### **Supplemental Information**

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#### Targeted Expression of Catalase to Mitochondria Prevents Age-Associated Reductions in Mitochondrial Function and Insulin Resistance

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Figure S1, Related to Figure 1. Antioxidant Protein Expression, Genomic DNA Damage, and Mitochondrial Respiration

(A) Catalase protein expression in various tissues.

(B) Expression of superoxide dismutase 2 (SOD2) and glutathione peroxidase 1 (GPX1) in quadriceps muscle.

(C) Genomic DNA damage assessed by QPCR (n=4-6).

(D and E) State IV oxygen consumption (D) and respiratory control rate of isolated mitochondria from skeletal muscle (E) (n=6). All data are mean  $\pm$  SEM. \*P<0.05; NS, not significant by ANOVA.



# Figure S2, Related to Figure 2. Whole-Body Energy Expenditure, Food Intakes and Locomoter Activity

(A-F) Energy expenditure (A), food intake (C), and locomotor activity (E) during 72 hr analysis; hour to hour average energy expenditure (B), food intake (D), and locomotor activity (F) for old groups during light/dark period. N=8 in young groups; n=12-15 in old groups. All data are mean  $\pm$  SEM. \*P<0.05; NS, not significant by ANOVA.



Figure S3, Related to Figure 4. VDAC Mitochondrial Membrane Protein Expression



**Figure S4, Related to Figure 4. EM of Muscle in Young and Old WT and MCAT Mice** In contrast to the young WT and old MCAT mice intramyofiblilar mitochondria are rarely found in A-band of the old WT mice and most of the mitochondria are singly located within the space between I band and Z-line, while young WT and MCAT mitochondria are located in A-band and connected together across the Z-line. Furthermore, the shape of old WT mitochondria are round or slightly elongated, while young WT and MCAT mitochondria are large and amorphous. Red spots denote mitochondria; far right panel is a close up view of the old WT and MCAT groups.

### **Table S1. Basal Characterization for Animals**

	Young		Old		P-value	P-value
	WT (n=8)	MCAT (n=8)	WT (n=10)	MCAT (n=10)	Young WT vs. old WT	Old WT vs. old MCAT
Age (month)	$6.6 \pm 0.3$	$6.9 \pm 0.3$	15.8 ± 0.1	15.7 ± 0.2	<0.001	NS
Body weight (g)	32.3 ± 0.7	31.6 ± 0.5	$34.9 \pm 0.5$	34.3 ± 1.2	<0.05	NS
Body fat (%)	12.5 ± 1.5	10.4 ± 1.1	13.6 ± 1.0	11.7 ± 1.8	NS	NS
Lean body mass (%)	73.9 ± 1.0	74.9 ± 1.1	$72.4 \pm 0.7$	74.0 ± 1.4	NS	NS
Fasting glucose (mg/dl)	98.1 ± 2.3	103.1 ± 4.0	102.2 ± 4.0	99.1 ± 3.9	NS	NS
Fasting insulin (µU/ml)	3.9 ± 1.3	3.7 ± 1.8	$5.7 \pm 0.6$	6.1 ± 1.8	NS	NS
Fasting FA (mg/dl)	1.5 ± 0.1	$1.4 \pm 0.1$	$1.0 \pm 0.1$	1.1 ± 0.1	<0.05	NS
Fasting EGP [mg/(kg-min)]	$9.5 \pm 0.4$	$11.0 \pm 0.4$	11.5 ± 0.5	$10.4 \pm 0.4$	<0.05	NS
Triglyceride in GAS muscle (mg/g tissue)	7.1 ± 0.9	5.1 ± 1.0	7.6 ± 0.9	5.7 ± 0.8	NS	NS

Data are expressed as mean values  $\pm$  SEM. *P-value* evaluated by one-way ANOVA followed by post hoc analysis using the Bonferroni's Multiple Comparison Test. NS, not significant.

	Young		Old		P-value	P-value
	WT (n=8)	MCAT (n=8)	WT (n=10)	MCAT (n=10)	Young WT vs. old WT	Old WT vs. old MCAT
Clamped glucose (mg/dl)	126 ± 3.1	127 ± 4.3	122 ± 1.9	124 ± 1.6	NS	NS
Insulin (μU/ml)	50.7 ± 3.4	54.9 ± 0.1	67.4 ± 3.9	59.9 ± 0.2	<0.05	NS
FA (mg/dl)	$0.35 \pm 0.07$	$0.29 \pm 0.03$	$0.37 \pm 0.04$	0.27 ± 0.04	NS	NS
FA suppression (%)	76 ± 4.7	77 ± 3.4	63 ± 3.3	73 ± 4.7	<0.05	0.09
EGP [mg/(kg-min)]	8.4 ± 1.0	8.7 ± 1.4	10.9 ± 1.1	$9.0 \pm 0.9$	NS	NS
EGP suppression (%)	10.1 ± 12.8	22.4 ± 11.2	0.4 ± 14.4	12.9 ± 7.9	NS	NS
Glycolysis [mg/(kg-min)]	$36.0 \pm 4.0$	29.1 ± 2.7	28.9 ± 1.4	31.8 ± 2.2	NS	NS
Glycogen synthesis [mg/(kg-min)]	23.3 ± 4.1	20.5 ± 2.7	9.3 ± 1.5	20.2 ± 3.13	<0.01	<0.05

## Table S2. Characterization of Animals during Hyperinsulinemic-Euglycemic Clamp Study

Data are expressed as mean values ± SEM. *P-value* evaluated by ANOVA followed by post hoc analysis using the Bonferroni's

Multiple Comparison Test. NS, not significant