# Angiotensin-converting enzyme 2 deficiency accelerates and angiotensin 1-7 restores age-related muscle weakness in mice

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### Abstract

**Background** A pharmacologic strategy for age-related muscle weakness is desired to improve mortality and disability in the elderly. Angiotensin-converting enzyme 2 (ACE2) cleaves angiotensin II into angiotensin 1-7, a peptide known to protect against acute and chronic skeletal muscle injury in rodents. Since physiological aging induces muscle weakness via mechanisms distinct from other muscle disorders, the role of ACE2-angiotensin 1-7 in age-related muscle weakness remains undetermined. Here, we investigated whether deletion of ACE2 alters the development of muscle weakness by aging and whether angiotensin 1-7 reverses muscle weakness in older mice.

**Methods** After periodic measurement of grip strength and running distance in male ACE2KO and wild-type mice until 24 months of age, we infused angiotensin 1-7 or vehicle for 4 weeks, and measured grip strength, and excised tissues. Tissues were also excised from younger (3-month-old) and middle-aged (15-month-old) mice. Microarray analysis of RNA was performed using tibialis anterior (TA) muscles from middle-aged mice, and some genes were further tested using RT-PCR.

**Results** Grip strength of ACE2KO mice was reduced at 6 months and was persistently lower than that of wild-type mice (p < 0.01 at 6, 12, 18, and 24-month-old). Running distance of ACE2KO mice was shorter than that of wild-type mice only at 24 months of age  $[371\pm26 \text{ vs. } 479\pm24 \text{ (m)}, p < 0.01]$ . Angiotensin 1-7 improved grip strength in both types of older mice, with larger effects observed in ACE2KO mice (% increase,  $3.8\pm1.5$  and  $13.3\pm3.1$  in wild type and ACE2KO mice, respectively). Older, but not middle-aged ACE2KO mice had higher oxygen consumption assessed by a metabolic cage than age-matched wild-type mice. Angiotensin 1-7 infusion modestly increased oxygen consumption in older mice. There was no difference in a wheel-running activity or glucose tolerance between ACE2KO and wild-type mice and between mice with vehicle and angiotensin 1-7 infusion. Analysis of TA muscles revealed that *p16INK4a*, a senescence-associated gene, and central nuclei of myofibers increased in middle-aged, but not younger ACE2KO mice. *p16INK4a* and central nuclei increased in TA muscles of older wild-type mice, but the differences between ACE2KO and wild-type mice remained significant (p < 0.01). Angiotensin 1-7 did not alter the expression of p16INK4a or central nuclei in TA muscles of both types of mice. Muscle *ACE2* expression of wild-type mice was the lowest at middle age (2.6 times lower than younger age, p < 0.05).

**Conclusions** Deletion of ACE2 induced the early manifestation of muscle weakness with signatures of muscle senescence. Angiotensin 1-7 improved muscle function in older mice, supporting future application of the peptide or its analogues in the treatment of muscle weakness in the elderly population.

Keywords Muscle weakness; ACE2; Angiotensin 1-7; p16INK4a

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### Introduction

Over the last few decades, skeletal muscle function has been recognized as a determinant of health condition and lifespan in the elderly population. Muscle weakness is a key component of two important age-related pathological conditions, sarcopenia and frailty, both of which are closely associated with mortality and disability in the elderly, causing social and economic burden worldwide.<sup>1</sup> Research efforts have been devoted to developing a therapeutic strategy for improving muscle weakness in the elderly. While physical training with nutritional support is the practical method for improving muscle weakness, its effect is limited because of impaired physical performance and fatigability of the fragile elderly population.<sup>2,3</sup> Thus, pharmacotherapy for age-related muscle weakness is a suitable alternative option. However, to date, no drugs have been found to improve muscle weakness in medical practice.

The mechanisms underlying muscle weakness are multifactorial and may involve multiple molecular pathways. Accumulating evidence suggests that the renin-angiotensin (RA) system, a main regulator of blood pressure, is involved in maintaining skeletal muscle homeostasis, making it a potential therapeutic target for muscle disorders.<sup>4</sup> Burks *et al.* reported that angiotensin II type 1 receptor blockade with the drug losartan improved muscle regeneration after muscle injury in mouse models of Marfan syndrome and Duchenne muscular dystrophy (DMD).<sup>5</sup> They also reported the preventive role of losartan against muscle regeneration after injury and immobilization-induced disuse atrophy in ageing mice.<sup>6</sup> Furthermore, another group reported that losartan treatment improved muscle strength and ameliorated fibrosis in a mouse model of congenital muscular dystrophy.<sup>7</sup>

