

Sox5 and Sox6 are required for notochord extracellular matrix sheath formation, notochord cell survival and development of the nucleus pulposus of intervertebral discs

Patrick Smits and Véronique Lefebvre*

Department of Biomedical Engineering and Orthopaedic Research Center, Lerner Research Institute, Cleveland Clinic Foundation, 9500 Euclid Avenue, Cleveland, OH 44195, USA

*Author for correspondence (e-mail: lefebvr@bme.ri.ccf.org)

Accepted 9 December 2002

SUMMARY

The notochord has major roles in vertebral column formation: indirectly by inducing sclerotome cell differentiation; and directly by forming the nucleus pulposus of intervertebral discs. *Sox5* and *Sox6* encode Sry-related HMG box transcription factors that act redundantly to promote chondroblast differentiation in all cartilages of the mouse embryo. We show that *Sox5* and *Sox6* are expressed in the notochord cell lineage and required for notochord late development.

In *Sox5*^{-/-}/*Sox6*^{-/-} embryos, the notochord formed a typical rod-like structure. It fulfilled its inductive functions, as indicated by expression of sonic hedgehog and sclerotome specification. However, the notochord failed to become surrounded with an extracellular matrix sheath. This phenotype was associated with a downregulation of extracellular matrix genes, including the genes for collagen 2, aggrecan and perlecan in both notochord cells and surrounding chondrocytic cells of presumptive inner annuli and vertebral bodies. The mutant notochord then underwent an aberrant, fatal dismantling after sclerotome cell migration. Its cells

became removed first from intervertebral spaces and then from vertebral bodies, and it progressively underwent apoptosis. Meanwhile, the development of inner annuli and vertebral bodies was dramatically impaired. Consequently, the vertebral column of *Sox5*^{-/-}/*Sox6*^{-/-} fetuses consisted of a very deficient cartilage and was devoid of nuclei pulposi. In *Sox5*^{-/-}/*Sox6*^{+/-} and more severely in *Sox5*^{+/-}/*Sox6*^{-/-} embryos, the notochord sheath was thinner, but cells survived. By birth, nuclei pulposi were rudimentary, and its cells poorly swelled and still expressing sonic hedgehog.

Hence, *Sox5* and *Sox6* are required for notochord extracellular matrix sheath formation, notochord cell survival and formation of nuclei pulposi. Through these roles and essential roles in cartilage formation, they are central transcriptional regulators of vertebral column development.

Key words: *Sox5*, *Sox6*, Notochord, Nucleus pulposus, Intervertebral disc, Cartilage, Vertebral column, Mouse

INTRODUCTION

The development of the vertebral column relies on the proper specification and differentiation of several cell lineages. In recent years, important transcription factors, growth factors and signaling molecules have been identified that control such cells as sclerotome cells, chondrocytes and osteoblasts (Christ et al., 2000; de Crombrughe et al., 2001; Karsenty and Wagner, 2002). However, little is known about notochord cells (Fleming et al., 2001).

The notochord is a midline structure of mesodermal origin that forms during gastrulation in chordate embryos (Hogan et al., 1994). By mouse embryonic day 9.5–11.5 (E9.5–E11.5), it consists of a solid rod underlying the neural tube. The cells are small, condensed and enveloped altogether within a sheath of extracellular matrix. This sheath contains many collagens, proteoglycans and glycoproteins, typically found in cartilage,

basement membranes, or mesenchymal and fibrous tissues (Gotz et al., 1995; Hayes et al., 2001).

The notochord has crucial roles in inducing ectoderm, endoderm, and mesoderm derivatives in the early embryo (Cunliffe and Ingham, 1999; Cleaver and Krieg, 2001). It may also have an important mechanical role, and thereby constitute a primitive axial skeleton (Adams et al., 1990). In vertebrate embryos, the notochord has major roles later on during vertebral column formation (Christ et al., 2000; Pourquie et al., 1993). Intervertebral discs are joint-like structures that connect vertebrae. They feature a highly hydrated nucleus pulposus core, surrounded with a cartilaginous inner annulus and a fibrous outer annulus. At E10.5–E11.5, signals from the notochord induce sclerotome cell migration, condensation, and differentiation around the notochord and neural tube. The so-formed perinotochordal tube has a metameric pattern with the alternation of more condensed and less condensed zones.

