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Characteristics of Muscle Fiber Type Are Predictive of Skeletal Muscle Mass and Strength in Elderly Men

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OBJECTIVES: To investigate the relationship between skeletal muscle fiber type-specific characteristics, circulating hormone concentrations, and skeletal muscle mass and strength in older men.

DESIGN: Cross-sectional analyses.

SETTING: University research center.

PARTICIPANTS: Forty-one community dwelling elderly men (≥ 65).

MEASUREMENTS: Leg strength (1-repetition maximum, 1RM) and whole-body and limb muscle mass were determined, and muscle fiber type composition, cross-sectional area (CSA), myonuclear content, and satellite cell (SC) content were assessed in skeletal muscle biopsy samples. In addition, blood samples were collected to determine serum testosterone, sex hormone-binding globulin, insulinlike growth factor (IGF)-1, and IGF binding protein-3 concentrations.

RESULTS: Muscle mass correlated with muscle strength ($0.41 \leq \text{correlation coefficient } (r) \leq 0.72$; $P < .01$). Muscle fiber CSA, myonuclear content, and SC content were significantly lower in type II than in type I muscle fibers. Myonuclear and SC content were positively correlated with muscle fiber CSA. Furthermore, greater muscle fiber CSA (type I and II) was associated with greater thigh muscle area and muscle strength ($0.30 \leq r \leq 0.45$; $P < .05$). Testosterone concentration was positively correlated with muscle mass and muscle fiber CSA. Regression analysis showed that SC content, myonuclear content, and testosterone concentration are predictive of muscle fiber CSA. Furthermore, muscle mass and type II muscle fiber CSA are predictive of muscle strength.

CONCLUSION: Skeletal muscle mass and strength in elderly men are positively correlated with muscle fiber type-specific CSA, myonuclear content, and SC content.

These findings support the assumption that a decline in SC content plays an important role in age-related decline in muscle mass and strength. *J Am Geriatr Soc* 58:2069–2075, 2010.

Key words: sarcopenia; muscle fiber size; satellite cells; aging; hypertrophy

Aging is associated with the gradual loss of skeletal muscle mass and function, termed sarcopenia. The prevalence of sarcopenia is approximately 25% after the age of 60, increasing to up to approximately 50% in people aged 80 and older.^{1–3} The latter is in accordance with the progressive nature of the loss of muscle mass^{4–8} and muscle strength^{6,9} with aging. The etiology of sarcopenia is generally studied in a cross-sectional manner within a large population of individuals ranging in age from 20 to 80, although few data are available on the determinants of muscle mass and strength within an elderly population. Some studies have reported that muscle mass and strength continue to decline after the age of 65.^{10,11} This seems consistent with the greater prevalence of sarcopenia in people aged 80 and older than in those aged 60 and older^{1–3} and agrees with longitudinal changes reported previously.^{12,13} Nevertheless, there is a large variability in muscle mass and strength within the elderly population that is not necessarily related to age per se.¹⁴ Therefore, studying determinants of muscle mass within the elderly population (independent of age) could provide better insight into the key etiological factors of sarcopenia.

Leg skeletal muscle mass seems to be the main determinant of leg muscle strength, independent of age.^{15,16} The loss of leg muscle mass in older adults has been attributed to a decline in the number of muscle fibers and, more specifically, to type II muscle fiber atrophy.^{8,17,18} These findings were recently extended by showing a type II muscle fiber-specific decline in satellite cell (SC) content in older muscle.¹⁹ It has been suggested that such a decline in muscle

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fiber SC content plays an important role in muscle fiber atrophy^{20,21} and the subsequent loss of muscle mass with aging. It could be speculated that SC content, muscle fiber size, and muscle mass are closely linked, independent of age. Therefore, it was hypothesized that, in an elderly population, muscle fiber size and SC content are correlated and are predictive of whole-body and limb skeletal muscle mass and strength.

