

Type II TGF β receptor modulates chondrocyte phenotype

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1	Type II TGFbeta receptor modulates chondrocyte phenotype.
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Abstract:

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- 2 Aging is one of the major risk factors of osteoarthritis. This pathology during which chondrocytes undergo
- 3 modifications of their phenotype may result from alteration of TGFβ signaling. This study investigates the role of
- 4 TGFß response in process of chondrocyte dedifferentiation/redifferentiation.
- 5 Dedifferentiation was induced by successive passages of human articular chondrocytes, whereas their
- 6 redifferentiation was performed by tridimensionnal culture in alginate. Human mesenchymal stem cells were
- 7 obtained from bone marrow and differentiated into chondrocyte-like phenotype by 3D-culture, embedded in the same
- 8 scaffold. Protein and mRNA levels were analyzed by Western-Blot and real time RT-PCR. Regulatory mechanism
- 9 was investigated using specific inhibitors (mithramycin), mRNA silencing or decoy oligonucleotides, and expression
- 10 vectors.
- 11 Chondrocyte dedifferentiation interfered with TGFB signalling by decreasing TBRII mRNA and protein levels and
- 12 subsequent TGF\$\beta\$ response. TBRII ectopic expression in passaged chondrocytes permitted to increase the expression
- 13 of several matrix genes, such as aggrecan or type II collagen. Redifferentiation of passaged chondrocytes permitted
- 14 to restore, at least in part, TBRII expression, and was related to differentiation of human bone marrow mesenchymal
- 15 stem cells toward chondrocytes, where both Sp1 and TBRII mRNA levels were increased. Moreover, Sp1
- 16 manipulation by silencing or ectopic expression and pharmacologic inhibition revealed a link between expression
- 17 levels of this transcriptional factor, which is crucial for constitutive expression of TβRII in cartilage, and TGFβ
- 18 response.

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- 19 Therefore, these data permit us to suggest an important role of TBRII expression in maintenance of chondrocyte
- 20 phenotype, which is altered with age, and bring new insights in our understanding of chondrogenesis process.

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Keywords: TGFbeta receptors, chondrocytes, dedifferentiation/differentiation, Sp1, osteoarthritis, aging

INTRODUCTION

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2 Articular cartilage consists of an abundant extracellular matrix, which is produced by chondrocytes. These highly 3 differentiated cells are responsible for the maintenance of the cartilage as a functional entity. Aging cause changes in 4 the properties of this tissue, notably by modifying the matrix composition or the growth factor responsiveness 5 (Kempson, 1982; Thonar et al., 1986; Dozin et al., 2002; Loeser et al., 2000; Rosen et al., 1997; Guerne, et al., 6 1995). During embryonic development, cartilage formation is initiated with condensation of mesenchymal cells, 7 followed by progression of chondrocyte differentiation toward proliferation, prehypertrophy and hypertrophy. 8 During osteoarthritis (OA), an articular disease strongly correlated with ageing (Buckwalter and Mankin, 1998, van 9 der Kraan et van den berg, 2008), chondrocytes undergo morphogenic alterations accompanied by profound 10 biochemical changes including the loss of production of cartilage-specific macromolecules. For instance, type II 11 collagen expression decreased wheareas type I collagen synthesis increases. Some of the alterations of chondrocyte 12 phenotype can be reproduced in vitro. Passages induce morphogenic alterations of cells, reduction of type II 13 collagen, and induction of type I collagen synthesis (Benya et al., 1978; Archer et al., 1990; Pei et al., 2000; 14 McDougall et al., 1996, Homicz et al., 2002). However, this model is not able to mimic others modifications 15 undergone by OA chondrocytes, and notably the induction of type X collagen, specific of hypertrophy stage. On the 16 other hand, the chondrocyte-specific phenotype can be recovered when the cells are cultured in three-dimensional 17 culture system, such as alginate matrices (Reginato et al., 1994; Bonaventure et al., 1994). 18 Several studies suggest that modifications of chondrocyte phenotype during OA result from alteration of 19 Transforming Growth Factor β (TGF β) signaling. Thus, the expression of TGF β and its signaling molecules is 20 altered with age, and the subsequent lack of TGF\$\beta\$ responsiveness might be at the root of OA development (Blaney-21 Davidson et al., 2005). Besides, TGFβ signaling is important for chondrogenesis. It promotes the differentiation of 22 embryonic chick limb cartilage (Schonherr et al., 2000), and upregulates a number of molecules associated with 23 prechondrogenic condensation (Serra et al., 1997; Chimal-Monroy et al., 1999; Grimaud et al., 2002). In addition, 24 primary chondrogenesis derived from mesenchymal stem cells (MSCs) needs TGFB signals (Reddi, 1994; 25 Andriamanalijaona et al., 2008). 26 Members of TGFß family transduce their signal via two serine/threonine receptors, type I (TßRI) and type II (TßRII) 27 receptors (Wrana et al., 1994). Upon ligand-binding, TßRI is phosphorylated by TßRII, resulting in activation of 28 intracellular proteins, named Smads. Receptor-regulated Smads (R-Smads: Smad2 and Smad3) are phosphorylated 29 by TBRI. Phosphorylated R-Smads form heterodimers with common-partner Smad (Co-Smads: Smad4), and 30 translocate into the nucleus where they bind to specific DNA sequences for activation of TGFB target gene 31 transcription (Shi et al., 2003). Inhibitory Smads (I-Smads: Smad7) inhibit phosphorylation of R-Smads by 32 competing with their binding to activated TBRI (Nakao et al., 1997). Recently, it was shown that besides signaling 33 via TßRI (Activin-like kinase 5, ALK5), TGF-beta can also signal via ALK1, thereby activating Smad1/5/8 instead 34 of Smad2/3 (Blaney-Davidson et al., 2009; Finnson et al., 2008). 35 We hypothesised that the maintenance of chondrocyte phenotype, which is altered with age, required a functional 36 TGFβ signalling. We examined the influence of members of TGFβ signalling on chondrocyte phenotype. We 37