Between E12.5 and E15.5, the more condensed zones develop into outer and inner annuli and the less condensed zones into vertebrae. The notochord thus has indirect roles in the formation of vertebrae and annuli. Simultaneously, it undergoes profound changes. Notochord cells located in vertebral bodies are removed and probably relocated into intervertebral regions (Aszódi et al., 1998; Rufai et al., 1999). In these latter regions, they proliferate and undergo hypertrophy to form the nuclei pulposi. The notochord thus also directly participates in vertebral column formation.

Several molecules are known to control the early notochord (Cunliffe and Ingham, 1999). For example, the T-box transcription factor brachyury (T) determines notochord cell differentiation and survival (Herrmann and Kispert, 1994), and the secreted factors sonic hedgehog (Shh) and noggin mediate inductive actions (Chiang et al., 1996; McMahon et al., 1998; Teillet et al., 1998). In vitro studies have suggested that the notochord sheath maintains the structural scaffold and internal pressure of the notochord (Carlson and Kenney, 1980; Adams et al., 1990), but its exact roles in vivo have not been determined. The cellular origin and molecular control of this sheath remain unknown. Moreover, the mechanisms that underlie the transformation of the notochord into nuclei pulposi are also largely unknown. Several mouse mutants have an abnormal development of the notochord and vertebral column. These include mice that lack collagen 2 (Li et al., 1995; Aszódi et al., 1998), the paired box transcription factors Pax1 and Pax9 (Wallin et al., 1994; Peters et al., 1999), and the homeobox transcription factor Bapx1 (Tribioli and Lufkin, 1999; Lettice et al., 1999; Akazawa et al., 2000). In these mice, vertebral body cartilages fail to develop properly. The notochord forms normally, but fails to be removed from vertebral bodies and to develop into nuclei pulposi. *Bapx1*, *Pax1* and *Pax9* are expressed in sclerotome cells, but not in notochord cells. It was therefore proposed that the transformation of the notochord into nuclei pulposi requires mechanical pressure from the vertebral body cartilage matrix. As of today, no regulatory factor expressed in notochord cells and controlling cell fate and differentiation beyond early embryonic stages has been identified.

Sox5 and *Sox6* encode two highly identical transcription factors, respectively, L-*Sox5* and *Sox6* (Lefebvre, 2002). These factors feature an Sry-related HMG box domain, which mediates DNA binding, and a coiled-coil domain, which mediates homo- and heterodimerization (Lefebvre et al., 1998). *Sox5* also encode a short protein (*Sox5*) that lacks the N-terminal half of L-*Sox5*, including the dimerization domain. The transcripts for L-*Sox5* and *Sox6* are co-expressed in all cartilages, and expressed in a few other tissues. The transcript for *Sox5* is expressed exclusively in testis. In vitro experiments have suggested that L-*Sox5* and *Sox6* cooperate with *Sox9*, a distant relative and master chondrogenic factor (Bi et al., 1999; Akiyama et al., 2002), in the direct activation of the collagen 2 gene (*Col2a1*) (Lefebvre et al., 1998). The roles of *SOX5* and *SOX6* in humans are unknown.

We recently inactivated *Sox5* and *Sox6* in the mouse and revealed that the two genes have essential, redundant roles in chondrogenesis (Smits et al., 2001). Whereas *Sox5*^{-/-} mice and *Sox6*^{-/-} mice are born with minor cartilage defects, *Sox5*^{-/-}/*Sox6*^{-/-} fetuses develop a generalized and dramatic chondrodysplasia. *Sox5*^{-/-}/*Sox6*^{-/-} prechondrocytes develop

normal precartilaginous condensations, but their differentiation into chondroblasts is delayed and impaired. These cells express cartilage extracellular matrix genes at low or undetectable levels and poorly proliferate. They are unable to persist as a pool of proliferating chondroblasts in epiphyses and to establish cartilage growth plates in metaphyses. Instead, they precociously undergo an aberrant maturation into hypertrophic chondrocytes. By E16.5, when the fetuses die, cartilages are rudimentary and matrix deficient, and are starting to be invaded by bone-forming cells.