The activation of the endogenous protective axis in the RA system is also a potential therapeutic target for muscle disorders. Angiotensin-converting enzyme 2 (ACE2) cleaves angiotensin II to angiotensin 1-7 (A1-7), a heptapeptide shown to have a protective role in multiple pathologies by blocking over-activation of the RA system. Recently, several studies have revealed that A1-7 attenuates muscle dysfunction in animal model.<sup>8–13</sup> For example, Acuña et al. reported that A1-7 decreases fibrosis and improves muscle function in DMD mice.<sup>10</sup> In addition, the genetic deletion or pharmacological inhibition of the A1-7 receptor Mas resulted in deteriorated muscular architecture and increased fibrosis with diminished muscle strength in DMD mice.<sup>10</sup> Therefore, it is currently recognized that pharmacological inhibition of the RA system and activation of the A1-7-Mas pathway contribute similarly to the improvement of muscle disorders in rodents. Nevertheless, caution should be taken when applying the current concept of the RA system to the treatment of ageing-associated muscle weakness, as the development of ageing-associated muscle weakness is distinct from that of other muscle disorders. For example, repair from acute or chronic injury in most muscle disorders largely depends on the proliferative capacity of skeletal muscle stem cells or satellite cells. However, the role of satellite cells in the development of ageing-associated muscle weakness remains controversial.<sup>14–16</sup> In addition, the possible improvement of ageing-associated muscle weakness by A1-7, believed to protect against over-activation of the RA system, remains unknown. To address this, we used ACE2 knockout (ACE2KO) and wild-type mice to investigate (i) whether ACE2 plays a pivotal role in ageing-associated changes in muscle function and (ii) whether A1-7 improves muscle weakness in ageing mice.

### Materials and methods

#### Experimental design

Experiments were performed using male ACE2KO mice (C57bl/6J background) and their wild-type littermates. The mice were fed standard chow and water ad libitum. The study consists of two lines of assays: longitudinal assay for observing muscle function in terms of the temporal change over the lifetime and the effect of A1-7 in older mice and cross-sectional assay for single time-point analysis using younger (3-month-old) and middle-aged (15-month-old) mice (Figure 1). For the longitudinal assay, grip strength, running endurance, and body weight were measured at 3, 6, 12, 18, and 24 months of age. Thereafter, A1-7 (4332-v; PEPTIDE INSTITUTE, Osaka, Japan) in 0.9% saline or vehicle (0.9% saline) was infused for 4 weeks at a dose of 0.1  $\mu$ g/kg body weight/min via a subcutaneously implanted osmotic minipump (Alzet model 1004; Durect, Birmingham, AL, USA) according to the previous study.<sup>17</sup> Three weeks after infusion, mice were subjected to an intraperitoneal glucose tolerance test (ipGTT), assessment of respiratory quotient using a metabolic cage, and assessment of physical activity using a running wheel. Finally, grip strength was measured, and tissues were extracted for further analysis. For the crosssectional assay, the ipGTT and assessment of respiratory quotient and grip strength were performed for experiments using middle-aged (15-month-old) mice. Thereafter, in situ analysis of tibialis anterior (TA) muscles was performed, and tissues were extracted for further analysis. For experiments using young (3-month-old) mice, tissues were extracted for further analysis. We measured food intake in 8-month-old mice for 7 days consecutively as previously described.<sup>18,19</sup> Briefly, we separated the mice one per cage and gave them preweighted amount of food. Seven days after, we weighed the remaining food and calculated their food intake. All experiments were approved by the Animal Care and Use Committee of Osaka University and were conducted in strict accordance with the US National Institutes of Health Guide for the Care and Use of Laboratory Animals.

Figure 1 The schematic overview of the experimental protocol. ACE2KO, angiotensin-converting enzyme 2 knockout; A1-7, angiotensin 1-7; ipGTT, intraperitoneal glucose tolerance test; RT-PCR, real-time PCR.

Male ACE2KO mice Male wild-type mice Longitudinal assay	3	6	12	15	18	24 25	Continuous Infusion of Angiotensin 1-7 or vehicle
<ul> <li>Grip strength</li> </ul>	0	0	0		0	00	
<ul> <li>Treadmill test</li> </ul>	0	0	0		0	0	
ipGTT						0	
<ul> <li>Metabolic cage</li> </ul>						0	
<ul> <li>Running wheel</li> </ul>						0	
<ul> <li>Histological analysis</li> </ul>						0	
<ul> <li>RNA expression</li> </ul>						0	
assay (rtPCR)						0	
Cross-sectional assay							
<ul> <li>Grip strength</li> </ul>				0			
<ul> <li>In situ assessment of</li> </ul>				$\sim$			
muscle contraction				0			
ipGTT				0			
<ul> <li>Metabolic cage</li> </ul>	_			0			
<ul> <li>Histological analysis</li> </ul>	0			0			
RNA microarray				0			
<ul> <li>RNA expression assay (rtPCR)</li> </ul>	0			0			

