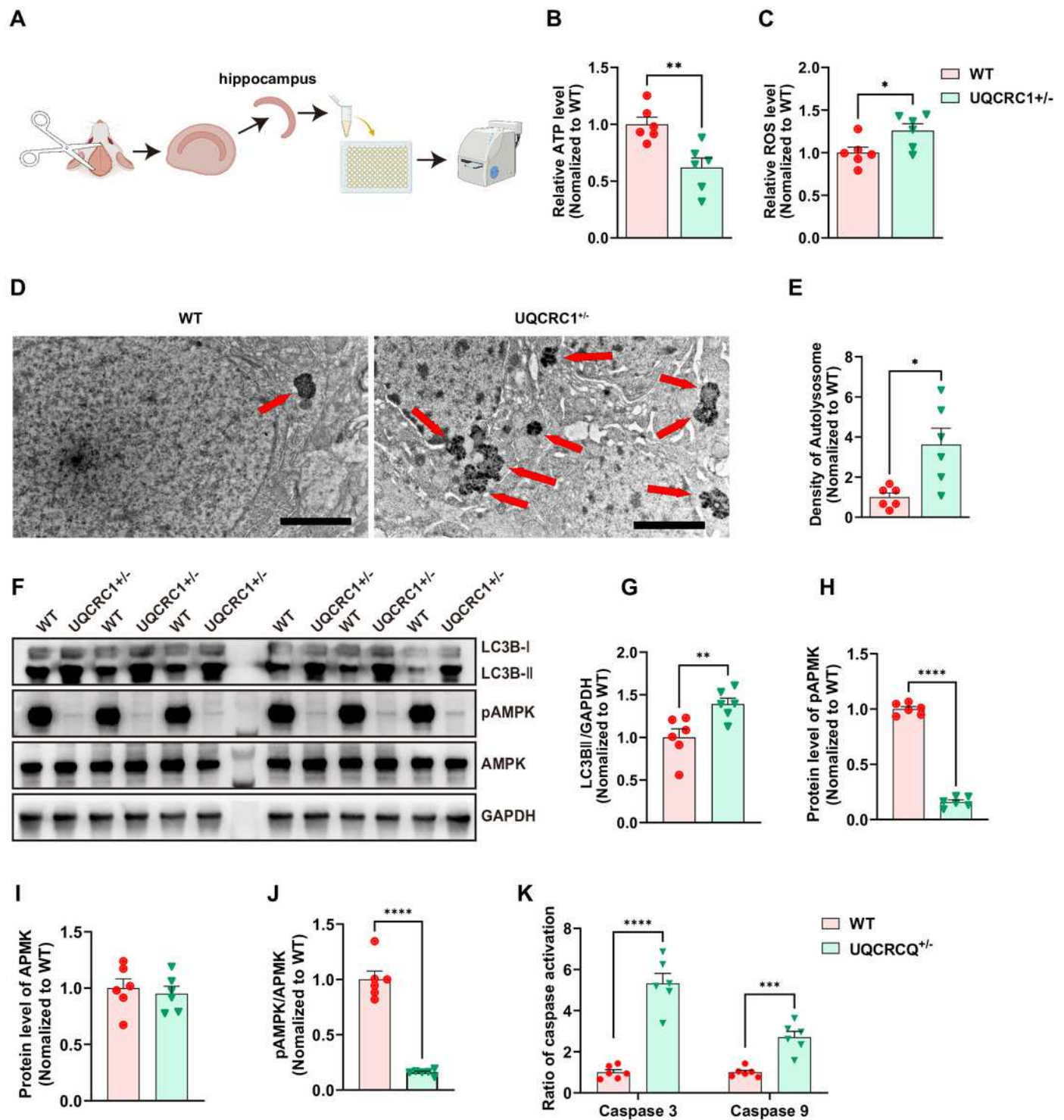
(A) Schematic representation of NOR, created in <https://BioRender.com>. (B) The recognition rate of both male and female *UQCRC1* +/- mice was lower than that of their conspecifics (n=8). (C) Schematic representation of NBT, created in <https://BioRender.com>. (D) The nesting score of male *UQCRC1* +/- mice was lower (n=8). (E) Schematic representation of the Barnes maze, created in <https://BioRender.com>. (F) Compared to male WT mice, male *UQCRC1* +/- mice exhibited longer escape latencies from day 3 during the training phase of Barnes maze (n=8). (G) Male *UQCRC1* +/- mice had prolonged escape latencies during the testing phase of the Barnes maze (n=8). (H) Both male and female *UQCRC1* +/- mice took more attempts to find the correct hole (n=8). Recognition rate in NOR and escape latencies in both the training and testing phases of the Barnes maze were quantified and expressed as mean  $\pm$  standard error of the mean (SEM). Nesting score in NBT and the numbers to identify target box during Barnes maze were recorded and presented using non-parametric descriptors (median with interquartile range). \*\*\*\*p < 0.0001; \*\*p < 0.01; \*p < 0.05.