There have been ample suggestions that age-related changes in hormonal profile can strongly modulate skeletal muscle mass and strength in older adults.²² In accordance, significant correlations have been reported between lean mass and muscle strength and serum testosterone and insulinlike growth factor (IGF)-1 concentrations in elderly men.^{2,14,23} Furthermore, dose-dependent increases in lean mass, strength, muscle fiber size, and SC content have been reported after testosterone supplementation in younger and older men,^{24,25} which is indicative of a role for anabolic hormones in regulating SC-induced muscle fiber hypertrophy. The goal of the present study was to extend previous findings^{24,25} by investigating the relationship between baseline serum hormone profiles and muscle characteristics. In line with potential hormonal regulation of SC, it was hypothesized that baseline hormone concentrations (particularly testosterone) are predictive not only of skeletal muscle mass and strength, but also of SC content and muscle fiber size.

The present study aimed to investigate the relationship between skeletal muscle fiber type-specific characteristics, circulating hormone concentrations, and skeletal muscle mass and strength in older men. Skeletal muscle characteristics were assessed at the whole-body, limb, and muscle fiber level, and circulating hormone concentrations were determined in a group of community-dwelling older men.

METHODS

Participants

Forty-one older men (65–86) were included in the present study. All participants participated in different studies that are part of a greater project investigating the clinical benefits of exercise and nutritional intervention in older men for which participants were recruited through advertisements in local newspapers. Medical history of all participants was evaluated, and an oral glucose tolerance test (OGTT) and resting electrocardiogram were performed. Participants with (silent) cardiac or peripheral vascular disease, orthopedic limitations, or type 2 diabetes mellitus²⁶ were excluded. All participants were living independently and had no history of participating in any structured exercise training program for at least 5 years. Furthermore, participants reported no problems in normal activities of daily living (e.g., walking, climbing stairs, rising from a chair) and did not need any assistive equipment (e.g., using a cane) while walking. Participants taking antihypertensive medications ($n = 13$), cholesterol- or lipid-lowering medications ($n = 8$), medications for prostate or urinary problems ($n = 8$), or no medications ($n = 20$) were included in the study. Individuals with more-severe medical problems (treatment by medical specialist more than twice a year) were excluded. Participants from the larger cohort ($n = 59$) were included in the present analysis when a muscle biopsy

was available, with at least 75 type I and 75 type II muscle fibers being analyzed, which is required to reliably assess myonuclear and SC content.²⁷ No differences were observed between the participants included in the present study and the larger cohort regarding age, body composition, and muscle mass and strength (Table 1). All participants were informed of the nature and possible risks of the experimental procedures before written informed consent was obtained. The Medical Ethics Committee of the Maastricht University Medical Center approved the study.

Study Design

After inclusion in this study, all participants underwent the same series of measurements (outlined below) to determine muscle characteristics at the whole-body, limb, and myocellular level. In addition, a single fasting blood sample was collected from every participant between 8:00 and 9:00 a.m. to assess hormonal profile. On all test days (muscle biopsy and blood sampling and dual energy X-ray absorptiometry (DXA) and computed tomography (CT) scanning), participants arrived at the laboratory by car or public transportation at 8:00 a.m. and in an overnight fasting state. In addition, participants were asked to maintain their normal physical activity pattern in the 3 days before muscle biopsy collection (no unaccustomed strenuous physical exercise or labor) to prevent any confounding transient perturbations in baseline SC content.

Strength Assessment

Maximum strength was assessed according to 1-repetition maximum (1RM) strength tests on leg press and leg extension machines (Technogym, Rotterdam, the Netherlands). During a familiarization trial, proper lifting technique was demonstrated and practiced. Maximum strength was estimated using the multiple-repetition testing procedure.²⁸ In an additional session, at least 1 week before muscle biopsy collection, each participant's 1RM was determined as described previously;²⁹ 1RM as determined from the second session was always the largest value recorded.