We show here that *Sox5* and *Sox6* are expressed in notochord cells and surrounding sclerotome-derived cells, and that *Sox5*^{-/-}/*Sox6*^{-/-} embryos have severe defects in notochord development. *Sox5* and *Sox6* are dispensable for early inductive functions of the notochord, but indispensable for notochord sheath formation, cell survival and development into nuclei pulposi. Hence, *Sox5* and *Sox6* are required to direct the fate and differentiation of notochord cells and chondrocytes, two cell types with major roles in vertebral column formation.

MATERIALS AND METHODS

Mouse mutants and embryo analysis

The generation of *Sox5/Sox6* mutant mice has been described (Smits et al., 2001). Mice were analyzed on the 129/SvEv × C57BL/6 genetic background, and skeletal phenotypes were fully penetrant. *Sox5*^{+/-} and *Sox6*^{+/-} mice were indistinguishable from wild-type littermates in all assays. Therefore, we refer to wild-type and single heterozygous mice as controls. Genotyping, skeletal preparations and histological analyses were performed as described (Smits et al., 2001). Unless otherwise indicated, embryo sections were stained with Alcian Blue and nuclear Fast Red. All experiments were repeated with two or more pairs of control and mutant littermates, and, unless otherwise indicated, figures show data from one representative experiment.

RNA in situ hybridization

RNA in situ hybridization was performed using ³⁵S-labeled antisense probes (Smits et al., 2001). Pictures were taken using a red filter for RNA signals and under blue fluorescence for nuclei stained with the Hoechst 33258 dye. *T* and *Car3* cDNA probes were generated by RT-PCR of ~500 bp of the 3' untranslated sequences. The *Fmod* cDNA probe corresponded to 100 bp of the open reading frame and 200 bp of the 3' untranslated sequence. The *Col3a1* (Metsäranta et al., 1991), *Lamc1* (Laurie et al., 1989), *Pax1* (Deutsch et al., 1988), *Shh* (Echelard et al., 1993) and other cDNA probes (Smits et al., 2001) were as described.