#### Physical activity tests

Physical activity was assessed by measurement of forelimb grip strength and running endurance. Forelimb grip strength was measured using a recently modified method in order to improve validity and reliability.<sup>20</sup> In brief, mice grasping the bar were pulled vertically by an inspector, and peak pull force was recorded on a digital force transducer (GPM-100; Melquest, Toyama, Japan). The gauge recorded tension at the time the mouse released its forepaws from the bar. We performed six consecutive measurements for 2 days, and the average of 12 measurements was used to represent grip strength for each animal. All test sessions were performed during the afternoon hours of the light cycle (11 a.m. to 5 p.m.) in the vivarium where the animals were housed. To assess running endurance, we used a treadmill system (Exer-3R; Columbus Instruments, Columbus, OH, USA). The procedure was performed as previously described.<sup>21</sup> In brief, after acclimation to moderate treadmill running for 5 days, we measured the running time until mice became exhausted and stopped running on the sixth day. The total running distance was calculated by multiplying running time by the speed of the treadmill system.

# RNA extraction, gene expression microarrays, and quantitative real-time PCR-based gene expression analysis

Total RNA was isolated from 0.1 g of mouse TA muscles with an RNeasy tissue fibrosis mini kit (74704; QIAGEN, Germany). Whole genome transcriptome profiling was assessed using a microarray system (SurePrint G3 Mouse 8 × 60K v2; Agilent, Santa Clara, CA, USA). For real-time PCR (RT-PCR), cDNA synthesis was performed with a ReverTra Ace qPCR RT kit (FSQ-101; TOYOBO, Osaka, Japan) according to the manufacturer's instructions. All genes were evaluated by the SYBR green qPCR system (Applied Biosystems 7900HT Fast Real-Time PCR System; Thermo Fisher Science, Waltham, MA, USA). Relative expression was calculated using the  $\Delta\Delta$ Ct method with normalization to constitutive genes, as indicated in each figure.

# *Evaluation of contractile properties of the tibialis anterior muscle in situ*

Tibialis anterior muscle contractile properties were measured as previously described.<sup>20</sup> Briefly, each mouse was anaesthetized and placed prone on a heated platform to maintain body core temperature at 37 °C. The sciatic nerve was carefully dissected, tied to a suture, and cut at the proximal end. The distal tendon of the TA muscle was tied at the muscle tendon junction using a 2-0 suture line, and the other end of the suture was tied to the load cell (UTA-100GR; MinebeaMitsumi, Nagoya, Japan). The contractile properties of the TA muscle were measured using the load cell connected to a directcoupled amplifier (SC20AZ-2; Labo support, Osaka, Japan), and the output was recorded using computer hardware (Power Lab; AD Instruments, Bella Vista, Australia). Optimal muscle length (Lo) and absolute twitch force (Pt) were determined with a resting tension between 10 and 20 g, when twitch force in response to 1 Hz supramaximal stimulation to the sciatic nerve was maximal. Maximal isometric tetanic force (Po) was determined in response to supramaximal stimulation between 50 and 200 Hz at Lo. Muscle fatigue was induced by repeated supramaximal stimulation once per second with 100 ms trains at a frequency of 100 Hz for

5 min. The data were recorded and analysed using Chart software version 5 (AD Instrument). The muscle cross-sectional area was determined on the basis of a muscle density of  $1.06 \text{ g/cm}^3$  and a fibre length of the Lo ratio of 0.61.

#### Intraperitoneal glucose tolerance test

Mice were fasted for 16 h and were anaesthetized with medetomidine (0.3 mg/kg), midazolam (4 mg/kg), and butorphanol (5 mg/kg). Blood was collected from the tip of the tail vein, and glucose levels were monitored with a glucometer at the indicated time points after intraperitoneal injection of glucose (2 g/kg body weight) for ipGTT. ACE2KO and wild-type mice that received the same treatment sequentially underwent measurements on the same day.