Table 1. Subject Characteristics

Characteristic	Mean \pm Standard Deviation (range)	
	Present Study (n = 41)	Total Cohort (n = 59)
Age	72 \pm 5 (65–86)	72 \pm 5 (65–86)
Body mass, kg	80.3 \pm 11.8 (63.1–102.2)	79.8 \pm 11.1 (63.0–102.2)
Height, m	1.73 \pm 0.06 (1.61–1.86)	1.72 \pm 0.05 (1.61–1.86)
Body mass index, kg/m ²	27.0 \pm 3.5 (21.6–34.6)	26.8 \pm 3.2 (21.6–34.6)
Fasting glucose, mmol/L	5.7 \pm 0.6 (4.5–6.9)	5.7 \pm 0.6 (4.5–6.9)
Body fat, %	25.4 \pm 6.8 (10.6–39.8)	25.3 \pm 6.5 (10.6–39.8)
Lean mass, kg	56.1 \pm 5.3 (49.0–71.1)	55.9 \pm 4.9 (48.8–71.1)
1RM leg extension, kg	84 \pm 13 (50–110)	85 \pm 12 (50–110)
1RM leg press, kg	170 \pm 28 (105–225)	170 \pm 27 (105–225)

No differences were observed between the subjects included in the present study, and the total cohort from which this group was selected.

1RM = 1-repetition maximum.

Thigh Muscle Cross-Sectional Area

Anatomical cross-sectional area (CSA) of the thigh muscles was assessed using CT scanning (IDT 8000, Philips Medical Systems, the Netherlands), as described previously.³⁰ Scans were performed at the mid-thigh level, and images were loaded onto a personal computer using AGFA IMPAX imaging software, version 5.2 (AGFA Healthcare, Brussels, Belgium). Muscle area of the right leg was selected between -29 and $+150$ Hounsfield units,³¹ after which the quadriceps muscle was selected using manual tracing. Total thigh and quadriceps muscle area were calculated using Lucia 4.81 software (Nikon Instruments Europe, Badhoevedorp, the Netherlands). Two investigators blinded to participant coding performed all analyses; intraclass correlation coefficients for inter- and intrainvestigator reliability were 0.997 and 0.998, respectively.

Body Composition

Directly after CT scanning, body composition and bone mineral content were measured using DXA (Lunar Prodigy Advance, GE Healthcare, Madison, WI). The system's software package (enCORE 2005, version 9.15.00) was used to determine whole-body and regional lean and fat mass. DXA scans were performed in a fasting state after participants had voided. The coefficient of variation (CV) for repetitive scans ($n = 4$, 2 weeks apart) were 0.4%, 1.0%, and 1.1% for whole-body lean mass, fat mass, and leg lean mass, respectively. Appendicular skeletal muscle mass (ASM) was calculated as the sum of lean mass of the arms and legs.¹ Body mass was measured to the nearest 0.1 kg using an electronic balance scale, and height was measured to the nearest 0.1 cm using a wall-mounted stadiometer. To control for the potential confounding effect of body size, all body composition measures were adjusted for knee height.³² Knee height was chosen instead of full height, because age-related disorders affecting the spine (e.g., osteoporosis) can confound the latter.³³

Blood Samples

Blood samples were collected in ethylenediaminetetraacetic acid-containing tubes and serum tubes. After centrifugation, aliquots of plasma and serum were frozen in liquid nitrogen and stored at -80°C . Plasma glucose concentrations were analyzed using a COBAS FARA analyzer (Uni Kit III, Roche, Basel, Switzerland). Total serum testosterone and sex hormone-binding globulin (SHBG) concentrations were measured using reagents from Roche Diagnostics (Mannheim, Germany), and assays were run on a Modular Analytics E170 analyzer (Hitachi Data Systems, Santa Clara, CA). The intraassay CVs are 2.7% to 1.8% and 1.1% to 1.7% for low to high concentrations of testosterone and SHBG, respectively. Bioavailable testosterone was calculated as non-SHBG-bound testosterone using a formula described and validated previously.³⁴ IGF-1 was analyzed on a Liaison system (DiaSorin, Brussels, Belgium) with reagents from DiaSorin. Intraassay CVs are 2.4% to 4.4% for high and low concentrations, respectively. IGF binding protein (BP)-3 was analyzed using a commercially available test kit (Biosource Europe, Nivelles, Belgium) on a Tecan/GENios analyzer (Tecan Group Ltd., Männedorf, Switzerland), with intraassay CVs of 4% to 8%.