showed that TBRII is implicated in the loss of TGFB responsiveness observed during chondrocytes passages. TBRII

- 1 ectopic expression permits to restore TGFB responsiveness as well as to increase the expression of some matrix
- 2 components. We also found that $T\beta RII$ expression level is intimately linked to differentiation status of chondrocytes.
- 3 Finally, we established a relationship between TGFβ-induced effects on matrix genes, and expression level of
- 4 Specific protein 1 (Sp1) transcription factor, which expression is decreased in the course of dedifferentiation process
- 5 (Duval et al., 2011) and is require to ensure constitutive expression of TβRII in chondrocytes (Baugé et al., 2008).

8

MATERIAL AND METHODS

9 Reagents.

- 10 Reagents were supplied by Invitrogen (Fisher Bioblock Scientific, Illkirch, France) unless otherwise noted.
- 11 Mithramycin (Sigma-Aldrich, St. Quentin Fallavier, France) and TGF\$1 (R&D Systems, Lille, France) were
- 12 resuspended in PBS or PBS-HCl, respectively. Alginate was provided by Sigma-Aldrich, and oligonucleotides were
- supplied by Eurogentec (Angers, France).

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Chondrocyte culture

- Human articular chondrocytes (HAC) were prepared from femoral head of post-mortem individuals with no known
- joint diseases or after femoral head fractures (ages between 45 and 80 years with median age of 68). The causes of
- donor's death were multiple and all of them signed agreement forms in the context of a body donation program
- according to local legislations. Similarly, donors from arthroscopy signed the agreement for this study according to
- 20 local ethical comitee (CPP). Chondrocytes were released by enzymatic digestion as previously described (Baugé et
- al., 2007). Cells were seeded at 4x10⁴ cells/cm² and cultured in Dulbecco's modified Eagle's medium (DMEM) (4.5
- 22 g/L D-glucose) supplemented with 10% heat-inactivated foetal calf serum (FCS), 100 IU/ml penicillin, 100 μg/ml
- 23 streptomycin and 0.25 µg/ml fungizone, in a 5 % CO₂ atmosphere. For dedifferentiation, HAC were passaged up to
- five times (P5) (about 7-8 population doubling) with trypsin-EDTA after reaching confluency.
- 25 For re-differentiation studies, chondrocytes, which had underwent three passages (about 4-5 population doubling),
- were encapsulated in alginate beads. Cells were suspended in alginate at the density of 2.5x10⁶ cells/ml, then slowly
- dropped into a 100 mM CaCl₂ solution. After instantaneous gelation, the beads were allowed to polymerize further
- 28 for a period of 10 min in CaCl₂ solution. After three washes in PBS, they were finally placed in DMEM
- supplemented with 10% foetal calf serum. After 3 weeks, the beads were rinsed with PBS and dissolved in a solution
- 30 composed of 55 mM sodium citrate and 25 mM EDTA. The suspension was incubated at 37°C until the beads were
- 31 completely dissolved (approximately 10 min.). Centrifugation of the suspension at 800 g for 10 min allows
- separating the cells from their alginate matrix.

