Exercise-stimulated interleukin-15 is controlled by AMPK and regulates skin metabolism and aging

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Summary

Aging is commonly associated with a structural deterioration of skin that compromises its barrier function, healing, and susceptibility to disease. Several lines of evidence show that these changes are driven largely by impaired tissue mitochondrial metabolism. While exercise is associated with numerous health benefits, there is no evidence that it affects skin tissue or that endocrine muscle-to-skin signaling occurs. We demonstrate that endurance exercise attenuates age-associated changes to skin in humans and mice and identify exercise-induced IL-15 as a novel regulator of mitochondrial function in aging skin. We show that exercise controls IL-15 expression in part through skeletal muscle AMP-activated protein kinase (AMPK), a central regulator of metabolism, and that the elimination of muscle AMPK causes a deterioration of skin structure. Finally, we establish that daily IL-15 therapy mimics some of the anti-aging effects of exercise on muscle and skin in mice. Thus, we elucidate a mechanism by which exercise confers health benefits to skin and suggest that low-dose IL-15 therapy may prove to be a beneficial strategy to attenuate skin aging.

Key words: aging; exercise; muscle; metabolism; mitochondria; skin.

Introduction

Skin is the largest organ in the human body and the primary physical barrier against infection and disease. Aging is associated with the deterioration of the dermal and epidermal layers of the skin, resulting from reductions in cell proliferation, collagen synthesis, extracellular matrix remodeling, and altered epidermal morphology (Fisher *et al.*, 2002). These pathophysiological changes are thought to be driven by

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aged and senescent cells that exhibit reduced energy metabolism, higher mitochondrial oxidative stress, and pronounced mitochondrial DNA (mtDNA) deletions (Isobe *et al.*, 1998; Lu *et al.*, 1999), reflecting characteristics of the mitochondrial free radical theory of aging (Harman, 1972). In agreement, the deletion of a free radical scavenger within the mitochondria, superoxide dismutase 2, from connective tissue results in premature skin aging (Treiber *et al.*, 2011; Velarde *et al.*, 2012) and depleting mtDNA in dermal fibroblasts mimics the gene profile of photoaging (Schroeder *et al.*, 2008). Conversely, treatment with PPAR agonists that stimulate mitochondrial metabolism and cell proliferation improves skin wound healing (Ham *et al.*, 2010) and retards age-related tissue degeneration (Dillon *et al.*, 2012). Therefore, interventions that improve skin metabolism and mitochondrial function are a promising means to maintain skin health in old age.

Endurance exercise is commonly prescribed to improve the health and function of the whole body via its ability to improve energy metabolism. Individuals that adhere to a regular exercise program have lower comorbidities and disease risk and exhibit greater survival in later life (Chakravarty et al., 2008), but the underlying mediators of these benefits are incompletely understood. Recently, we have shown that exercise training can prevent the systemic mitochondrial dysfunction and progeroid symptoms in the polymerase-gamma mutator mouse, including a prevention of skin deterioration (Safdar et al., 2011). Given the systemic nature of the benefits of physical activity, it seems reasonable that circulating factors induced by exercise are likely to mediate at least a portion of these adaptations. Indeed, several proteins secreted from skeletal muscle (myokines) have been identified that can mediate important physiological functions in other organs (Pedersen & Febbraio, 2012), demonstrating a tremendous potential for peripheral tissue metabolic control by exercise-mediated factors.

While facets of skin aging have been ascribed to extracellular hormones that are altered during the lifespan, such as the growth hormone/IGF-1 axis (Zouboulis & Makrantonaki, 2012) or cytokines (Coppé *et al.*, 2008), the endocrine effects of exercise on skin or its aging process have not been studied. In the current study, we provide the first knowledge that regular exercise can attenuate skin aging in humans and mice. We show that exercise regulates skin mitochondrial metabolism and that mice lacking muscle AMPK have reduced serum IL-15 and accelerated skin aging. Finally, we partially rescue age-associated skin deterioration to a similar extent as exercise by injecting aged mice with recombinant IL-15, demonstrating the critical role of this factor in skin function and metabolism.