Muscle Biopsy Sampling

Muscle biopsy samples were collected from the right leg in the morning after an overnight fast. After local anesthesia, percutaneous needle biopsies (50–80 mg) were taken from the vastus lateralis muscle, approximately 15 cm above the patella.³⁵ Any visible nonmuscle tissue was removed immediately, and biopsy samples were embedded in Tissue-Tek (Sakura Finetek, Zoeterwoude, the Netherlands), frozen in liquid nitrogen-cooled isopentane, and stored at -80°C .

Immunohistochemistry

From all biopsies, 5- μm -thick cryosections were cut at -20°C . Samples from two participants were mounted together on uncoated glass slides. Care was taken to align the samples properly for cross-sectional fiber analyses. Serial cross-sections were stained for muscle fiber typing and myocellular SC content. Details of the analytical procedures have been described previously.³⁰ In short, muscle fiber typing (type I vs II) was determined based on myosin heavy chain staining, and a CD56 antibody was used to determine SC content. Laminin was used to visualize the basement membrane, and nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI). After staining, all images were digitally captured using fluorescence microscopy (Nikon Instruments Europe). Image processing and quantitative analyses were done using the Lucia 4.81 software package. An investigator blinded to participant coding performed all image recordings and analyses. Within each image, the number of fibers, the mean fiber CSA, the number of myonuclei per fiber, and the myonuclear domain (fiber CSA/number of myonuclei) were measured for the type I and type II muscle fibers separately. For the SC slides, fiber typing was determined by matching the serial fiber typing slides. SCs were determined at the periphery of each fiber and stained positive for deoxyribonucleic acid (DAPI) and CD56. The number of SCs per muscle fiber and the percentage of SC [number of SCs/(number of SC+number of myonuclei) $\times 100$] were determined for the type I and II muscle fibers separately. A mean total of 338 ± 152 muscle fibers (178 ± 90 type I and 160 ± 77 type II muscle fibers) were analyzed for each participant. To determine interobserver reliability for the number of SCs per fiber, two investigators blinded to participant coding analyzed five randomly chosen biopsy specimens; the CV was 1.5%. Furthermore, the agreement in determination of the fiber type to which SCs belong was shown to be 99%.

Statistics

Sample size was calculated based on previous findings;^{9,36} 20 to 39 participants were needed to detect significant correlations between muscle fiber CSA, muscle mass, and muscle strength (with power = 0.80 and $\alpha = 0.05$). Kolmogorov-Smirnov tests showed that age was the only variable that was not normally distributed. To determine the relationship between age and all other variables (measured using DXA, CT, muscle biopsy, and blood sample analyses), Spearman rank correlation coefficients (r) were calculated. The relationship between the various muscle characteristics and between muscle characteristics and hormonal profiles were determined by calculating bivariate

Table 2. Characteristics of Lean Body Mass and Their Correlations with Age and Muscle Strength

Characteristic	Mean ± SD	<i>r</i> Age,		<i>r</i> 1RM-LE,		<i>r</i> 1RM-LP,	
		Mean ± SEE	<i>P</i> -Value	Mean ± SEE	<i>P</i> -Value	Mean ± SEE	<i>P</i> -Value
Whole-body lean mass, kg	56.1 ± 5.3	−0.24 ± 0.16	.07	0.41 ± 0.15	.005	0.48 ± 0.14	.001
Leg lean mass, kg	17.8 ± 1.9	−0.36 ± 0.15	.01	0.44 ± 0.14	.002	0.55 ± 0.13	<.001
Arm lean mass, kg	6.5 ± 0.8	−0.26 ± 0.16	.05	0.54 ± 0.13	<.001	0.45 ± 0.14	.002
Appendicular skeletal muscle mass, kg*	24.3 ± 2.5	−0.36 ± 0.15	.01	0.51 ± 0.14	<.001	0.57 ± 0.13	<.001
Quadriceps muscle CSA, cm ²	73.0 ± 11.3	−0.50 ± 0.13	<.001	0.72 ± 0.11	<.001	0.67 ± 0.12	<.001
Thigh CSA, cm ²	159.2 ± 19.2	−0.52 ± 0.13	<.001	0.67 ± 0.12	<.001	0.72 ± 0.11	<.001