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Culture of human bone marrow-derived stem cells

- Human bone marrow-derived stem cells were collected from the same patients as donors used for HAC isolation.
- The harvested cells were washed twice with α -MEM, and isolated by adherence over 24h to tissue culture plastic and
- 37 were expanded in monolayer culture with α-MEM supplemented with glutamine, FGF (1 ng/ml), 10% FCS and

- 1 antibiotics. After amplification by successive passages (until P5), the absence of hematopoietic markers (such as
- 2 CD45) was checked by RT-PCR (data not shown). Thereafter, bone marrow-derived stem cells were directed toward
- 3 chondrogenesis by culture in 3-D scaffold (alginate bead).
- 4 For chondrogenic differentiation, isolated human bone marrow derived stem cells were encapsulated in alginate
- 5 beads, according to the same protocol than HAC, and cultured with a chondrogenic medium consisting of DMEM
- 6 containing 100 nM dexamethasone, 50 μg/ml ascorbic acid-2-phosphate, 40 μg/ml proline, 1mM sodium pyruvate
- and Insulin Transferin Selenium supplement (ITS, Sigma), for 21 days and the media were changed twice a week.

Plasmids and transfection experiments

- 10 p3TPLux, pCMV5-TßRII, pEVR2-Sp1 expression vectors were described earlier (Baugé et al., 2008). Chondrocytes
- were transiently transfected by the calcium phosphate precipitation method as previously described (Baugé et al.,
- 12 2007). After overnight transfection, the culture medium was replaced by DMEM containing 10% FCS, and
- maintained in culture for 24 h.

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Luciferase assay

- 16 Luciferase assay was performed with Luciferase Assay kit from Promega (Charbonnières, France), as previously
- 17 described (Baugé et al., 2007).

18 19

Gene silencing experiments

- 20 1.5 μg of double strand siRNA targeting Sp1, provided from Santa-Cruz Biotechnology (through Tebu, Le Perray en
- 21 Yvelines, France), were nucleofected in HAC, according to the manufacturer's protocol with human chondrocyte
- Nucleofector solution (Amaxa, Köln, Germany). Negative control siRNA known to not lead to specific degradation
- 23 of any cellular message was also used. After 6 h of transfection, the culture medium was replaced with DMEM
- containing 10% FCS and the chondrocytes cultures were incubated for 24 h.

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RNA extraction and real-time RT-PCR.

- 27 Total RNA from primary HAC cultures were extracted using Trizol. Following extraction, 1 µg of DNase-I treated
- 28 RNA was reverse transcribed into cDNA in the presence of random hexamers (Applied Biosystems, Courtaboeuf,
- France) and Moloney murine leukaemia virus reverse transcriptase. The reaction was carried out at 37°C for 1h
- followed by a further 10-min step at 95°C. Amplification of the generated cDNA was performed by real-time PCR in
- 31 Applied Biosystems SDS7000 apparatus with appropriate primers designed with Primer Express software. The
- 32 relative mRNA level was calculated with the $2^{-\Delta\Delta CT}$ method.

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Protein extraction and Western-blot.

- 35 Cells were rinsed, scrapped in RIPA lysis buffer supplemented with phosphatase and protease inhibitors. The
- 36 extracts (30 µg protein) were subjected to fractionation in a 8 or 10% SDS-PAGE, transferred to PVDF membranes
- 37 (Amersham Biosciences, Orsay, France), and reacted with rabbit aggrecan, TβRII or Sp1 polyclonal antibodies.

1 Subsequently, membranes were incubated with anti-rabbit secondary peroxidase-conjugated antibody. The signals 2 were revealed with SuperSignal West Pico Chemiluminescent Substrate (Pierce Perbio Science, Brébières, France) 3

and exposed to X-ray film. The membranes were also reacted with anti β -actin to verify equal loading.

4 5

Synthesis and transfection of decoy oligonucleotides.

- 6 Sp1 decoy oligonucleotides sequences are: Forward 5'- ATTCGATCGGGGGGGGGGGGGG-3' and Reverse 5'-
- 7 CGACGCCCGGCCCGATCGAAT -3' (consensus sequences are underlined). This decoy oligonucleotide was
- 8 previously shown to bind Sp1 transcriptional factor (Baugé et al., 2008). These oligonucleotides were annealed
- 9 overnight and transfected into cells using the Oligofectamine reagent according to the manufacturer's instructions.

10

11 **Nuclear extracts and EMSA**

- 12 Nuclear extracts were prepared as described previously (Dignam et al., 1983). For EMSA, 5 fentomoles of (γ-
- 13 ³²P)ATP-labelled probes (Sp1 binding sequence) and 4 µg of nuclear extracts were incubated at room temperature
- 14 for 15 min in binding buffer (20 mM Tris-HCl pH 7.5, 1 mM DTT, 100 mM NaCl, 5% bovine serum albumin, 0.1%
- 15 Nonidet P-40, 10% glycerol, 25 µM ZnCl₂) and 2 µg of poly(dI-dC). (dI-dC). Samples were resolved on native
- 16 polyacrylamide gel and exposed for autoradiography.