Results

Exercise robustly alters whole-body metabolism and protects from ageassociated physical deterioration and disease, although the most wellestablished positive adaptations involve metabolic tissues such as liver, skeletal muscle, or the cardiovascular system (Lanza *et al.*, 2008; Seals *et al.*, 2009). To determine whether parallel adaptations occur in skin, we assessed changes in skin structure and physiology in habitually active subjects (ACT, \geq 4 h week⁻¹ of high-intensity aerobic exercise) compared to sedentary controls (SED, \leq 1 h week⁻¹ of exercise) across the human lifespan. Importantly, these subjects were recruited based on

© 2015 The Authors. Aging Cell published by the Anatomical Society and John Wiley & Sons Ltd. This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited. their regular duration of physical activity and not their performance to make our conclusions more relevant to the general population. We found ACT subjects had a thinner stratum corneum epidermal layer throughout the lifespan (Fig. 1A,B; for subject characteristics see Table S1). Additionally, the oldest (65–86 years) ACT subjects also had attenuated thinning of the stratum spinosum layer of the epidermis compared to the SED subjects (Fig. 1C). However, habitual exercise did not appear to influence the age-related loss of reticular dermis collagen (Fig. 1D,E). As others have observed similar improvements to the skin of aged mice using interventions that enhance mitochondrial biogenesis

(Dillon *et al.*, 2012), we assessed mtDNA content and gene expression in ACT and SED subjects in skin as well as commonly available buccal cell (cheek swab) samples. We found that mtDNA copy number in buccal cells and skin was greater in ACT vs. SED subjects (Fig. 1F,G). Similarly, basal levels of buccal cell mitochondrial genes were only maintained with age in ACT subjects (Fig. 1H–J). These findings indicate that exercise can attenuate some aspects of skin aging in humans and that these changes are associated with improvements in tissue mitochondria.

As our cross-sectional comparisons could be influenced by unknown confounding variables across the study groups, we chose to enroll a



Fig. 1 Chronic exercise adherence is associated with attenuated aging in human skin and greater mitochondrial content. (A) Stratum corneum thickness and (B) representative images of skin cross sections stained using H&E from 20- to 86-year old subjects that were highly active exercisers (ACT) or were largely sedentary (SED). Black arrow indicates the stratum corneum layer, double asterisk indicates the stratum spinosum, and single asterisk indicates the dermis. (C) Stratum spinosum layer thickness in SED and ACT subjects in various age groups. (D) Reticular dermis collagen content in each age group and (E) representative trichrome-stained cross sections of human skin. Data from SED and ACT groups have been combined as there was not a group effect. (F) mtDNA copy number in skin and (G) buccal cell samples from habitually sedentary (SED) and highly active (ACT) individuals. (H) mRNA expression of *Cox7b*, (I) *Nd1*, and (J) *Pgc*-1α in basal buccal cell samples from 20- to 86-year old SED and ACT subjects. Data are mean \pm SE. For all analyses, n = 11-19 per age and activity group. Cross-sectional analyses and comparisons were made using a two-way ANOVA. *indicates a significant (P < 0.05) difference from the indicated group(s). #indicates a significant (P < 0.05) overall activity group effect. [†]indicates a difference (P < 0.05) from 20-39 year old individuals.

subset of the sedentary elderly adults into a 3-month cycling exercise program to more directly determine the ability of exercise to reverse agerelated changes to human skin. We found that 12 weeks of endurance exercise training in sedentary elderly adults reduced stratum corneum thickness (Fig. 2A) and, unexpectedly, decreased stratum spinosum thickness (Fig. 2B). This contrasting effect of exercise in the stratum spinosum compared to our cross-sectional cohort may be due to our use of a relatively short exercise intervention, whereas a longer intervention may promote more extensive remodeling that could restore stratum spinosum thickness. Additionally, endurance exercise training in elderly SED subjects increased collagen content (Fig. 2C) and skin mtDNA copy number (Fig. 2D).