#### Histochemistry

After excision, TA muscles of mice were immediately frozen in liquid nitrogen-cooled isopentane, covered in embedding agent (Tissue-Tek O.C.T. Compound; Sakura Finetek, Torrance, CA, USA), and stored at  $-80^{\circ}$ C until cryosectioning (10  $\mu$ m). Basic muscle morphology was assessed with haematoxylin and eosin staining, according to standard protocol. All stained, sectioned images were captured under a microscope (Keyence BZ-X700; Keyence, Osaka, Japan) with a minimum of four fields of view per muscle section at ×20 magnification. All histological images were analysed using ImageJ software (ImageJ version 1.51; NIH, Bethesda, MD, USA) as previously described.<sup>22</sup> We counted muscle fibre cross sections with central nuclei and normalized to the number of muscle fibres, as previously reported.<sup>23</sup>

# Assessment of oxygen consumption, carbon dioxide elimination, and physical activity

Oxygen consumption and carbon dioxide elimination were measured using an O<sub>2</sub>/CO<sub>2</sub> metabolism measuring system for small animals (MK-5000RQ; Muromachi Kikai, Tokyo, Japan). Mice were placed in the calorimeter chambers and stabilized overnight, and we used data from the next light/dark cycle. In these chambers, mice were able to eat and drink *ad libitum*. The amount of exercise was measured using a wireless running wheel system (ENV-044&SOF-860; Med Associates Inc., St Albans City, VT, USA). We placed the running wheel in the ordinary mouse cage and counted the number of rotations of the running wheel.

#### Statistical analysis

All data are presented as the means  $\pm$  SEM. Temporal changes of the physical tests in individual mice were analysed

with a paired *t*-test. Differences between two independent groups were analysed by Student's *t*-tests. Differences among wild-type mice with vehicle and A1-7 and ACE2KO mice with vehicle and A1-7 were analysed with two-way ANOVA and Bonferroni testing to determine the difference between the genotypes and between the treatments. Difference in *ACE2* expression among wild-type mice with different ages was analysed with one-way ANOVA and Bonferroni testing (*Figure* 6).

#### Results

#### Angiotensin-converting enzyme 2 knockout mice exhibited more prominent age-dependent declines in motor function than wild-type mice

For the longitudinal assay, mice were fed a normal diet with periodic tests of physical strength until 24 months of age (Figure 1). There was no difference in grip strength between wild-type and ACE2KO mice at 3 months of age (Figure 2A). ACE2KO mice had significantly reduced grip strength at 6 and 12 months of age compared with the previous time point, while wild-type mice had reduced grip strength at 24 months of age, suggesting that age-associated declines in muscle function progressed earlier in ACE2KO mice. The grip strength of 6-month-old ACE2KO mice was lower than that of wild-type mice of the same age, and the difference remained significant until 24 months of age. Running distance in the treadmill test gradually decreased with age in both types of mice, and a difference between the groups was observed only at 24 months of age (Figure 2B). Body mass increased less during ageing in ACE2KO mice than in wild-type mice. ACE2KO mice had lower body mass than wild-type mice at 12 and 18 months of age, while the difference disappeared at 24 months of age because of reduction in body weight in the wild-type mice (Figure 2C). During the whole experiments, 2 of the 33 ACE2KO mice and 5 of the 32 wild-type mice died naturally before 24 months of age.

# Angiotensin 1-7 restored age-associated muscle weakness in mice

As shown in *Figure* 3A, 4 weeks infusion of A1-7 but not vehicle increased the grip strength of ACE2KO mice at 24 months of age. A1-7 tended to increase the grip strength of wild-type mice, but the difference did not reach statistical significance (P = 0.06). The percentage change of grip strength after A1-7 infusion was higher than that after vehicle infusion in both ACE2KO and wild-type mice (*Figure* 3B). Grip strengths of ACE2KO and wild-type mice were similar after A1-7 infusion and were equivalent to that in 6-month-old ACE2KO mice and 12-month-old wild-type mice.

**Figure 2** Serial changes in (A,B) motor function and (C) body mass during the ageing process in angiotensin-converting enzyme 2 knockout (ACE2KO) and wild-type mice. (A) Forearm grip strength was measured six times per day for 2 days, and the data were averaged as individual values. (B) Running distance was measured with the treadmill test. The number of animals was 25 and 24 in wild-type mice and ACE2KO mice, respectively. Difference from the previous test was analysed with a paired *t*-test, and difference between the genotypes was analysed with a Student's *t*-test. \*p<0.01 vs. age-matched ACE2KO mice, †p<0.05 vs. the previous test.



**Figure 3** Effect of A1-7 infusion on grip strength of aged mice. Forearm grip strength was measured before and after infusion of A1-7 (100 ng/kg/min) and vehicle for 4 weeks in 24-month-old mice. (A) Individual value before (pre) and after (post) infusion. Differences were analysed with a paired *t*-test. (B) Percentage change after infusion. Differences were analysed with a Student's *t*-test. ACE2KO, angiotensin-converting enzyme 2 knockout; A1-7, angiotensin 1-7.