Spearman rank (for age) and Pearson (for one-repetition maximum (1RM) for leg extension (LE) and leg press (LP)) correlation coefficients (*r*).

*Sum of lean mass in arms and legs.

SEE = standard error of the estimate; CSA = cross-sectional area.

Pearson correlation coefficients. In addition, partial correlations (adjusting for the effect of age) were calculated, eliminating age as a contributing factor to the correlation. Likewise, the correlation between age and muscle strength was adjusted for the effect of quadriceps CSA. Differences between the various correlation coefficients were tested for statistical significance.³⁷ Differences between type I and II muscle fibers were analyzed using paired-samples *t*-tests. Forward linear regression modeling was used to identify the main predictors for muscle mass and strength (quadriceps and thigh CSA, and 1RM leg press and leg extension) and muscle fiber size. Based on the results of the correlation analysis, a number of independent variables were included as potential predictors. Because age was not normally distributed, this variable was fit as a categorical variable (<70, 70–74, and ≥75). All analyses were performed using SPSS version 15.0 (SPSS, Inc., Chicago, IL). An α -level of .05 was used to determine statistical significance. Differences between type I and II muscle fibers were determined using two-sided tests. Correlation coefficients were determined using one-sided tests because specific directions were expected for the correlations studied. All data are presented as means ± standard deviations.

RESULTS

Participant characteristics are provided in Table 1. The mean age of the participants was 72 ± 5 (range 65–86). Basal blood glucose concentrations, glycosylated hemoglobin levels, and oral glucose tolerance were within the normal range for healthy older men.

Age-Related Changes

Age did not correlate with whole-body lean mass, although there was a significant negative correlation between age and regional measures of muscle mass, with the strongest correlations observed between age and total thigh and quadriceps muscle CSA (Table 2). In addition, for the leg press and leg extension exercises, muscle strength was negatively correlated with age ($r = -0.38$ and -0.37 , $P = .008$ and $.009$, respectively). When adjusted for the effect of quadriceps muscle CSA, the correlation between age and muscle strength disappeared (leg press: $r = -0.10$, $P = .52$, leg extension: $r = -0.14$, $P = .41$).

At the myocellular level, type II muscle fiber CSA was shown to be smaller than type I muscle fiber CSA (Table 3).

In addition, myonuclear and SC content were lower in the type II than the type I muscle fibers. Whereas the percentage of type I and II muscle fiber was similar, the percentage of type II muscle fiber area was smaller than of type I muscle fiber area. In contrast to type I muscle fiber CSA, type II muscle fiber CSA correlated significantly with age ($r = -0.23$; $P = .04$).

Whole-Body, Regional, and Myocellular Characteristics

All measures of lean mass (whole-body and regional lean mass) and thigh and quadriceps CSA showed positive correlations with each other and with leg extension and leg press strength (Table 2). Correlations between quadriceps CSA and leg extension strength ($r = 0.72$) and between thigh CSA and leg press strength ($r = 0.72$) were most pronounced ($P < .001$). Partial correlation coefficients showed that the relationships between muscle mass and strength did not change when adjusted for age.

For the type I and II muscle fibers, greater muscle fiber CSA was associated with more myonuclei per fiber, a greater myonuclear domain, and more SCs per muscle fiber (Table 4). In addition, the number of myonuclei per muscle fiber correlated positively with the number of SCs per muscle fiber ($r = 0.40$ for type I and II muscle fibers; $P = .006$).