To determine whether ACT individuals produced a unique mitochondrial response to exercise, we measured the expression of the master regulator of mitochondrial biogenesis, Pgc-1a, in buccal cells from each group immediately after exercise and found that only ACT subjects increased Pgc-1a expression postexercise (Fig. 3A). We hypothesized that a circulating factor was mediating our observed changes in mitochondria: therefore, we incubated human primary dermal fibroblasts in media conditioned with 10% human serum acquired from young SED and ACT individuals prior to and following acute exercise. Similar to our findings regarding buccal swab Pgc-1a mRNA, we found that only ACT postexercise serum-conditioned media increased mitochondrial content in fibroblasts (Fig. 3B), an effect that was completely ablated when serum proteins were precipitated from solution (Fig. S1A, B). We then evaluated plasma samples from these subjects before and after exercise using a panel of known cytokines and chemokines to screen for proteins that were uniquely altered in the ACT postexercise condition (Fig. 3C, complete results in Table S2). This analysis produced four candidate proteins: TNF-β, IL-15, IL-10, and RANTES; however, as RANTES has only been associated with inflammation, we focused on the remaining analytes (Figs 3D and S1C). We then tested which of these analytes were essential to the mitochondrial response by pretreating the serum with neutralizing antibodies to TNF- β , IL-10, or IL-15 and found that only anti-IL-15 antibodies mitigated the exercise-stimulated increase in mitochondrial complex IV (Fig. 3E) and citrate synthase activity (Fig. S1D) in the fibroblasts. We next determined whether IL-15 stimulates mitochondrial function in a dose-dependent and cell-autonomous manner by measuring mitochondrial respiration in human dermal fibroblasts treated with recombinant human IL-15 (rhIL-15) between 0 and 1000 pg mL⁻¹. Mitochondrial respiration increased with rhIL-15 doses up to 10 pg mL^{-1} , then declined (Fig. 3F), suggesting a narrow window of therapeutic effectiveness that corresponds closely to our observed postexercise levels of IL-15 in human plasma (5.6 \pm 1.2 pg mL $^{-1}$). Higher rhIL-15 levels of 100 and 1000 pg mL $^{-1}$ of rhIL-15 caused an increase in dermal fibroblast proliferation vs. control (Fig. 3G), similar to previously described growth effects in liver (Suzuki *et al.*, 2006). Additionally, when we co-incubated primary human fibroblasts with the rhIL-15 and PPAR γ antagonist GW9662 or the STAT5 inhibitor pimozide, the increase in cytochrome c oxidase (COX) activity due to rhIL-15 was ablated (Fig. 3H). These data are consistent with previous work showing a mitigation of progeroid aging due to mtDNA mutations in skin using the PPAR γ agonist bezafibrate (Dillon *et al.*, 2012) as well as a requirement for both STAT5 and PPAR γ in keratinocyte differentiation (Dai *et al.*, 2007).

Exercise improves skeletal muscle mitochondrial capacity via signaling cascades that promote mitochondrial biogenesis (O'Neill et al., 2013). We sought to determine whether a regulator of mitochondrial biogenesis, $Pqc-1\alpha$, was elevated in skin following exercise. We found that a single session of treadmill running in mice caused a transient elevation in Pgc-1a mRNA in both skeletal muscle and skin, peaking at 1 h postexercise (Fig. 4A) in parallel to the rise in serum IL-15 (Fig. 4B). To determine whether IL-15 was necessary for exercise-stimulated mitochondrial biogenesis in vivo, we exercised wild-type mice that were given a venous injection of IL-15-neutralizing antibody or IgG control prior to and following 1 h of treadmill exercise. We found that neutralizing circulating IL-15 partially reduced the exercise-induced elevation in skeletal muscle Pgc-1 α expression, but not skin Pgc-1 α expression (Fig. 4C). Skin Pgc-1B expression was also not affected by the IL-15neutralizing antibody (Fig. 4D). However, IL-15 neutralization prevented an exercise-stimulated reduction in skin PPARy coactivator-related protein 1 (Pprc1) mRNA (Fig. S4E) and importantly abolished an increase in downstream mitochondrial gene expression of cytochrome b (Fig. 4F). These results suggested that IL-15 partially mediates exercise-stimulated mitochondrial biogenesis in skin and skeletal muscle.

Next, we sought to determine whether IL-15 was required for basal mitochondrial function, so we evaluated tissue mitochondria in mice lacking whole-body expression of IL-15 (IL-15 KO) and found lower COX activity in skin and skeletal muscle tissue from IL-15 KO mice (Fig. 5A). IL-15 is known to be highly expressed in skeletal muscle (Pedersen & Febbraio, 2012) and also secondarily in skin when compared across metabolic tissues (Fig. 5B). As acute exercise capacity was predictive of the positive effects on skin tissue in our human subjects (20-39 year VO₂ peak, SED: $34 \pm 2 \text{ mL kg}^{-1} \text{ min}^{-1}$, ACT: $57 \pm 2 \text{ mL kg}^{-1} \text{ min}^{-1}$), this implicated the involvement of muscle AMP-activated protein kinase (AMPK), a well-described energy-sensing molecule that regulates exercise capacity (O'Neill et al., 2011). We tested whether AMPK was involved in the expression and/or induction of IL-15 via exercise using mice that lacked both the β 1- and β 2-subunits of AMPK in skeletal muscle (AMPK DMKO). These mice have exercise intolerance due to reduced muscle mitochondria and impaired contraction-stimulated glucose uptake but appear phenotypically normal at rest compared to

Fig. 2 A short-term aerobic exercise intervention improves skin structure and increases tissue mitochondria. (A) Stratum corneum thickness, (B) stratum spinosum thickness (C) reticular dermis collagen content, and (D) mtDNA copy number in skin samples acquired before (Pre) and after 3 months of aerobic exercise training (Post) in previously sedentary elderly (65–86 year) adults. Data are mean \pm SE. n = 10. Pre/ post-training comparisons were made using a paired *t*-test. *indicates a significant (P < 0.05) difference from the indicated group(s).