17 18

Statistical analysis:

- 19 All experiments were repeated at least with 3 different donors, and similar results were obtained. Representative
- 20 experiments are shown in the figures. Data are presented as the mean \pm SD. Statistical significance was determined
- 21 by Student's t-test. P values less than 0.05 were considered significant. ***= P < 0.001; *= P < 0.01; *= P < 0.05
- 22 versus controls.

RESULTS

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- 2 Chondrocyte passages reduce the TGFB responsiveness and decrease TBRII expression.
- 3 In order to ascertain that subculturing chondrocytes led to the loss of cartilaginous phenotype in our hands, we
- 4 monitored the decrease of type II collagen and aggrecan mRNA expression after successive passages. Their
- 5 expressions were strongly decreased as soon as the first passage (about 2 cell doubling). The maximal reduction was
- 6 obtained at P2 (approximatively 3 doubling) and was maintained at least until P5. At opposite, collagen type I and III
- 7 mRNA levels were higher at P2 (suppl figure 1A). Therefore, we limited the following experiments to 2 or 3
- 8 passages (3-5 cell doubling).
- 9 To investigate chondrocyte responsiveness to TGFB during dedifferentiation process, we focused on the expression
- 10 of Plasminogen Activator Inhibitor-1 (PAI-1), which is rapidly induced by TGFβ in chondrocytes (Baugé et al.,
- 11 2007). Human articular chondrocytes which underwent three passages were incubated for 4 h with TGFβ1 (2 ng/ml).
- 12 Figure 1A shows that passages induced a significant decrease of PAI-1 induction by TGFB.
- 13 TGF\$\beta\$ initiates its cellular action by binding to T\$\beta\$RII, which in turn activates T\$\beta\$RI. Thus, we investigated the effect
- 14 of passages on mRNA expression of both receptors. Whereas TBRI transcripts were increased in comparison to
- 15 primary chondrocytes (P0) (fig. 1B), TβRII is dramatically decreased following dedifferentiation (fig. 1C). This
- 16 reduction of TBRII expression was also observed at protein level (fig 1D).

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Passages increases the expression of the R-Smad, Smad3, and decreases that of the I-Smads, Smad7.

- 19 Since signalling by TGF\$\beta\$ is mediated by Smads, we explored the potential alteration of Smads expression during
- 20 chondrocyte dedifferentiation. Our result (fig 2) showed that serial passages of chondrocytes in serum supplemented
- 21 medium did not modulate significantly Smad2 and 4 mRNA levels. Moreover, they increased Smad3 mRNA levels
- 22 whereas they decreased the expression of the I-Smad, Smad7. These data suggest that the loss of TGFB
- 23 responsiveness observed in dedifferentiated chondrocytes is independent of Smads expression level, since an up-
- 24 regulation of Smad3 and down-regulation of Smad7 is usually associated with activation of TGFß signalling.

25 26

Ectopic expression of TBRII restores TGFB responsiveness in passaged chondrocytes and increases expression

27 of matrix genes.

- 28 To confirm the crucial role of TBRII expression alteration in the reduction of TGFB sensitiveness of passaged
- 29 chondrocytes, we examined the effect of forced TBRII expression on the induction of PAI-1 by TGFB (fig 3A). HAC
- 30 were transfected with a TβRII expression vector (pCMV5-TβRII) or its corresponding insertless vector (pCMV5) for
- 31 24h and further incubated with TGFβ. Dedifferentiation clearly causes reduced TGFβ-induced PAI-1 expression,
- 32 which was restored by TβRII ectopic expression. In addition, transfection of TβRII expression vector in passaged
- 33 chondrocytes induces an increased mRNA level the expression of aggrecan and collagen type II (x2-3) (fig 3B-C).

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TBRII expression increases during chondrocyte differentiation

- 36 Since TBRII levels decreased upon dedifferentiation, we hypothesized that its expression could also be modulated
- 37 during chondrocyte differentiation process. Therefore, we pursued by examination of TBRII expression in re-
- 38 differentiating chondrocytes. For that, the TBRII mRNA level was evaluated prior dedifferentiation (P0), after

- 1 dedifferentiation (P3) and after redifferentiation in alginate. TBRII mRNA level was increased after a 3 week-culture
- 2 in alginate (fig 4A), time which permit to restore (at least partially) a chondrocyte phenotype in this system (Duval et
- 3 al., 2011).
- 4 Similar experiments were performed using human bone marrow-derived mesenchymal stem cells, which underwent
- 5 chondrogenesis by culture in alginate microbeads (Duval E/Boumediene K, unpublished data; suppl data 2B). This
- 6 system is well-known to induce chondrogenic differentiation from mesenchymal stem cells from different origins
- 7 (Andriamanalijaona et al., 2008). As shown in figure 4B, TBRII mRNA levels strongly increased with the culture
- 8 time, and subsequently increased during differentiation of hBM derived-MSC into chondrocyte-like phenotype.