Type I and II muscle fiber CSA showed a positive correlation with leg extension strength (Figure 1) but not with leg press strength. Furthermore, greater muscle fiber CSA

Table 3. Muscle Fiber Type Characteristics

Characteristic	Mean ± Standard Deviation	
	Type I	Type II
Fiber, %	52 ± 13	48 ± 13
CSA, μm^2	6,460 ± 1,662	5,276 ± 1,402*
CSA%	57 ± 14	43 ± 14*
Nuclei/fiber	3.3 ± 0.8	2.7 ± 0.7*
Nuclear domain, μm^2	2,036 ± 75	1,970 ± 79
SCs/fiber	0.087 ± 0.031	0.050 ± 0.016*
SC%	2.7 ± 0.9	1.8 ± 0.6*

*Significantly different from type I muscle fibers.

CSA = cross-sectional area; CSA% = percentage of total area occupied per fiber type; SC = satellite cell; SC% = percentage of SCs (number of SCs / (number of SCs + number of myonuclei) × 100%).

Table 4. Correlations Between Skeletal Muscle Characteristics and Muscle Fiber Cross-Sectional Area (CSA)

Characteristic	Pearson Correlation Coefficient \pm Standard Error of the Estimate, <i>P</i> -Value	
	Type I CSA	Type II CSA
Nuclei/fiber	0.56 \pm 0.13, <.001	0.56 \pm 0.13, <.001
Nuclear domain	0.41 \pm 0.15, .005	0.50 \pm 0.14, .001
Satellite cells/fiber	0.55 \pm 0.13, <.001	0.50 \pm 0.14, <.001
Quadriceps muscle CSA	0.33 \pm 0.15, .02	0.39 \pm 0.15, .006
Thigh CSA	0.30 \pm 0.15, .03	0.39 \pm 0.15, .006
One-repetition maximum		
Leg extension	0.32 \pm 0.15, .02	0.45 \pm 0.14, .002
Leg press	0.07 \pm 0.16, .33	0.14 \pm 0.16, .19

was associated with greater quadriceps CSA and greater thigh CSA (Table 4). Adjusting for age did not modulate any of the correlations.

Hormonal Profiles and Muscle Characteristics

Average serum concentrations of total testosterone, bioavailable testosterone, SHBG, IGF-I, and IGF-BP3 were 16.1 \pm 4.8 nmol/L, 5.8 \pm 1.6 nmol/L, 52.3 \pm 12.7 nmol/L, 15.1 \pm 4.8 nmol/L, and 76.1 \pm 11.8 nmol/L, respectively. Higher total testosterone concentrations were associated with higher serum SHBG concentrations ($r = 0.52$, $P < .001$). Bioavailable testosterone concentrations correlated positively with IGF-1 ($r = 0.31$, $P = .02$). In addition, a positive correlation was observed between IGF-1 and IGF-BP3 ($r = 0.46$, $P = .001$).

Higher levels of bioavailable testosterone were associated with greater ASM ($r = 0.26$; $P = .04$) and greater muscle mass measured using CT ($r = 0.27$; $P = .04$). Whereas type II muscle fiber CSA was greater with higher concentrations of bioavailable testosterone expressed as a percentage of total testosterone ($r = 0.30$; $P = .03$), this correlation was not significant for type I muscle fibers ($r = 0.23$; $P = .07$). Furthermore, higher bioavailable testosterone levels were associated with a larger percentage of fiber area occupied by type II muscle fibers ($r = 0.31$; $P = .02$). No positive correlations were observed for type I and II muscle fiber myonuclear or SC content with any of the hormones measured.

Linear Regression Analysis

Based on the outcome of the correlation analysis, potential predictors of muscle mass and strength were analyzed in a regression model. For muscle strength, the independent variables were age, quadriceps and thigh CSA, bioavailable testosterone concentration, and type I and II muscle fiber CSA. For muscle mass, the independent variables were age, bioavailable testosterone concentration, and type I and II muscle fiber CSA. For muscle fiber type I and II CSA, the independent variables were age, bioavailable testosterone concentration, and myonuclear and SC content (type I and II, respectively).