Fig. 3 IL-15 is responsible for exercise-stimulated mitochondrial biogenesis in skin *in vitro*. (A) Expression of *Pgc-1* α mRNA in buccal swabs prior to and immediately following a single session of aerobic exercise using *Gapdh* as a stable housekeeping gene. *n* = 8 per group. (B) Representative immunoblots of mitochondrial protein subunits of complex III (core 2, CIII) and complex IV (COX II, CIV) in primary dermal fibroblasts that were incubated for 48 h with media containing 10% human serum from the indicated blood sample. *n* = 4 replicates per condition. (C) Criteria used to analyze the results of a plasma cytokine panel for determination of factors induced by exercise that are likely to signal peripheral tissue mitochondrial biogenesis. Numbers in parentheses indicate the number of analytes that pertain to the successive criteria. (D) The percent change in plasma IL-15, IL-10, and TNF- β in SED and ACT groups in response to a single bout of exercise. *n* = 8 per group. (E) Immunoblots of mitochondrial complex IV protein in primary dermal fibroblasts incubated for 48 h in enriched media containing 10% basal or postexercise (Post-ex) serum from ACT subjects that was pretreated with control IgG, anti-TNF- β , anti-IL-15, or IL-10-neutralizing antibidies. *n* = 4 replicates per condition. (F) Oxygen consumption rate (OCR) and (G) cell counts of nucubated with 10 pg mL⁻¹ / hlL-15 with and without 1 μ M GW9662 or 5 μ M pimozide for 48 h. Data are mean \pm SE. Data in A and D were compared using a two-way repeated-measures ANOVA. Data in E were compared using a paired *t*-test. Data in F–H were compared using a one-way ANOVA. **P* < 0.05 from basal or control condition. ND, not detectable.

wild-type littermates (O'Neill *et al.*, 2011). We found that muscle *ll15* mRNA expression and plasma IL-15 were reduced in AMPK DMKO mice (Fig. 5C,D) and, similar to the IL-15 KO mice, AMPK DMKO mice had reduced COX activity in skin tissue (Fig. 5E). To further explore whether muscle AMPK was necessary for the stimulation of *ll15* expression, we incubated skeletal muscle from WT and AMPK DMKO mice *ex vivo* in the presence or absence of the AMPK activator AICAR. AICAR produced an increase in *ll15* in both soleus and EDL muscles in wild-type, but not in AMPK DMKO mice (Fig. 5F). To mimic exercise, we examined *ll15*

expression in *tibialis anterior* (TA) muscles following a 30-min contraction protocol, matching the work performed by each group (average force of contraction – WT: 236 ± 11 mN, AMPK DMKO: 232 ± 10 mN). We found no changes in *ll15* mRNA were present at the 0 h time point, but that *ll15* expression was specifically increased in wild-type but not in AMPK DMKO mice 3 h after the muscle contractions compared to the resting limb (Fig. 5G), indicating that contraction-stimulated *ll15* expression is dependent on AMPK activity. Furthermore, we analyzed skin morphology in older (18 month old) WT and AMPK DMKO mice and



Fig. 4 IL-15 partially regulates exercise-mediated mitochondrial signaling in skin and skeletal muscle. (A) qPCR of $Pgc-1\alpha$ mRNA in tissues from wild-type mice that were sacrificed at the indicated times after cessation of exercise or that did not exercise (SED). Muscle is *quadriceps* tissue, n = 9 per group. (B) Corresponding serum IL-15 concentration at the indicated times postexercise. n = 9 per group. (C) $Pgc-1\alpha$ mRNA expression in *quadriceps* muscle and skin from mice that remained sedentary or exercised in conjunction with injection of IgG control or anti-IL-15-neutralizing antibody. n = 6-10 per group. (D) $Pgc-1\beta$ and (E) Pprc1 mRNA expression in skin tissue from mice that remained sedentary or exercised in conjunction with injection if IgG control or anti-IL-15-neutralizing antibody. n = 6-10 per group. (F) Cytb mRNA mRNA is in acquired from wild-type mice that rested (SED) or were subjected to treadmill exercise in conjunction with tail vein injection of a total of 5 μ g of IgG control or anti-IL-15 antibody. n = 6-10 per group. Mice in the neutralization experiments were injected with 2.5 μ g antibody immediately before and after the exercise bout and sacrificed at 1 h following cessation of exercise. All acute mouse exercise experiments were performed at a 10-degree uphill grade at a speed of 16 m per minute for 1 h. mRNA expression analyses used β 2-microglobulin as a stable housekeeping gene. Data are mean \pm SE. Data were compared using a one-way ANOVA. *P < 0.05 from SED group. †P < 0.05 from EX + IgG. #P < 0.05 from all other conditions.