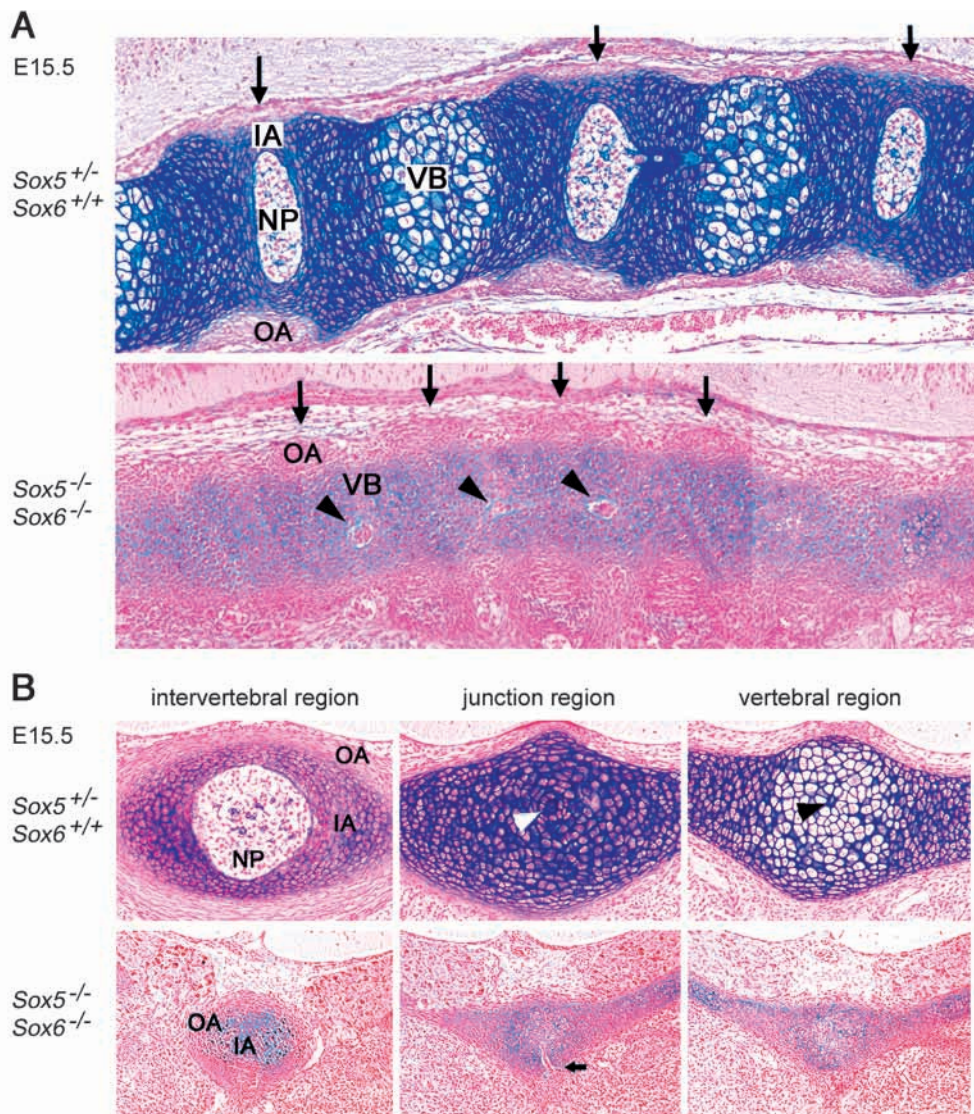
Cell proliferation assay

Pregnant mice were injected with 5-bromo-2'-deoxyuridine (BrdU) (Zymed Laboratories) (10 µl/g of mouse) and sacrificed 2 hours later. Embryos were fixed in 4% paraformaldehyde, dehydrated through a graded ethanol series and embedded in paraffin wax. BrdU-labeled DNA was detected in 7 µm sections using a BrdU immunostaining kit (Zymed laboratories). Sections were counterstained with Hematoxylin.

TUNEL assay

The Alkaline Phosphatase In Situ Cell Death Detection kit (Roche) was used according to the manufacturer's instructions, with the following modifications. Before incubation with the terminal deoxynucleotidyl transferase (TdT), sections were blocked for 20 minutes with 0.3 mg/ml bovine serum albumin in phosphate buffered saline. The TdT enzyme was used at a 1:20 dilution and the converter alkaline phosphatase at a 1:2 dilution.

Fig. 1. Lack of nuclei pulposi in *Sox5*^{-/-}/*Sox6*^{-/-} fetuses. (A) Histological analysis of mid-sagittal sections of the thoracic vertebral column of E15.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} fetuses. In the control (top), intervertebral discs (arrows) feature a nucleus pulposus (NP), a cartilaginous inner annulus (IA) and a mesenchymal outer annulus (OA). Vertebral bodies (VB) exhibit a core of hypertrophic chondrocytes, surrounded with chondroblasts juxtaposed to inner annulus chondroblasts. In the mutant (bottom), the segmentation of the vertebral column is only visible in outer annuli. Nuclei pulposi are lacking. The cartilage matrix of inner annuli and vertebral bodies is deficient (pale Alcian Blue staining), and chondroblasts are all small and round. Groups of condensed cells (arrowheads), remnants of the notochord, are seen in vertebral body regions, but not in intervertebral regions. (B) Histology of cross sections of E15.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} fetuses in the upper lumbar vertebral column. In control intervertebral regions (top), the nucleus pulposus is surrounded with inner and outer annulus. Notochord cells are no longer seen in vertebral bodies or between intervertebral and vertebral regions (arrowheads). The mutant intervertebral and vertebral regions (bottom) show no nucleus pulposus or notochord cells. Residual notochord cells form trains between the center of cartilages and the outer mesenchyme between vertebral and intervertebral regions (arrow).