#### Angiotensin-converting enzyme 2 deletion and angiotensin 1-7 did not affect tissue mass, glucose tolerance, and wheel-running activities

We found that ACE2 deficiency did not alter the mass of the fast-twitch TA muscle or slow and fast-twitch gastrocnemius

muscle at the age of 3, 15, and 25 months (*Figure* 4A). We previously reported that ACE2 deficiency exacerbated and A1-7 infusion improved muscle insulin resistance induced by a high-fat diet.<sup>17</sup> However, ACE2 deficiency did not alter glucose tolerance in middle-aged and older mice fed a normal diet (*Figure* 4B). ACE2 deficiency did not alter voluntary

**Figure 4** Tissue weight, metabolic function, and physical activity in aged mice. (A) Wet weight of average tibialis anterior (TA) muscle and average gastrocnemius medialis (GM) muscle adjusted by body weight at the age of 3 months (n = 3-4), 15 months (n = 5), and 25 months (n = 8-9). (B) Intraperitoneal glucose tolerance test (ipGTT) was performed by intraperitoneal injection of glucose (2 g/kg body weight) under anaesthesia after 16 h of fasting at the age of 15 months (n = 6-7) and 25 months (n = 4-7). Left panel indicates serial change in blood glucose level, and right panel represents area under the curve (AUC). (C) Physical activity was assessed with a running wheel system at the age of 25 months. n = 3-6 in each group. (D) Oxygen consumption (left panel) and carbon dioxide elimination (right panel) were assessed with a metabolic cage at the age of 15 months (n = 6-7) and 25 months (n = 7-9). (E) Food consumption during 7 days in 15-month-old angiotensin-converting enzyme 2 knockout (ACE2KO) and wild-type (WT) mice. Differences between genotypes at the age of 3 and 15 months were analysed with Student's *t*-test. Differences between the treatments.



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wheel-running activity in older mice (*Figure* 4C). A1-7 infusion also did not alter muscle mass, glucose tolerance, and the wheel-running activity in older mice.

#### Angiotensin-converting enzyme 2 deletion and angiotensin 1-7 increased oxygen consumption in older mice

There was no difference in oxygen consumption and carbon dioxide elimination assessed by metabolic cages in middleaged ACE2KO and wild-type mice (*Figure* 4D). In contrast, older ACE2KO mice had higher oxygen consumption and carbon dioxide elimination in both light and dark cycle than older wild-type mice (*Figure* 4D). A1-7 infusion to older mice increased oxygen consumption significantly in light cycle and non-significantly (P = 0.07) in dark cycle. A1-7 infusion did not increase carbon dioxide elimination in both light and dark cycle (*Figure* 4D). Food consumption measured in 8-month-old mice did not differ between the genotypes (*Figure* 4E).

#### A senescence-associated gene, p16INK4a, increased in skeletal muscle of angiotensin-converting enzyme 2 knockout mice

We performed RNA microarray analysis using TA muscles from middle-aged mice to clarify alteration of the gene

expression profile induced by ACE2 deletion. While middleaged ACE2KO mice had lower forelimb grip strength than age-matched wild-type mice, the contractile force of the TA muscle did not differ between ACE2 and wild-type mice (Figure 5A and 5B). Fatigue of the TA muscle during repetitive contractions was more prominent in ACE2KO than wild-type mice (Figure 5C). RNA microarray analysis revealed that, among the total 42 405 coding and non-coding RNA analysed, only 14 genes in TA muscles of ACE2KO mice increased or decreased more than two-fold compared with those of wildtype mice (P < 0.01; Table 1). Representative genes including those with high false discovery rate (q-value) were validated by RT-PCR using different samples from 15-month-old mice (Npnt, Pdgfd, MYBPH, p16INK4a, and Ppp1r1a). Consequently, the results confirmed the difference between the genotypes, with the exception of MYBPH (Figure 6). Among the genes, the largest difference was observed in p16INK4a, known as an inducer of cellular senescence by promoting cell cycle arrest. We found that p53 and p21, which are involved in other cellular senescence pathways, were unaltered and decreased in ACE2KO mice compared with wild-type mice, respectively. We also confirmed that p19, encoded by the same genetic locus as p16INK4a, was not altered by ACE2 deficiency. We did not find differences in expression of the tested genes in TA muscles of young mice at 3 months of age (Figure 6). In the older mice after vehicle infusion, the difference in expression between ACE2KO and wild-type mice

**Figure 5** Grip strength and contractile force of tibialis anterior (TA) muscle in middle-aged mice. (A) Forearm grip strength of 15-month-old mice. (B) Pt (absolute twitch force) and Po (maximal isometric tetanic force) in the TA muscles of 15-month-old mice. (C) Change in force during fatigue-inducing conditions in the TA muscles. Values were compared by using Student's *t*-test and are expressed as the mean  $\pm$  SEM. *n* = 6 in each group.