# Figure 2

The downregulation of *UQCRC1* led to autophagy impairment.

(A) Schematic illustration of the experimental design for ATP assays, reactive oxygen species assessments, and caspase 3 and caspase 9 evaluations, created in <https://BioRender.com>. (B) The ATP level in *UQCRC1* +/- mice was lower. (C) The ROS level in *UQCRC1* +/- mice was higher. (D) Representative transmission electron microscope (TEM) images of WT mice and *UQCRC1* +/- mice (red arrows indicate autolysosomes; scale bar: 2  $\mu$ m). (E) The density of autolysosomes was higher in *UQCRC1* +/- mice. (F) Western blot images of LC3B I, LC3B II, pAMPK, AMPK in WT mice and *UQCRC1* +/- mice. (G-H) Quantification of protein expression of LC3B II (G), pAMPK (H), AMPK (I) and pAMPK/AMPK ratio (J). (K) Activation of both caspase 3 and caspase 9 was higher in *UQCRC1* +/- mice. Data are represented as mean  $\pm$  SEM. \*\*\*\*p < 0.0001; \*\*\*p < 0.001; \*\*p < 0.01; \*p < 0.05.

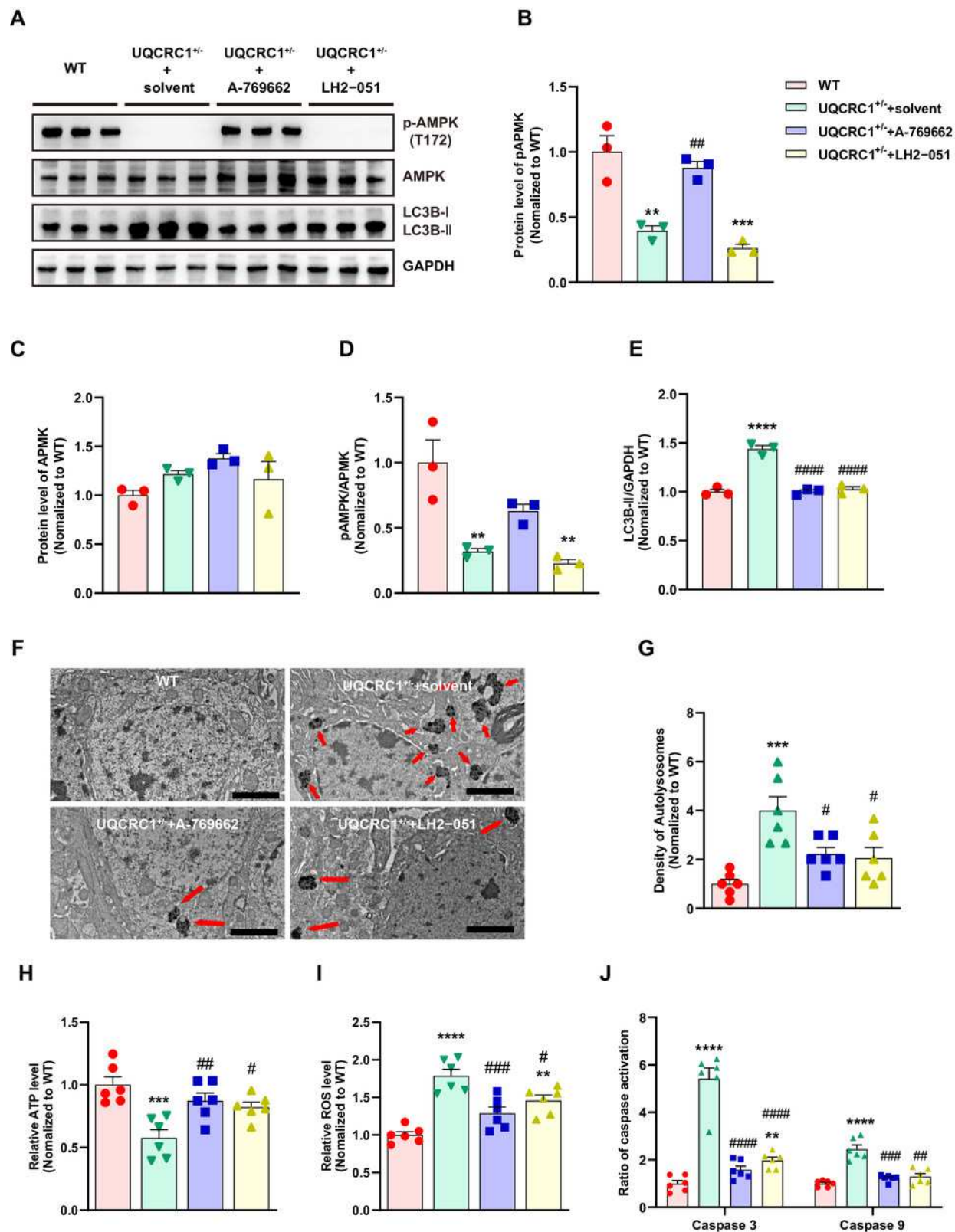




# Figure 3

Both lysosomal function improvement and AMPK activity enhancement ameliorated autophagy in *UQCRC1* mice.

(A) Western blot image of pAMPK, AMPK, LC3B I and LC3B II. (B-E) Quantification of protein expression of pAMPK (B), AMPK (C), pAMPK/AMPK (D) and LC3B II (E) . (F) Representative transmission electron microscope (TEM) images (red arrows indicate autolysosomes; scale bar: 2  $\mu$ m). (G) The density of autolysosomes in *UQCRC1* +/- mice reduced after the administration of A-769662 or LH2-051. (H) The ATP level in *UQCRC1* +/- mice increased after the administration of A-769662 or LH2-051. (I) The ROS levels in *UQCRC1* +/- mice diminished following the treatment with A-769662 or LH2-051; however, after LH2-051 administration, the ROS levels did not return to those observed in WT mice. (J) Following administration of A-769662 or LH2-051, activation of caspase 3 and caspase 9 in *UQCRC1* +/- mice