Sp1 is important for $T\beta RII$ expression and the maintenance of chondrocyte phenotype

- Given that Sp1 expression decreases during chondrocyte dedifferentiation and is restored by alginate culture (Duval
- et al., 2011), we proposed that Sp1 expression and activity could increase during differentiation of stem cells. Sp1
- mrnA level was raised during human bone marrow derived stem cells differentiation into chondrocytes (Fig 5A).
- To define whether DNA binding activity of Sp1 is also increased in these conditions, EMSAs were performed using
- 15 the consensus DNA binding site as a probe (fig 5B). One delayed protein-DNA complex was strongly enhanced by
- 16 hBM-MSC culture in alginate beads after 72h or 96h. The binding activity of the complex to the probe was
- decreased in the presence of Sp1 antibody (and not in the presence of irrelevant antibodies, namely against cKrox or
- Sp3), indicating that this complex contains Sp1 factor.
- 19 Then, we investigated the effect of inhibition of Sp1 on TBRII expression. First, we blocked the binding activity of
- 20 Sp1 family members using decoy oligonucleotides or mithramycin (Jia et al., 2007) and showed that the loss of Sp1
- 21 binding to DNA inhibited TBRII mRNA level-expression (fig 6A-B). Furthermore, we established that the specific
- 22 reduction of Sp1 expression by RNA silencing, checked by Western-blot, reduced TBRII expression (fig 6C).
- 23 Mithramycin also reduced TBRII protein expression. Interestingly, aggrecan expression was also decreased by
- 24 mithramycin (figure 6D).
- 25 Finally, in figure 7, we showed that Sp1 ectopic expression is able to restore TBRII expression in dedifferentiated
- 26 chondrocytes and counteracts inhibition of aggrecan expression induced by passages.

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DISCUSSION

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Modifications of chondrocyte phenotype occur with aging and during osteoarthritis. They can be mimicked, in part, in vitro by subculturing monolayer cultures of chondrocytes (Homicz et al., 2002; Pei et al., 2000; McDougall et al., 1996; Duval et al., 2011). This culture model provides a way to understand some of the molecular mechanisms responsible for chondrocyte senescence and dedifferentiation, and to elucidate the transition from young/healthy to old/osteoarthritic cartilage. Conversely, studies of mechanism of re-differentiation of chondrocyte or differentiation of mesenchymal cells into chondrocytes can permit to bring new insights in order to develop the therapeutic use of mesenchymal stem cells (MSC) in cartilage disorders. In this study, we showed that chondrocyte dedifferentiation is accompanied by TBRII expression alteration, whereas redifferentiation of passaged chondrocytes or differentiation of MSC into chondrocytes are accompanied by TBRII production. Moreover, we demonstrated that forced TBRII expression is able to restore totally or partially the expression of matrix genes, suggesting a key-role of this receptor and TGFß signal in the maintenance of chondrocyte phenotype. Our in vitro experiments show the key-role of TBRII expression in chondrocyte phenotype maintenance. This result completes in vivo data which showed TBRII-deficient mice have a reduced proliferation of chondrocytes and an accelerated early hypertrophic differentiation relative to control (Seo et al., 2007). In addition, conditional transgenic mice in which a dominant negative form of TBRII is expressed, develop a hypoplastic cartilage (Hiramatsu et al., 2011), confirming that TBRII is required for normal cartilage development, and subsequently chondrocyte phenotype. Chondrocyte dedifferentiation, which occurs with aging, is likely associated to alteration of cell responsiveness to TGFB by reduction of TBRII expression. Of interest, we have recently demonstrated that interleukin-1B, a proinflammatory cytokine which is increased in OA process, rendered cells insensitive to TGF\$\beta\$ by reducing T\$\beta\$RII expression (Baugé et al., 2007). Together, these results corroborate in vivo data, which indicated that TBRII expression is dramatically reduced in old mice (Blaney Davidson et al., 2005) and during OA (Boumediene et al., 1998; Verdier et al., 2005), and confirmed the relationship between TBRII expression and the reduction of cell responsiveness to TGFB observed in OA and old cartilage. Concerning TBRI expression, it increases with passages, suggesting that this receptor is not involved in the reduction of cell responsiveness to TGFB in passaged chondrocytes. This result was unexpected since TBRI expression is reduced during experimental OA induced in mice by destabilization of the medial meniscus model (Blaney-Davidson et al., 2009). In addition, we examined the expression levels of TGFB signalling intracellular mediators, namely Smads. We showed that Smad3 expression is increased during chondrocytes dedifferentiation whereas that of Smad2 is not modified. Similar effects were obtained in vivo with aging (Blaney-Davidson et al., 2005). Smad3 acts as one important TGFB signalling pathway members to develop and/or maintain the phenotype of chondrocytes. Smad3, but not Smad2, stimulates primary chondrogenesis (Furumatsu et al., 2005). In addition, Smad3 deficiency accelerates chondrocytes maturation and leads to osteoarthritis (Yang et al., 2001; Van de Laar et al., 2011). Therefore, we speculate that the increased Smad3 expression observed during dedifferentiation and aging could be a compensatory mechanism to promote the chondrocyte phenotype. Moreover, these results obtained on Smad expression could be due to the loss of TGFß response since TGFß reduces Smad3 expression (Mori et al., 2000; Baugé et al., 2011).