Total thigh CSA alone was shown to predict 1RM leg press (coefficient of determination (R^2) = 0.52), and quad-

iceps CSA and type II muscle fiber CSA predicted 1RM leg extension ($R^2 = 0.54$). Age and type II muscle fiber CSA significantly predicted total thigh ($R^2 = 0.28$) and quadriceps CSA ($R^2 = 0.32$). Finally, the number of myonuclei per fiber, the number of SCs per fiber, and bioavailable testosterone concentration significantly predicted type I ($R^2 = 0.49$) and type II ($R^2 = 0.48$) muscle fiber CSA.

DISCUSSION

The present study shows that muscle fiber type characteristics, bioavailable testosterone concentration, and muscle mass are closely correlated, providing important predictors of skeletal muscle mass and strength in older men. The findings indicate that SC content and muscle fiber CSA are tightly coupled, concurrent with the assumption that a decline in SC content plays an important role in type II muscle fiber specific atrophy and, as such, in the loss of skeletal muscle mass and strength in older men.

The present study shows that skeletal muscle mass correlates strongly with muscle strength in older men. Although longitudinal changes were not studied, this finding supports the concept that the loss of muscle mass is associated with muscle weakness, resulting in functional impairment in older adults.^{16,38} Previous studies with participants selected over a wide age range (18–80) reported significant correlations between strength and muscle fiber size.^{9,36} The current study extends these findings by showing that, independent of age, smaller type I and II muscle fiber size is associated with a smaller quadriceps CSA and lower leg strength in an older male population. The correlation between muscle fiber size and muscle strength tended to be stronger for type II than type I muscle fibers (Figure 1). Although speculative, this might be partly attributed to the observation that type II muscle fibers can generate greater specific tension, explaining their importance for explosive force production.³⁹ As such, specific type II muscle fiber atrophy with aging contributes to the development of muscle weakness from a quantitative (loss

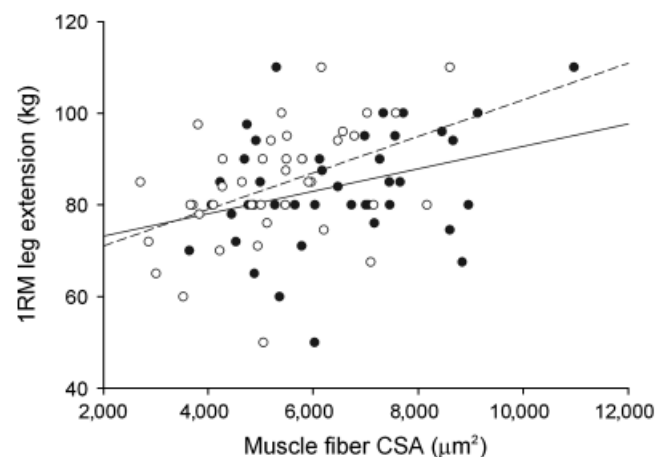


Figure 1. Scatter plot for the correlation of type I (filled circles) and type II (open circles) muscle fiber cross-sectional area (CSA) with one-repetition maximum (1RM) leg extension strength. Lines represent the fitted regression. Pearson correlation coefficients (r) tended to be stronger ($P = .06$) for type II (dashed line, $r = 0.45$, $P = .002$) than type I muscle fibers (solid line, $r = 0.33$, $P = .02$).

of muscle fiber area) and a qualitative (relative increase in type I muscle fiber area) perspective. The finding that quadriceps muscle fiber size was significantly correlated with leg extension but not leg press strength (Table 4) is probably related to the fact that leg extension represents isolated quadriceps and knee extension strength, whereas leg press represents combined hip, knee, and ankle extension strength.²⁹ As such, other factors, such as the ability to coordinate force production in different muscles and joints, may have obscured the potential correlation between muscle fiber size and leg press strength.