RESULTS

Sox5^{-/-}/*Sox6*^{-/-} embryos lack nuclei pulposi

Histological analysis of E15.5 *Sox5*^{-/-}/*Sox6*^{-/-} fetuses indicated that in addition to developing severe cartilage deficiencies, these fetuses were also failing to develop intervertebral discs (Fig. 1). In the thoracic region, which, together with the lumbar region, corresponds to the vertebral column segment that is the most developmentally advanced, vertebral bodies could be distinguished from intervertebral discs in control embryos (Fig. 1A). Vertebral bodies contained a core of hypertrophic chondrocytes, flanked with prehypertrophic cells and proliferating chondroblasts. Intervertebral discs contained a nucleus pulposus core surrounded with inner annulus cartilage and outer annulus mesenchyme. In *Sox5*^{-/-}/*Sox6*^{-/-} fetuses, intervertebral areas could be identified only by the presence of outer annulus. There were no nuclei pulposi, and inner annulus cartilage was indistinguishable from vertebral body cartilage. Both cartilages had a deficient extracellular matrix, and vertebral bodies featured no hypertrophic chondrocytes. Interestingly, small groups of condensed cells were present in

vertebral body regions and at the junction with intervertebral regions, but never within intervertebral regions. Their aspect strongly suggested that these cells were remnants of the notochord (see Fig. 4). In the most developed (upper lumbar) region (Fig. 1B), these notochord cells were found only at the junction between vertebral and intervertebral spaces. They were often forming trains heading to the outer mesenchyme. Because a similar phenotype, with abnormal removal of the notochord and absence of nuclei pulposi, had not been described in mouse mutants, we embarked upon a detailed analysis.

Sox5 and *Sox6* are co-expressed in the notochord, nucleus pulposus and cartilage

The expression pattern of *Sox5* and *Sox6* during vertebral column development was analyzed by RNA in situ hybridization of mouse embryo sections (Fig. 2). At E11.5, at the onset of vertebral column formation, *Sox5* and *Sox6* were both expressed in notochord cells and in sclerotome cells surrounding the notochord and neural tube. From E11.5 to E15.5, and until birth (data not shown), *Sox5* and *Sox6*