Genbank accession	Gene symbol	Wild type	ACE2KO	Fold change (log <sub>2</sub> )	P-value	<i>q</i> -value
NM 033525	Npnt	38.3 ± 2.5	102.7 ± 3.4	1.42	0.0001	0.043
XM_006509885	Pdgfd	152.8 ± 6.6	41.8 ± 3.4	-1.87	0.0001	0.043
NM_016749	Mybph	1305.1 ± 71.7	278.4 ± 100	-2.23	0.0011	0.206
NM_009877	Cdkn2a(p16INK4a)	27.5 ± 5.7	369.9 ± 47.1	3.75	0.0020	0.264
NR 028261	Rian	425.0 ± 32.6	198.1 ± 5.4	-1.1	0.0024	0.264
NR_040707	1110046J04Rik	650.9 ± 36.1	1349.9 ± 117.2	1.05	0.0047	0.429
NM_021391	Ppp1r1a	634.4 ± 62.7	272.5 ± 2.52	-1.22	0.0059	0.434
NR_045633	A230056J06Rik	15.9 ± 3.6	41.4 ± 3.5	1.38	0.0069	0.434
NM 001291046	Nudt3	21.7 ± 1.8	$44.6 \pm 4.4$	1.04	0.0085	0.434
NM_007492	Arx	61.6 ± 10.3	127.5 ± 9.0	1.05	0.0086	0.434
NM_173396	Tgif2	148.0 ± 7.8	68.1 ± 15.2	-1.12	0.0097	0.434
XM_011247076	Cd226	42.5 ± 3.3	102.3 ± 12.4	1.27	0.0098	0.434
NM_009014	Rad51b	46.5 ± 5.9	17.6 ± 1.9	-1.4	0.0098	0.434
XR_406760	Mus musculus predicted gene 5590 (Gm5590)	$44.9 \pm 6.7$	93.1 ± 8.1	1.05	0.0100	0.434

Table 1 List of genes with  $\log_2$  fold change > |1| in expression and P < 0.01 between TA muscles of 15-month-old ACE2KO and wild-type mice in microarray analysis

ACE2KO, angiotensin-converting enzyme 2 knockout; TA, tibialis anterior.

n = 3, each group. *P*-value and *q*-value were calculated using a Student's *t*-test and a Benjamini–Hochberg procedure, respectively. Genes filled with grey were tested by quantitative real-time PCR analysis.

remained significant only for *p16INK4a*, which was significantly increased compared with middle-aged mice. A1-7 infusion did not alter expression of any of the tested genes in the older ACE2KO and wild-type mice (*Figure* 6). Finally, *ACE2* reduced significantly in 15-month-old wild-type mice compared with 3-month-old mice, while the expression non-significantly reincreased at 25 month. A1-7 did not alter *ACE2* expression in 25-month-old wild-type mice.

#### Central nuclei in tibialis anterior muscle fibres were frequently observed in angiotensin-converting enzyme 2 knockout mice

Histological analysis of TA muscles revealed no significant differences in the cross-sectional area of myofibres between middle-aged ACE2KO and wild-type mice (Figure 7A). There was also no increase in fibrosis and immune cell infiltration in TA muscles from ACE2KO mice. The only difference found in the histological analysis was an increase in central nuclei in muscle fibres from middle-aged ACE2KO mice, which were rarely observed in wild-type mice of the same age (Figure 7B). Central nuclei were not increased in the TA muscle from younger ACE2KO mice (Figure 7B). Central nuclei of myofibres increased in the older wild-type mice compared with the middle-aged mice (P < 0.01). The difference between the genotypes remained significant in the older mice (Figure 7B). A1-7 infusion did not alter the cross-sectional area or central nuclei in the older ACE2KO and wild-type mice (Figure 7B).

### Discussion

In this study, we found that ACE2 deficiency resulted in the early manifestation of muscle weakness in mice.

Comprehensive analysis of the TA muscle in middle-aged (15-month-old) mice revealed that the senescence-associated gene *p16INK4a* and central nuclei of muscle fibres were increased in ACE2KO mice. We found that a continuous A1-7 infusion improved muscle strength in older ACE2KO and wild-type mice without any alternation in the senescent phenotypes of skeletal muscle. To the best of our knowledge, this is the first study to show that ACE2 may influence the physiological process of skeletal muscle ageing and that A1-7 may restore ageing-associated muscle weakness.