found reduced collagen content (Fig. 5H,I) and reduced dermal thickness in AMPK DMKO mice (Fig. 5H,J), consistent with their skin mitochondrial defects. Overall, these results suggest that IL-15 is regulated by muscle AMPK and that skeletal muscle metabolism can regulate skin morphology.

As IL-15 appears to regulate exercise-stimulated mitochondrial signaling in skin and skeletal muscle, we sought to clarify its therapeutic potential in young (5 month) and old (23 month) mice with daily intravenous injections of recombinant mouse IL-15 (rmIL-15) that mimicked the physiologic elevation of endogenous IL-15 observed following acute exercise (Fig. S2A). Both young and old mice that received rmIL-15 injections or exercise had significantly higher skin and muscle COX activity, and mtDNA copy number (Fig. 6A-D). However, in contrast to previous findings from muscle IL-15-overexpressing mice (Quinn et al., 2009), there was no change in body or tissue mass in any of the groups (Fig. S2B-F) and only the exercise group had an increase in treadmill running capacity (Fig. S2G). Moreover, oxygen consumption, cage movement, and food intake were similar among 5-month-old mice (Fig. S3A-C), indicating that behavioral changes did not account for the changes in mitochondria. However, both rmIL-15 and exercise treatment resulted in higher cage activity and oxygen use in 23-month-old mice (Fig. 6E,F), indicating a partial restoration of physical function specifically in the aged mice, although there was no difference in food consumption (Fig. S3C). Despite no change in muscle mass, rmlL-15 and exercise treatment also improved muscle grip strength in 5-month-old mice (Fig. 6G). Finally, rmIL-15 and exercise treatment resulted in higher stratum spinosum thickness and dermal collagen content in 23-monthold mice compared with PBS-treated mice, which partially reversed the effects of aging (Fig. 6H,I). These findings are in agreement with reports showing stimulation of collagen production *in vitro* by rhlL-15 (Kim *et al.*, 2014) and PPAR agonists (Ham *et al.*, 2010) and demonstrate that physiologic elevations of circulating IL-15 can mimic some of beneficial effects of exercise training on aging skin and skeletal muscle metabolism. Furthermore, as inflammation is a hallmark of aging and metabolic dysfunction (Green *et al.*, 2011), we measured plasma cytokines from each treatment group and found that rmlL-15 and exercise treatment reduced circulating levels of the inflammatory cytokines IL-6 and MCP-1 (Fig 6J,K).

Discussion

The broad, systemic benefits of exercise on the aging process (Chakravarty *et al.*, 2008) are likely complexly regulated among many organs. We identify circulating IL-15 as an exercise-stimulated hormone that mediates the health of skin tissue. While IL-15 is one of many factors secreted from skeletal muscle that is elevated following exercise in mice and humans (Pedersen & Febbraio, 2012), the regulation of musclederived IL-15 and its physiological consequences has remained unclear. IL-15 has been shown to be elevated in humans after exercise in some studies (Tamura *et al.*, 2011), but not all (Nielsen *et al.*, 2007). The relatively transient increase in IL-15 in circulation immediately following exercise seen in the current work as well as the work of Tamura *et al.* (2011), suggests that the timing of blood sampling is important in detecting the exercise response. Moreover, an immediate change in