The correlation between age and muscle strength disappeared when corrected for muscle CSA. In accordance, regression analysis showed that muscle mass and type II muscle fiber size are the main predictors of muscle strength (explaining >50% of the variance). In general, muscle mass and quality determine muscle strength. Because age was not shown to contribute significantly to the variability in strength, it seems that age does not affect muscle quality per se. Other factors, such as type II muscle fiber atrophy, greater intramuscular lipid or connective tissue content,^{40,41} impaired neural function,⁴ and a lower physical activity level^{4,14} have been associated with poorer muscle quality and, as such, lower muscle strength. Because these aspects were not specifically addressed in the present study, their relative importance for muscle strength remains to be determined.

Type II muscle fiber CSA was shown to be smaller than type I muscle fiber CSA (Table 3). In addition, myonuclear and SC content were lower in type II than type I muscle fibers, which is consistent with previous findings.^{8,17,19,30,42} Type I and type II muscle fiber SC and myonuclear content were positively correlated with muscle fiber CSA, consistent with the myonuclear domain theory.^{43–45} It has been proposed that changes in SC content play an important role in age-related skeletal muscle atrophy^{19–21,46–48} and exercise-induced muscle hypertrophy.^{30,43} Therefore, it was speculated that skeletal muscle SC content acts as an important regulator of muscle fiber size, as the outcome of the regression analysis supports, showing that SC and myonuclear content can be predictive of muscle fiber size. As such, the muscle fiber type-specific decline in SC content observed with aging seems to represent an important factor in the age-related loss of muscle mass and strength.

Consistent with previous findings,^{2,23} a positive correlation was found between bioavailable testosterone levels and muscle mass (CT, DXA) and muscle fiber size. Together with SC and myonuclear content, bioavailable testosterone was also shown to be predictive of muscle fiber size, although despite previous findings,²⁵ no correlation was seen between testosterone and myonuclear or SC content. Although this might imply that the primary action of testosterone is not directed toward SCs, more research is warranted to address this question. It has been suggested that local and systemic IGF-1 play a role in regulating the SC cycle,^{20,49} although no correlations were observed between IGF-1 concentrations and SC content. Moreover, in contrast to previous studies,^{14,23} no correlations were observed between IGF-1 and various measures of muscle mass. The small variability in IGF-1 and muscle mass in the participant population included in the present study might explain these findings. Alternatively, it might be that

muscle-specific IGF-1 is a stronger predictor of muscle mass and myogenic potential, and future research should further delineate the specific roles of muscle-specific and systemic IGF-1 in regulating skeletal muscle mass.⁴⁹

The present cross-sectional analysis does not allow any direct causal relationships to be determined. Age-related changes in muscle characteristics (e.g., muscle fiber atrophy) generally represent slowly progressing processes, taking 20 to 30 years to become apparent.⁸ As such, longitudinal studies are difficult to assess because of methodological problems regarding follow-up duration and high dropout rates. In the present study, whether the known age-related changes in muscle fiber characteristics can explain the large variance in muscle mass and strength in a population of community-dwelling elderly men was therefore assessed. This is the first study to show predictive associations between SC content, muscle fiber size, and skeletal muscle mass and strength in elderly men.

The sample size of the present study and the inclusion of only relatively healthy elderly men may limit its potential for generalization to other subpopulations such as women or frail older men. Nonetheless, combined with the findings from recent intervention studies^{25,30} and *in vivo* animal experiments,^{21,43,46} the current data indicate that SCs play an important role in the age-related loss of muscle mass and function.

In conclusion, muscle mass and muscle fiber size are predictive of skeletal muscle strength in elderly men. Furthermore, SC content, myonuclear content, and bioavailable testosterone concentration seem to play important roles in regulating muscle fiber size and, as such, skeletal muscle mass and strength in older men. These findings provide further support to the idea that a decline in type II muscle fiber SC content plays an important role in the loss of muscle mass and strength with aging.

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