We also observed the TBRII expression during in vitro chondrogenesis. Three-dimensional scaffold was used to redifferentiate dedifferentiated chondrocytes or to differentiate MSC toward chondrogenic phenotype. This culture system was described earlier to permit the induction of chondrogenic markers (Reginato et al., 1994; Bonaventure et al., 1994; Andriamanalijaona et al., 2008; Lemare et al., 1998; Cai et al., 2007; Duval et al., 2011). We showed that TBRII expression is induced during redifferentiation of subcultured chondrocytes and during chondrogenic process of stromal stem cells. In this regard, Mizuta and coll. found that expression of both TBRI and TBRII receptors was upregulated during periosteal chondrogenesis induced by exogenous TGF\$1 treatment (Mizuta et al., 2002). Our data corroborate another in vitro study which proposes that TGFB signal transduction may be involved in the early stage of bone marrow-MSC chondrogenesis (Huang et al., 2005), and are consistent with recent studies, which demonstrated that TBRII down-regulation induced by retinoid acid treatment inhibits chondrogenesis of mouse embryonic palate mesenchymal cells (Yu et Xing, 2006) and that TGF\$\beta\$ signalling is needed in the chondrogenic differentiation of MSC (Longobardi et al., 2006). Finally, we investigated the role of Sp1 role in the regulation of TBRII expression and matrix production. This ubiquitous zinc-finger protein has earlier been proposed to regulate matrix production by adult articular chondrocytes maintained in primary and expansion culture (Duval et al., 2011; Dharmavaram et al., 1997; Oh et al., 2007). Furthermore, we have demonstrated loss of Sp1 gene expression as a function of chondrocyte passage number and that Sp1 protein synthesis is restored in part by re-differentiation in 3D-alginate culture (Duval et al., 2011). Here,

maintained in primary and expansion culture (Duval et al., 2011; Dharmavaram et al., 1997; Oh et al., 2007). Furthermore, we have demonstrated loss of Sp1 gene expression as a function of chondrocyte passage number and that Sp1 protein synthesis is restored in part by re-differentiation in 3D-alginate culture (Duval et al., 2011). Here, we showed that the decrease of Sp1 induced by serial passaging of chondrocytes is responsible for TßRII reduction. However, we cannot exclude that Sp1 acts indirectly. Sp1 might permit the re-expression of TßRII in dedifferentiated chondrocytes through Sox9, which can be induced by Sp1 (Piera-Velazquez et al., 2007). Sox9 was first identified as a transcription factor which specifies the chondrocyte lineage and is absolutely required for complete cartilage differentiation (Bi et al., 1999). In our study, Sp1 was not able to counteract the decrease of type II collagen (data not shown) whereas it increases aggrecan mRNA level in passaged chondrocytes. Since it was previously established that TßRII gene expression occurred earlier than that of type II collagen gene expression during differentiation of bone marrow-MSC under dynamic compressive loading (Huang et al., 2005), we hypothesize that longer time of Sp1 overexpression could restore the aggrecan or type II collagen expression in dedifferentiated chondrocytes. In order to validate this speculation, it will be necessary to reproduce this experiment using stable transfection system, which could permit to maintain a high level of Sp1 during a long period.

In conclusion, we demonstrated that TßRII expression level is essential to maintain and/or promote chondrocyte phenotype, and established a key-role of Sp1 in this process. These findings may help in understanding the behaviour of aged chondrocytes. In addition, it brings original insights in the process of MSC differentiation into chondrocytes and subsequently permits to envisage innovative strategies to optimize the therapeutic use of chondrocytes or MSC in cartilage disorders.

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- 9 CB conceived, carried out experiments, analyzed data and wrote paper. ED participated to data collection and
- analysis. DO, NG and SL participated to data collection. PG participated to data interpretation. KB conceived
- experiments, analyzed data and was involved in writing the paper. All authors had final approval of the submitted
- 12 and published versions.

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DISCLOSURE

The authors confirm that there are no conflicts of interest.

