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## Tendon-Derived Stem/Progenitor Cell Aging: Defective Self-renewal and Altered Fate

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### Summary

Aging is a major risk factor for tendon injury and impaired tendon healing, but the basis for these relationships remains poorly understood. Here we show that rat tendon-derived stem/progenitor cells (TSPCs) differ in both self-renewal and differentiation capability with age. The frequency of TSPCs in tendon tissues of aged animals is markedly reduced based on colony formation assays. Proliferation rate is decreased, cell cycle progression is delayed and cell fate patterns are also altered in aged TSPCs. In particular, expression of tendon lineage marker genes decreased while adipocytic differentiation increased. Cited2, a multi-stimuli responsive transactivator involved in cell growth and senescence, was also downregulated in aged TSPCs while CD44, a matrix assembling and organizing protein implicated in tendon healing, was upregulated, suggesting that these genes participate in the control of TSPC function.

### Keywords

aging; tendon-derived stem/progenitor cells; self-renewal; differentiation; Cited2; CD44

### Introduction

Age is a major risk factor for tendon disorders. Age-related changes in structural and mechanical properties may predispose tendons to injury; moreover, tendon healing is often impaired in the elderly (Vogel 1978; Birch *et al.* 1999; Magnusson *et al.* 2003; Coupee *et al.* 2009). Tenoblasts and tenocytes, the major tendon cellular elements that produce and organize tendon extracellular matrix (ECM) (Kannus 2000), also undergo age-dependent changes in number and activity (Ippolito *et al.* 1980; Nakagawa *et al.* 1994), but the mechanisms behind these changes remain unclear. Recently, tendons of several species (Zhang & Wang 2010; Bi *et al.* 2007; Rui *et al.* 2009) were shown to contain a small population of cells with stem cell properties, termed tendon-derived stem/progenitor cells (TSPCs). Since adult or tissue-resident stem/progenitor cells are considered essential for tissue maintenance and repair, we investigated whether the self-renewal capacity and differentiation potential of rat TSPCs are influenced by age.

### Results and discussion

TSPCs were prepared from rat tendon tissues based on their preferential attachment and potent clonal expansion over the majority of resident tendon cells (Bi *et al.* 2007). We found

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that nearly 100% of both young and aged TSPCs stained positively for three stem cell markers: nucleostemin, Oct-4, and SSEA-4 (Suppl. Fig. 1), indicating that this population retains features reflecting their stemness regardless of age. Rat TSPCs also expressed surface antigens CD44 (Fig. 1 A, left panels) and CD90.1 (Fig. 1 A, middle panels), but not the endothelial cell marker CD106 (VCAM-1) (Fig. 1 A, right panels), as shown for murine and human TSPCs (Bi *et al.* 2007) and recently reported for rat TSPCs (Rui *et al.* 2009). Aged TSPCs expressed lower levels of CD90.1 than young cells, but higher levels of CD44 as determined by both the percentage of positive cells (Fig. 1 A, left panels) and mean fluorescence intensity (Fig. 1B). These differences in CD44 expression were also seen at mRNA levels by RT-PCR (Fig. 1C). CD44, implicated in healing of many tissues, is downregulated during scarless fetal tendon healing (Favata *et al.* 2006); moreover, mice genetically deficient in CD44 showed improved patellar tendon healing (Ansoerge *et al.* 2009). These findings suggest that the increased CD44 expression in aged TSPCs may contribute to reduced TSPC repair capacity with age.

TSPCs accounted for a dramatically smaller fraction of total tendon cells in old rats ( $0.171\pm 0.06\%$ ) than young rats ( $6.26\pm 0.55\%$ ,  $P<0.001$ ) (Fig. 1D) based on the number of colonies formed by whole tendon-derived cells (TDC) (Bi *et al.* 2007; Delorme & Charbord 2007). This corresponded to a 70% reduction in the total number of cells recovered from old versus young tendons. However, colony formation assays using P0 TSPCs from aged and young rats showed only small differences (data not shown). These results indicate that diminished colony formation by aged TSPCs primarily were due to low TSPC numbers likely reflecting a depleted TSPC pool in aged tendon tissues.

Proliferation of aged and young TSPCs also differed. Cell numbers after 8 days of culture were lower in aged TSPCs ( $1.08\pm 0.09\times 10^6$  vs.  $1.91\pm 0.16\times 10^6$ ,  $P<0.05$ , Fig. 1E, upper panel), with a corresponding increase in estimated mean population doubling time (DT) ( $18.8\pm 0.22$  hrs vs.  $17.4\pm 0.20$  hrs,  $P<0.05$ , Fig. 1E, lower panel). Similar differences were observed in CFSE-dilution assays (Lyons 2000) assessed by FACS (data not shown). Analysis of cell cycle phase distribution using propidium iodide further showed that aged TSPCs contained a higher fraction in G2/M (Fig. 1F), suggesting that aged TSPCs were preferentially subject to late cell cycle arrest. This could result from accumulated genetic and/or epigenetic damage, as has been reported in other stem/progenitor cell populations (Rossi *et al.* 2007). Additionally, differences in apoptotic rates between young and old TSPCs could also contribute to the observed disparities in population size. However, to our knowledge, neither issue has yet been examined in TSPCs. Collectively, these data indicate that TSPCs undergo age-related declines in self-renewal capacity (Nishimura *et al.* 2005; Levi & Morrison 2008) that could account for reduced stem cell numbers with aging.