Fig. 5 IL-15 expression and skin metabolism are regulated by skeletal muscle AMPK. (A) COX activity relative to WT mice in skeletal muscle (*quadriceps*), and skin tissue from WT and IL-15 KO mice. n = 9. (B) Relative //15 mRNA expression in body tissues. n = 8-10. (C) //15 mRNA in *gastrocnemius* muscle (n = 4) and (D) plasma IL-15 levels in 3-month-old wild-type littermates (WT) and AMPK $\beta1\beta2$ double-muscle knockout (AMPK DMKO) mice (n = 5). (E) Cytochrome c oxidase activity in skin from WT and AMPK DMKO mice. (F) //15 mRNA expression in isolated *EDL* and *soleus* muscles from WT and AMPK DMKO mice that were incubated with 2 mM AICAR or vehicle for 2 h. n = 3. (G) //15 mRNA expression in *tibalis anterior* muscle from WT and AMPK DMKO mice that were subjected to a 30-min *in situ* electrical stimulation protocol (STIM) or served as the contralateral control (CON). n = 4-5. (H) Histological images of skin from 18-month-old wild-type and AMPK DMKO littermates stained with H&E (top) or trichromes (bottom). Black arrowhead is the stratum spinosum and the white asterisk indicates the dermis. (I) Quantification of skin dermal collagen and (J) stratum spinosum thickness in aged WT and AMPK DMKO mice. n = 5. mRNA expression was normalized to *Gapdh* or β -*actin* as a stable housekeeping gene. Results in A, C–E, and I–J were compared using an unpaired *t*-test. Data in F were analyzed using a three-way ANOVA. G was compared using two-way repeated-measures ANOVA. *Significantly different (P < 0.05) from the indicated group or from the control condition. ⁴Main effect of genotype. [#]Main effect of muscle type. Data are mean \pm SE.

circulating levels of IL-15 suggests a pool of readily secretable IL-15 that would not necessarily be reliant on transcriptional changes in IL-15 mRNA, as noted by others (Bamford *et al.*, 1998).

Muscle IL-15 has been previously associated with whole-body metabolic changes. Work using muscle IL-15-overexpressing mice has suggested that IL-15 negatively regulates adiposity (Quinn *et al.*, 2009), enhances muscle mass, and improves exercise capacity; however, these mice are hyperactive at a young age (Quinn *et al.*, 2009), which may account for much of their improved energy metabolism. Additionally, as circulating IL-15 is ~5000-fold higher in these mice compared to our basal levels, their effects ascribed to IL-15 may not be physiologically relevant. In particular, we see an increase in muscle strength in young mice, but no changes in muscle mass with rmIL-15 treatment, possibly

due to our use of a much lower dose than the mouse muscle overexpression model. However, it is possible that a more prolonged treatment period with low-dose rmIL-15 would have produced more robust changes in muscle size in light of the fact that a splice variant of $Pgc-1\alpha$ has been shown to enhance muscle mass (Ruas *et al.*, 2012).

IL-15 is necessary for natural killer immune cell activation and expansion in response to viral pathogens, primarily through the IL-15 receptor $\beta\gamma$ complex (Waldmann, 2006). IL-15 binds to its β - and γ -receptors (Kd: 10^{-9} M) (Giri *et al.*, 1994) but has higher affinity for its α -receptor (Kd: 10^{-11} M) (Anderson *et al.*, 1995), indicating that expression well below 10^{-9} M could avoid overt pathogen-like immune system activation. Exercise and our injected dose of IL-15 in wild-type mice were suitable to transiently reach a peak of ~100 pg mL⁻¹

Fig. 6 Daily IL-15 therapy in mice mimics the anti-aging effects of exercise on skin structure and mitochondria. Chow-fed mice were injected once daily via tail vein with PBS, recombinant mouse IL-15 (rmIL-15), or underwent forced exercise training (EX) for 33 consecutive days and were sacrificed at the age indicated. (A) *Quadriceps* muscle mtDNA copy number and (B) cytochrome *c* oxidase (COX) activity. n = 6–8 per group. (C) Skin mtDNA copies and (D) COX activity. n = 6–8 per group. (E) Cage activity and (F) oxygen uptake (VO₂) over a 24-h period in 23-month-old mice treated with PBS, rmIL-15, or EX. (G) Peak grip strength in each treatment group normalized to body weight. (H) Quantification of dermal collagen content (left) and representative images of trichrome-stained skin cross sections from each treatment group (right). n = 4–8 per group. White asterisk indicates the dermis. (I) Quantification of startum spinosum thickness and representative images of H&E-stained cross sections of skin from each group. (L) Graphical illustration of the effects of exercise-induced muscle IL-15 signaling to skin tissue. Five-month-old and 23-month-old mice were injected with 500 and 1000 pg of rmIL-15, respectively. All data were compared using a two-way ANOVA. *indicates a significant difference (P < 0.05) from PBS or the indicated group. [†]indicates a noverall effect of age. Data are mean \pm SE.



(~0.76 × 10⁻¹¹ M) in circulation, possibly low enough to avoid the negative consequences of chronically elevated IL-15 that may trigger leukemia (Fehniger *et al.*, 2001) or pancreatic β-cell death (Chen *et al.*, 2013) in mice. IL-15 has been administered to mice (1200–7200 µg kg⁻¹) (Munger *et al.*, 1995) and rhesus macaques (10–50 µg kg⁻¹) (Waldmann *et al.*, 2011) at far higher doses than our injected dose (~17 ng kg⁻¹). However, the ability of doses of IL-15 higher than those experienced during exercise to mediate changes in skin mitochondrial function remains questionable, as levels higher than 10 pg mL⁻¹ did not stimulate mitochondrial respiration in dermal fibroblasts.