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FIGURE LEGENDS

- 2 Figure 1: Chondrocyte passages reduces the TGFB responsiveness and decreases TBRII expression.
- 3 A: Primary (P0: about 1-2 population doubling) or passaged (P1 (~2-3 population doubling), P2 (~3-4 population
- 4 doubling) and P3 (~ 4-5 population doubling)) HAC were cultured for 5-6 days before incubation with TGFβ1 (2
- 5 ng/ml) for 4 h. Total RNA was isolated and analyzed by real-time RT-PCR, using PAI-1 and GAPDH primers.
- 6 B-C: Total RNA and proteins from freshly isolated chondrocytes or from chondrocytes dedifferentiated by repeated
- 7 passages were extracted. TBRI and TBRII mRNA levels were defined by real-time RT-PCR.
- 8 D: TBRII protein level was analyzed by Western-blot.

9

1

- 10 Figure 2: Passages up-regulates the R-Smad, Smad3, and down-regulates the I-Smads, Smad7.
- A-D: HAC were cultured as described in figure 1 and Smad2, 3, 4 and 7 mRNA levels were analyzed by real-time
- 12 RT-PCR. Results are shown as histograms of relative expression versus GAPDH signal.

13

- 14 Figure 3: Ectopic expression of TBRII restores TGFB responsiveness in passsaged chondrocytes and increases
- 15 expression of matrix genes.
- 16 A: Primary HAC (P0) or passaged twice (P2) were transfected with TβRII expression vector (pCMV5-TβRII) or
- 17 with the insertless plasmid (5 μ g). After 15 h, the medium was changed for DMEM+2% FCS for 24 h and cells were
- 18 incubated with TGFβ1 addition (2 ng/ml) for 4 h. PAI-1 expression was analyzed by real-time RT-PCR.
- 19 B-C: HAC were cultured and transfected as above. 24 h later, col2A1 and aggrecan mRNA expression was analyzed
- 20 and normalized with GAPDH level (B). Aggrecan and TβRII protein levels were also analysed by Western-blot (C).

21

- 22 Figure 4: TBRII expression increases during chondrocyte differentiation
- A: Primary chondrocytes (P0) or HAC dedifferentiated by 3 passages (P3) were cultured in monolayer or transferred
- into alginate beads for 21 days. Thereafter, TBRII mRNA level was analyzed by real-time RT-PCR.
- 25 B: Human bone marrow-derived stem cells were cultured in alginate beads for different incubation-period and TBRII
- 26 mRNA levels were assayed. Results are shown as relative gene expression versus GAPDH signal.

2728

- Figure 5: Sp1 expression increases during chondrocyte differentiation
- A: Human bone marrow-derived stem cells were cultured in alginate beads for different incubation-period and Sp1
- 30 mRNA levels were assayed. Results are shown as histograms of relative gene expression versus GAPDH signal.
- 31 B: Electrophoretic mobility shift assay (EMSA) reactions were performed using probe corresponding to Sp1
- 32 consensus DNA binding site, incubated with 4 µg nuclear extract from hBM-MSC cultured in alginate for 0h, 72h or
- 33 96h. EMSA reactions were also performed after incubation of nuclear extracts (from hBM cultured 72h in alginate)
- with Sp1, Sp3 and cKrox antibodies.

3536

Figure 6: Sp1 regulates TβRII expression.

- 1 A: Primary HAC, transfected with a Sp1 consensus binding sequence (Sp1 decoy), were incubated in DMEM+2%
- 2 FCS for 24 h. Thereafter, TβRII mRNA versus GAPDH level was determined by RT-PCR.
- 3 B: Subconfluent cultures of chondrocytes were treated for 48 h in the presence or absence of mithramycin (100 and
- 4 150 nM). TβRII expression was analyzed at mRNA level by real-time RT-PCR.
- 5 C: HAC were nucleofected with siRNA oligonucleotides targeted Sp1 or with control siRNA. Thereafter, the
- 6 medium was replaced with DMEM+10% FCS for 24 h. Sp1 and TBRII protein levels were defined by Western-Blot.
- 7 D: Chondrocytes were treated for 48 h in the presence or absence of mithramycin (100 and 150 nM). TβRII and
- 8 aggrecan expression was analyzed at protein level by Western-Blot.

Figure 7: Sp1 regulates chondrocyte phenotype

- 11 A-B: HAC passaged twice (P2) were transfected with Sp1 expression vector (pEVR2-Sp1) or with the insertless
- 12 plasmid (5 μg). After 15 h, the medium was changed for DMEM+2% FCS for 24 h. Total RNA was isolated and
- 13 TBRII and aggrecan mRNA levels were analyzed and normalised with GAPDH level, in comparison to HAC with no
- passage (P0).