was reduced; however, caspase 3 activation did not recover to WT levels after LH2-051 treatment. Data are represented as mean  $\pm$  SEM. \*\*\*\* p < 0.0001 vs. WT; \*\*\* p < 0.001 vs. WT; \*\* p < 0.01 vs. WT. #### p < 0.0001 vs. *UQCRC1* +/- +solvent; ###p < 0.001 vs. *UQCRC1* +/- +solvent; ##p < 0.01 vs. *UQCRC1* +/- +solvent; #p < 0.05 vs. *UQCRC1* +/- +solvent.



# Figure 4

The cognitive deficits in *UQCRC1* +/- mice were ameliorated by augmenting AMPK activation and improving lysosomal function.

(A) Representative heat maps showing the duration and location of the subject during novel object recognition (“N” and “F” represent novel object and familiar object, respectively). (B) The recognition rate of *UQCRC1* +/- mice increased after the administration of A-769662 or LH2-051. (C) Representative figures of the nest-building test. (D) The nesting score of *UQCRC1* +/- mice increased after the administration of A-769662 or LH2-051. (E) Representative trajectory chart during testing phase of the Barnes maze (gray circle indicates the target hole). (F) Over the training phase of the Barnes maze, the escape latencies of *UQCRC1* +/- mice were restored to levels similar to WT mice. (G) A-769662 or LH2-051 treatment lowered the escape latencies of *UQCRC1* +/- mice during the Barnes maze testing phase. (H) *UQCRC1* +/- mice treated with A-769662 or LH2-051 showed fewer attempts to find the target hole. Recognition rate in NOR and escape latencies in both the training and testing phases of the Barnes maze were quantified and expressed as mean  $\pm$  SEM. Nesting score in NBT and the numbers to identify target box during Barnes maze were presented as median and interquartile range. \*\*\*\*  $p < 0.0001$  vs. WT; \*\*  $p < 0.01$  vs. WT, \*  $p < 0.05$  vs. WT. ###  $p < 0.001$  vs. *UQCRC1* +/- +solvent; ##  $p < 0.01$  vs. *UQCRC1* +/- +solvent; #  $p < 0.05$  vs. *UQCRC1* +/- +solvent.