We hypothesize that fewer 'pulses' of IL-15 from skeletal muscle and other tissues during sedentary living are in part responsible for the accelerated degeneration of dermal fibroblasts that drive the clinical symptoms of skin aging. These findings are bolstered by prior reports that mitochondrial energetics play a central role in aging (Trifunovic *et al.*, 2004) and that mitochondrial biogenesis can retard cellular aging (Sahin & Depinho, 2010; Safdar *et al.*, 2011). Moreover, the finding that muscle AMPK deletion causes skin deterioration is in line with previous work, showing that the circulating environment can strongly affect brain and muscle aging (Villeda *et al.*, 2011; Katsimpardi *et al.*, 2014; Sinha *et al.*, 2014). As low muscle AMPK activity is commonly observed in a wide range of health disorders (Steinberg & Jørgensen, 2007), reduced basal or exercise-stimulated IL-15 secretion and other AMPK-dependent factors may exacerbate skin or muscle tissue disease progression.

As mitochondrial degeneration is widely implicated in disease, there are many potential conditions where the exercise mimicking effects of IL-15 therapy might be useful, such as in frail patients or those with metabolic disease that have a low exercise capacity. Additionally, IL-15 may prove beneficial for cutaneous wound healing as stimulating mitochondrial metabolism improves skin responses to injury (Ham *et al.*, 2010). Given the prior findings that IL-15 may stimulate browning of white adipose tissue (Boström *et al.*, 2012) and promote liver repair (Suzuki *et al.*, 2006), exercise-induced IL-15 may also be responsible for improvements in other organs and prove a multifaceted strategy to treat health disorders.

Experimental procedures

Human subject recruitment and testing

A subset of the human subjects in this study have been previously described (Crane *et al.*, 2013b), and all experimental procedures were approved by the institutional research ethics board. Tissue samples from human subjects were collected following an overnight fast. Skin samples were acquired from the upper portion of the non-sun-exposed buttocks below the waistline using a 4-mm punch biopsy under local anesthetic without norepinephrine. Subcutaneous fat was separated from the bottom portion of the dermis following tissue collection. Buccal swabs and plasma and serum samples were collected at rest and immediately following acute exercise. All samples were flash-frozen in liquid nitrogen. After baseline tissue collections, subjects underwent a bout of exercise testing to confirm aerobic fitness, as described (Crane *et al.*, 2013b). Within 5 min of finishing, each subject then underwent 30 min of exercise at 50% of their cycling power maximum. In total, the subjects cycled for approximately 45 min each.

In addition to the acute exercise trial, a portion of the 65-86 year old sedentary subjects underwent 12 weeks of twice-weekly aerobic exercise training on a Monark Cardio Care 827E cycle ergometer. The training program started with participants maintaining 65% of HRmax for 30 minutes and progressed by 5% every other week until intensity reached 75% of HR max, at which point the duration of exercise increased by 5 minutes every other week reaching 75% of HR max for 45 minutes at the end of the training intervention.

Sample analysis

Plasma (EDTA) cytokines from human subjects were initially analyzed using a 42-analyte multiplex ELISA assay (Millipore, Millipore, Billerica, MA, USA), and subsequent mouse IL-15 analyses were performed as a single-plex ELISA. GM-CSF, IFN-γ, IL-10, IL-12 (p70), IL-13, IL-1β, IL-2, IL-4, IL-5, IL-6, IL-7, IL-8, and TNF- α were assessed using high-sensitivity ELISAs (0.13–2000 $pg mL^{-1}$), and all other analytes were analyzed using standard detection limits (3.2–10 000 pg mL⁻¹). Proteins in human serum were precipitated using ethanol, followed by centrifugal pelleting of the precipitant and removal of the supernatant that contained soluble metabolites. The supernatant fraction was then dried using a vacuum centrifuge, and the metabolites were resuspended in sterile water to the original serum volume. The removal of all proteins was confirmed using a protein assay, and this fraction was added to intact serum for cell culture experiments as indicated. Primary human dermal fibroblasts were cultured according to standard explant methods (Villegas & McPhaul, 2001) and all cell treatments were for 48 h. The recombinant human IL-15 was purchased from R&D Systems, and the GW9662 compound and pimozide were acquired from Sigma-Aldrich Sigma-Aldrich, St. Louis, MO, USA Abcam, Cambridge, UK R&D Systems, Minneapolis, MN, USA.