Glucose intolerance and insulin resistance are closely related to ageing-associated muscle weakness.<sup>24</sup> We previously found that ACE2 deficiency exaggerated glucose tolerance and insulin resistance in mice fed a high-calorie diet.<sup>17</sup> However, in the present study, we did not find any differences in glucose tolerance between ACE2KO and wild-type mice fed a normal diet until old age. A1-7 infusion in older mice also did not alter glucose tolerance, suggesting that insulin sensitivity is not associated with the influence of ACE2 deficiency and A1-7 infusion on muscle function.

We found that ACE2KO mice exhibited smaller body weights than wild-type mice without affecting food consumption and muscle mass in lower limb at middle age. This implies the possibility that ACE2 deficiency alters metabolic function, leading to change in body composition that we did not measure in this study. However, the change in oxygen consumption between the genotypes in the older mice but not in the middle-aged mice implicates that long-term ACE2 deficiency somehow alters metabolic function, but the effect is not associated with the early manifestation of muscle senescence in ACE2KO mice. We found that A1-7 infusion modestly increased oxygen consumption in older mice without affecting wheel-running activity. Further investigation will be required to elucidate the precise mechanisms, but our findings suggest that A1-7 increases systemic metabolism together with the improvement of muscle function in aged mice.

**Figure 6** Gene expression of tibialis anterior muscle. Gene expression relative to *GAPDH* was analysed by quantitative real-time PCR analysis of tibialis anterior muscles in mice at the age of 3 months (n = 5-8) and 15 months (n = 5-6) and at the age of 25 months (n = 6-9). Expression level was relative to that of 15-month-old wild-type mice. *MYBPH, p53*, and *p19* in which no differences were observed between 15-month-old ACE2KO and wild-type mice were not tested in 25-month-old mice. For the genes except *ACE2*, differences between genotypes at the age of 3 and 15 months were analysed with Student's *t*-test. Differences among groups at the age of 25 months were analysed with two-way ANOVA to detect the difference between the genotypes and the difference between the treatments. For *ACE2*, the difference among wild-type mice with different ages was analysed with one-way ANOVA, and the difference between the treatments in 25-month-old mice was analysed with Student's *t*-test. ACE2KO, angiotensin-converting enzyme 2 knockout; A1-7, angiotensin 1-7; WT, wild-type.



**Figure 7** Histological analysis of tibialis anterior muscle. (A) Representative pictures and (B) central nuclei per muscle fibre (%) and cross-sectional area of single muscle fibres ( $\mu$ m<sup>2</sup>) of haematoxylin–eosin-stained cross sections of tibialis anterior muscles in mice at the age of 3 and 15 months and at the age of 25 months. Circles indicate the centrally located nuclei. *n* = 4–6 in each group. Differences between genotypes at the age of 3 and 15 months were analysed with Student's t-test. Differences among groups at the age of 25 months were analysed with two-way ANOVA to detect the difference between the genotypes and the difference between the treatments. \**P* < 0.01 vs. 15-month-old wild-type mice by a Student's t-test. ACE2KO, angiotensin-converting enzyme 2 knockout; A1-7, angiotensin 1-7; WT, wild-type.



The only specific morphological finding observed was a prominent increase in central nuclei of muscle fibres from ACE2KO mice. It is well recognized that central nuclei appear during the regeneration process of injured muscle in several pathologies, including trauma and myopathies. Central nuclei in skeletal muscle also increase to a small extent during physiological ageing in mice, consistent with our finding that ~3% of muscle fibres had central nuclei in 25-month-old wild-type mice.<sup>23</sup> While muscle stem cells (i.e. satellite cells) are considered the origin of central nuclei in the regeneration process, the pathophysiological basis of central nuclei in aged muscle remains largely undetermined. It has been known that muscle denervation is associated with increased central nuclei.<sup>25</sup>

However, it is unlikely that decreased innervation, if existed in ACE2KO mice, solely caused the increased central nuclei in TA muscle, because decreased innervation should alter muscle size as well.<sup>25</sup> Because of technical reasons, we used TA muscles as a representative fast-twitch muscle for morphological and gene analysis instead of muscles that generate forearm grip strength. The discrepancy between the reduced forearm grip strength and the unaltered contractile force and muscle size of TA muscles in 15-month-old ACE2KO mice suggests that the magnitude of impact of ageing on muscle performance and morphology differs among muscles. Therefore, it is conceivable that the reduced grip strength in ACE2KO mice is not caused by the increased central nuclei in myofibres but by other deteriorations in muscle morphology of muscles promoting forearm grip strength.