15

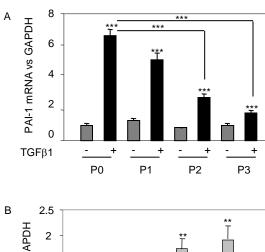
16 Suppl data 1:

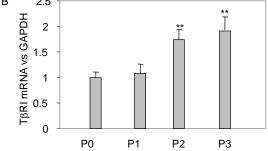
- 17 Primary (P0) or passaged (until P5 ~ 6-7 population doubling) HAC were cultured in monolayer for 5-6 days.
- 18 Thereafter, COL1A1, COL2A1, COL3A1 and aggrecan mRNA levels were analyzed by real-time RT-PCR.

19

20 Suppl data 2:

- A: P3 (~ 4-5 population doubling) chondrocytes were cultured in monolayer or transferred into alginate beads for 21
- days. Thereafter, COL1A1, COL2A1, COL3A1 mRNA levels were analyzed by real-time RT-PCR.
- B: Human bone marrow-derived stem cells were cultured in alginate beads for 7 days and COL2A1 and aggrecan
- 24 mRNA levels were assayed. Results are shown as histograms as relative gene expression versus GAPDH signal.





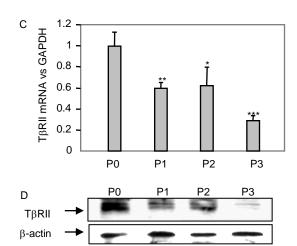


Figure 1:

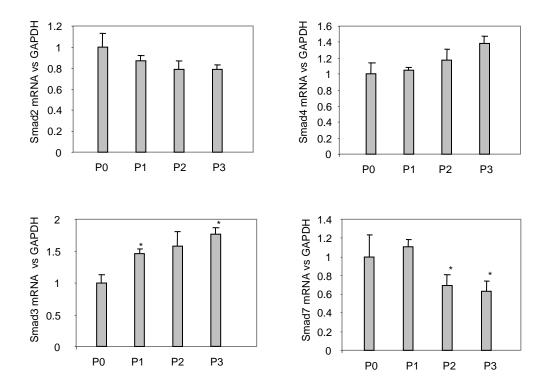


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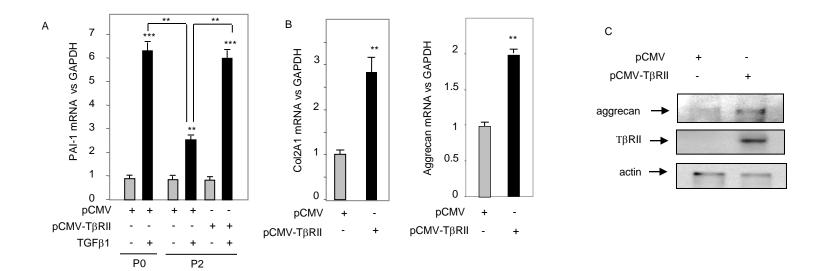
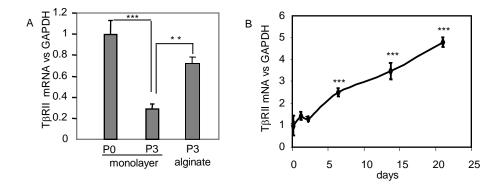


Figure 3:



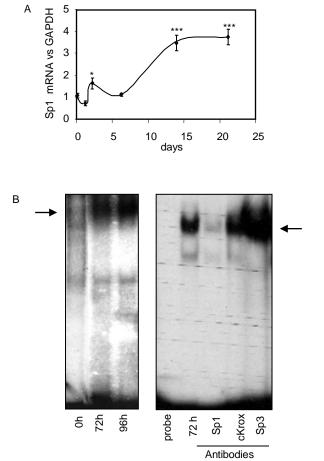


Figure 5

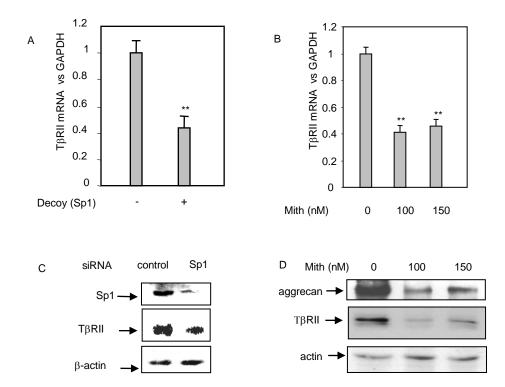
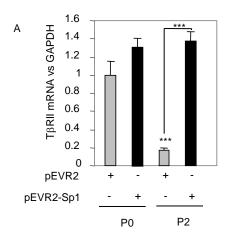


Figure 6



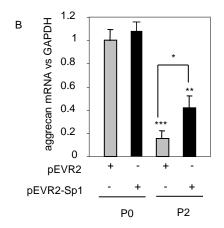


Figure 7