Cell and tissue lysates were prepared using 0.05 M potassium phosphate buffer, and mitochondrial protein immunoblotting was performed as described using human or rodent antibodies from Abcam (#ab110411, #ab110413). Cytochrome c oxidase activity was performed by incubating sample lysates with reduced cytochrome c and measuring the change in absorbance at 550 nm over 90 s in a 96-well plate. DNA and RNA isolation and qPCR of tissues were performed as described (Crane *et al.*, 2013a).

All mouse experiments were performed using female mice on a chow diet. IL-15 knockout mice (IL-15 KO) and C57/BL6 control mice were obtained from Taconic and sacrificed at 12 weeks of age. Wild-type mice for acute exercise experiments were obtained from Jackson laboratories and sacrificed at 16 weeks of age. AMPK DMKO mice, and wild-type littermates were bred and housed in the McMaster University Animal Facility under standard housing conditions with a 12-h light/dark cycle. Acute exercise experiments were performed for 1 h at a speed of 16 m per minute at an uphill grade of 10 degrees. All neutralizing antibodies were purchased from R&D Systems.

Ex vivo and in situ muscle experiments

Soleus and EDL muscles were excised under anesthesia from AMPK DMKO mice and wild-type littermates, incubated in 1 mL of media at 30 °C for 2 h with and without AICAR before being removed, and snap-frozen. *In situ* contraction experiments were performed by first isolating the tibialis anterior (TA) muscle under anesthesia (ketamine/xylazine) from both legs and connecting the distal tendon of one leg to a force transducer with string. The sciatic nerve of that leg was then exposed and enervated using pulsed electrical stimulation 400 time at 1.5–4 milliamps at a frequency of 10 Hz for 0.5 s over the course of 30 min and the contralateral leg served as a nonstimulated control. Mice were maintained on a 33 °C peltier warmed block under isofluroane gas anesthetic during the contraction and recovery periods using a carrier

gas of 95% oxygen. TA muscles were harvested immediately following (0 h) the contraction protocol or after 3 h.

Animal testing

Grip strength was tested in mice as described (Ogborn *et al.*, 2012). Mouse behavior (cage activity, food intake) and metabolic parameters (VO₂, VCO₂, RER) were analyzed using Columbus Lab Animal Monitoring System (CLAMS; Columbus instruments, Columbus, OH, USA) over a 24-h period with lights on at 07:00 and off at 19:00 hours.

C57/BL6 mice that were 4 and 22 months old were randomly allocated to vehicle (PBS), rmIL-15, or exercise daily treatment for 33 consecutive days. rmIL-15 (5-month-old group: 500 pg; 23-month-old group: 1000 pg) or vehicle control (PBS) was injected via the tail vein using a tuberculin syringe. This dosing was used based on pilot work in 3- to 4-month-old mice (Fig. S3A) and because the 23-month-old mice weighed approximately twice as much as the 5-month-old mice at the start of the experiment.

Exercise-trained mice underwent 33 consecutive days of progressive forced treadmill running. The first week of exercise training occurred at a treadmill speed of 16 m per minute for 5-month-old mice and 10 m per minute for 23-month-old mice at a 10-degree uphill grade for 1 h. After one week, the 23-month-old group was increased to 12 m per minute and after 2 weeks increased to 14 m per minute. After the first week, 5-month-old mice were increased to 18 m per minute for the remainder of the training period. Grade and duration remained unchanged for the training period.

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Author contributions

J.D.C. and M.A.T. designed the study and J.D.C. performed most of the experiments. L.G.M., I.K.B., and M.A.T. assisted with the human studies and analyses. J.S.L., R.J.F., and A.L.B. assisted with the AMPK experiments. G.R.S. and B.E.K. provided materials and animals for the experiments and assisted with study design and discussion. J.D.C., G.R.S., and M.A.T. analyzed and interpreted data. J.D.C. wrote the manuscript.

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Conflict of interests

The authors have no conflict of interests to declare.

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Supporting Information

Additional Supporting Information may be found in the online version of this article at the publisher's web-site.

Fig. S1. Exercise serum deproteinization and plasma analyte screen results.

Fig. S2. Recombinant mouse IL-15 dosing and body and tissue weights from daily PBS, rmIL-15 and EX treatments.

Fig. S3. Metabolic and behavioral characteristics of 5 month old treated mice.

 Table S1.
 Subject physical characteristics.

Table S2. Results of the exercise-stimulated plasma cytokine and chemokine screen in young individuals.

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