The ability of TSPCs to differentiate into tenocytes was also diminished by age. Expression of two tendon lineage-specific genes (Perez *et al.* 2003; Shukunami *et al.* 2006; Murchison *et al.* 2007), Scleraxis (Scx) and Tenomodulin (Tnmd), was lower in aged TSPCs than in young cells (Fig. 2A). Aged TSPCs also showed diminished induction of these markers by TGF- $\beta$ 3 (Kovacevic & Rodeo 2008). Whether these differences also extend to other tenocyte lineage-inductive stimuli like mechanical loading (Juncosa-Melvin *et al.* 2006; Kuo & Tuan 2008), and whether they reflect differentiation capacity *in vivo* remain to be determined. Interestingly, however, aged TSPCs formed adipocytes more readily than younger cells (Fig. 2B) and expressed higher levels of adipogenic markers PPAR $\gamma$ 2 (PPARGC1A), C/EBP $\alpha$  (Cebpa/CEBPA), and leptin following induction (Fig. 2C). Young and old TSPCs showed no apparent difference in the ability to form osteoblasts or chondrocytes (Suppl. Fig. 2). These data may help explain the higher levels of adipose tissue normally associated with older tendons (Kannus & Jozsa 1991), a pattern similar to that

observed in bone marrow, where adiposity was found to correlate inversely with the functionality of hematopoietic stem/progenitor cells (Naveiras *et al.* 2009).

Finally, we explored age-dependent changes in a potential regulator of TSPC function. Cited2 is a transcription factor implicated in the control of growth and senescence in several cell types (Sun *et al.* 1998; Kranc *et al.* 2003; Yokota *et al.* 2003; Sun 2009). Recent studies further revealed that Cited2 is required to maintain adult hematopoietic stem cells (Chen *et al.* 2007; Kranc *et al.* 2009). We found that Cited2 expression in aged TSPCs was reduced at both the mRNA (Fig. 2D) and protein levels (Fig. 2E). These data are consistent with positive roles for Cited2 in TSPC self-renewal. Furthermore, the coordinated expression of Cited2 and Scx suggests that it may also regulate TSPC differentiation.

The present study has demonstrated remarkable changes in number and function of tendon stem/progenitor cells with advancing age. It remains to be determined whether these age-related changes are also influenced by factors such as activity, which could affect tendon loading history. Second, while the functional characterizations of rat TSPCs *in vitro* may reflect properties of these cells, including the adaptation to an *in vitro* environment, further studies are also needed to determine whether TSPCs exhibit age-dependent differences in the ability to repopulate functional tenocyte pools *in vivo*.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

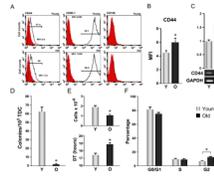
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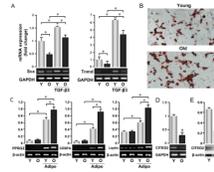
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### Figure 1. Aging-related cellular changes in TSPC self-renewal

Young and aged TSPCs were prepared, as described by Bi et al (Bi *et al.* 2007), from 3–4 and 24–26 month Sprague-Dawley male rats, respectively. Patellar tendons were digested with collagenase A/dispase (2 h, 37 °C) and cultured in DMEM plus 10% FBS for 7–9 days. Adherent cells (passage 0, P0) at the end of culture were used as TSPCs for all assays unless otherwise specified. (A–C) Age-related surface antigen expression changes in TSPCs. A: Representative histogram plots. TSPCs stained with fluorescein-conjugated anti-rat antibodies (open) or isotype control antibodies (filled) were analyzed by FACS. B: Mean fluorescence intensity of CD44 based on FACS analysis of surface expression as shown in A (n=3, independent experiments, \* $P < 0.05$ ). C: CD44 mRNA levels determined by RT-PCR. Upper Panels represent the quantification of band intensity from the gel shown in lower panels. (D) Colony formation. Total tendon-derived cells (TDC) were plated at  $1 \times 10^3$  cells/well (6-well plate) and grown for 9 days. Colony forming units (CFUs) were scored after methylene blue staining. Error bars represent SD (n=3, \* $P < 0.05$  versus aged TSPC). (E) Proliferation. TSPCs were seeded at  $1 \times 10^3$  cells/well (6 well plate) and cultured for 8 days. Cell numbers were counted and cell population doubling times were estimated from start (0) and end (1) points. Error bars represent SD (n=3, \* $P < 0.05$  versus aged TSPC). (F) Cell cycle distribution. TSPCs were fixed with 70% ethanol, stained with propidium iodide (PI), and analyzed by FACS. Error bars represent SD (n=3 independent experiments; \*  $P < 0.05$ )



### Figure 2. Altered cell fate of aged TSPCs

P0 TSPCs were prepared as described in Figure 1 and used for all assays. (A) Expression of tendon lineage-specific genes. TSPCs were cultured with or without TGF- $\beta$ 3 for 3 days, and total RNA was extracted after culture. (B, C) Adipocyte-skewed differentiation of aged TSPCs. Cells were cultured in specific induction medium for 16 days. B: Oil Red O staining per (Gimble *et al.* 1995); C: mRNA expression of adipogenic marker genes in untreated and adipogenic-induced TSPCs. (D, E) Expression of Cited2 at mRNA (D) and protein (E) levels in young and old TSPCs. mRNA expression of indicated genes in A, C, D was assessed by RT-PCR. Upper Panels represent the quantification of band intensity from the gel shown in lower panels. Data are representative of 3 experiments and confirmed by real-time PCR.