While there were few group differences in gene clusters of TA muscles between middle-aged ACE2KO and wild-type mice in the RNA array analysis, the expression of a tumour suppressor protein, p16INK4a, also known as cyclin-dependent kinase inhibitor 2A, was dramatically increased by ACE2 deficiency. The difference between the genotypes remained significant until old age. p16INK4a prevents mammalian cells from oncogenic stimuli by promoting retinoblastoma-dependent cell cycle arrest, which also contributes to systemic ageing in mammals.<sup>26</sup> In ageing skeletal muscle, p16INK4a is mainly expressed in satellite cells, or muscle stem cells, which surround the plasma membrane of muscle fibres. Senescent satellite cells expressing p16INK4a lose their ability to proliferate and differentiate upon muscle injury, yet it remains controversial if and how this dysfunction of satellite cells causes the development of sarcopenia.<sup>14–16</sup> Recent findings in mice support the contribution of satellite cells to ageing-associated changes in muscle fibrosis and myofibre maintenance<sup>14,27</sup> but not to the ageingassociated loss of skeletal muscle size and strength, that is, sarcopenia.<sup>16</sup> In contrast, Baker *et al.* reported that clearance of p16INK4a-positive cells in skeletal muscle increased exercise ability in mice.<sup>15</sup> It was also recently reported that p16INK4a is not a marker of senescence but has a causal role in the dysfunction of aged satellite cells.<sup>28</sup> In this study, we found that ACE2 deficiency was associated with early induction of p16INK4a in skeletal muscle without alteration of other senescence genes, including p53 and p21. Further investigation is required to determine whether the increased p16INK4a seen in ACE2KO mice is a marker or inducer of organ ageing.

While continuous A1-7 infusion improved grip strength in older ACE2KO and wild-type mice, A1-7 did not appear to reverse skeletal muscle ageing, as shown by the lack of alternation in p16INK4a expression and central nuclei with A1-7 infusion. One explanation of this finding is that decreased A1-7 caused by ACE2 deficiency induced skeletal muscle ageing, and A1-7 improved the contractile function of muscle fibres but not the irreversible senescence of satellite cells and muscle fibres. Further investigation will be required to elucidate whether A1-7 infusion to younger ACE2KO mice could restore the muscle senescence. Another possibility is that the accelerated skeletal muscle ageing seen in ACE2KO mice was induced via an A1-7-independent pathway. In addition to its function in the RA system, ACE2 interacts with the apelin system<sup>29</sup> and functions as a collectrin homologue in intestinal tissue.<sup>30,31</sup> In this context, while we found that ACE2 expression reduced in TA muscles of middle-aged wild-type mice, it remains unclear whether the alternation in muscle ACE2 is responsible for age-related muscle weakness. Further studies, including those that utilize mice lacking Mas, a receptor of A1-7 or mice with tissue-specific deletion of ACE2, will be required to further elucidate these possible mechanisms. The protective role of A1-7 in skeletal muscle function has been shown with the use of several mouse models, including the chronic infusion of angiotensin II,<sup>11,13</sup> endotoxin exposure,<sup>12</sup> disuse atrophy,<sup>8</sup> and genetic induction of DMD.<sup>10</sup> Underlying mechanisms proposed in these studies regarding the effects of A1-7 include phosphorylation of AKT, prevention of the increase in atrogin-1 and MuRF-1,<sup>9,11</sup> reduction in phosphorylation of p38 MAPK, and inhibition of transforming growth factor- $\beta$  signalling.<sup>10,12,13</sup> We did not find any influence of A1-7 on these signalling pathways in TA muscles of aged mice (data not shown). We also found no change in mass and myofibril size of TA muscle by A1-7 infusion. The improved grip strength and the lack of alteration in TA muscle might imply that the impact of A1-7 is different among skeletal muscles. Further investigation will be required to elucidate the underlying mechanisms involved in the improvement of muscle weakness by A1-7 in aged mice.

Despite the early manifestation of senescent phenotypes, ACE2KO mice exhibited a normal lifespan. This is in contrast to conventional mouse models of senescence, including klotho mice and a series of senescence-accelerated mice, which die earlier than normal mice.<sup>32,33</sup> Whole-body analysis of ageing phenotypes in ACE2KO mice will be required to clarify the mechanisms of early organ senescence with the normal lifespan. Finally, it is clinically relevant that A1-7 improved ageing-induced skeletal muscle weakness not only in ACE2KO mice but also in wild-type mice. Currently, there is no established pharmacological therapy for the treatment of ageing-associated muscle weakness. A1-7 has been shown to play a protective role in multiple organs without any adverse effects in rodents.<sup>34,35</sup> Further investigation will be required to investigate a fascinating question whether a longer administration of A1-7 would extend lifespan in older mice. Finally, our results provide an important basis for the future clinical application of A1-7 and its analogues as safe drugs for the improvement of muscle weakness in the elderly population.

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### **Conflict of Interest**

All authors declare that they have no conflict of interest regarding the publication of this article.

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