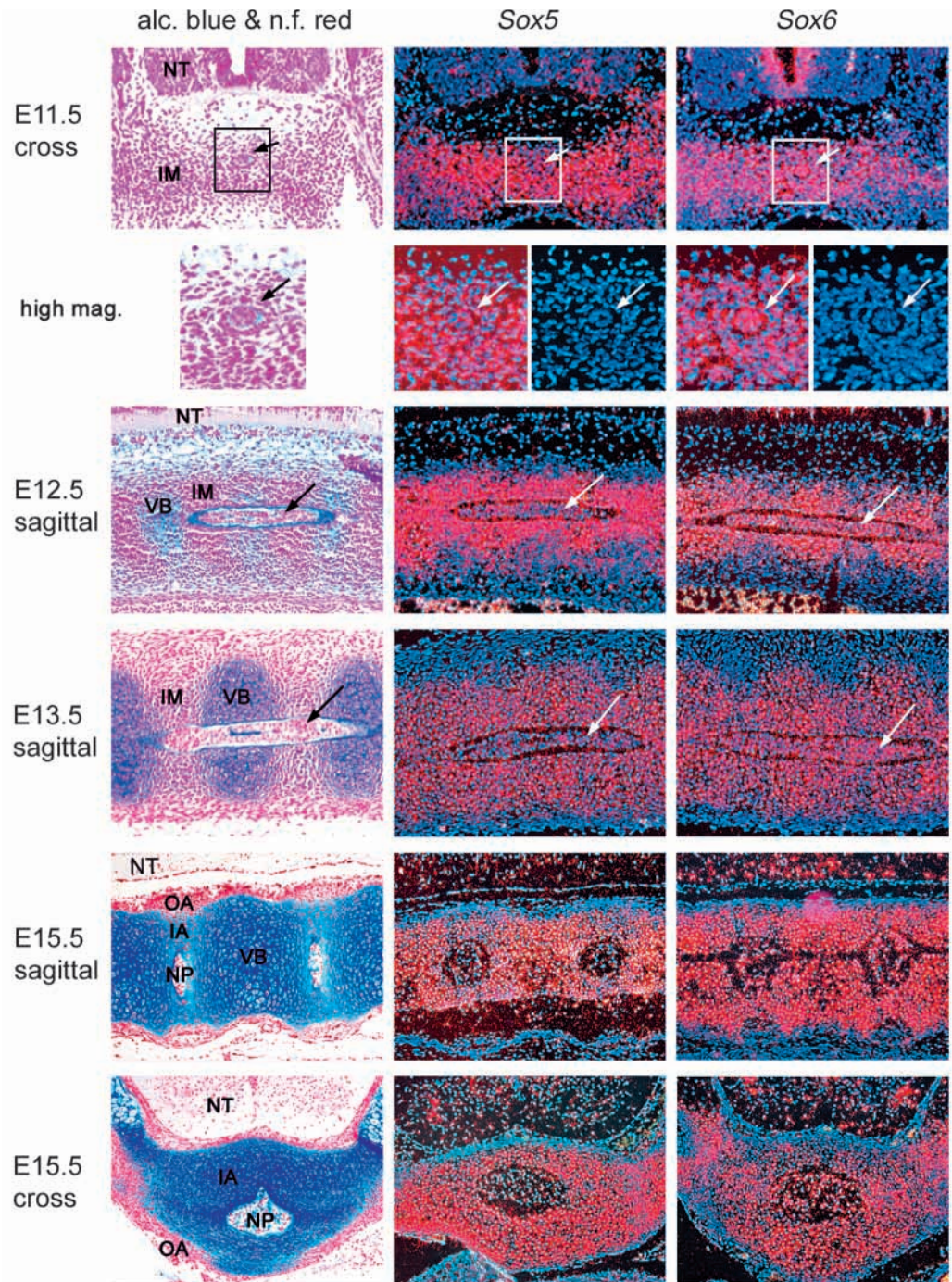


Fig. 2. Co-expression of *Sox5* and *Sox6* in the notochord and nucleus pulposus. Mid-sagittal and cross-sections were made in the lumbar vertebral column of E11.5-E15.5 wild-type embryos. Cross-sections were chosen in intervertebral regions. Each row shows pictures of adjacent sections. Left-hand panels are sections stained with Alcian Blue and nuclear Fast Red to identify the neural tube (NT), notochord (arrows), intervertebral mesenchyme (IM), inner annulus (IA), outer annulus (OA), nucleus pulposus (NP) and vertebral bodies (VB). The other panels show *Sox5* (middle) and *Sox6* (right) RNA in situ hybridization (arrows indicate notochord). The second row displays high-magnification pictures of the boxed areas in the first row. The left-hand pictures for each hybridization were photographed under standard conditions of blue fluorescence (cell nuclei) and dark-field exposure (RNA signals). Fluorescence only was used for the right-hand pictures to visualize the notochord better. At all stages, *Sox5* and *Sox6* are co-expressed in the notochord, nucleus pulposus, prechondrocytes (IM) and chondroblasts (IA, VB). The signal for *Sox5* RNA is slightly weaker in the notochord and nucleus pulposus than in surrounding tissues, whereas the signal for *Sox6* RNA is similar in all these tissues.

remained expressed in notochord cells throughout differentiation into nucleus pulposus cells. They also remained expressed in sclerotome-derived prechondrocytes and chondroblasts in vertebral bodies and inner annuli, but not in outer annuli. At all stages, the signal for *Sox5* RNA was slightly weaker in the notochord and nucleus pulposus than in precartilaginous and cartilage, whereas the signal for *Sox6* RNA was similar in all these tissues.

***Sox5* and *Sox6* are required for notochord sheath formation**

At E11.5, the notochord of both control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos was a rod of condensed cells, with the mutant

notochord generally containing fewer cells (Fig. 3A). Control embryos had started to deposit a notochord sheath, which stained with Alcian Blue, whereas sclerotome-derived cells had not yet accumulated any cartilage matrix. The notochord sheath was totally missing in mutant embryos. By E13.5, the control notochord sheath had become much thicker (Fig. 3B). It was evenly thick in vertebral and intervertebral areas, despite a strong difference in abundance of cartilage matrix between these two areas. In mutant littermates, vertebral bodies had started to deposit some cartilage matrix, but the notochord sheath was still undetectable. *Sox5* and *Sox6* thus control notochord sheath formation.

We tested the RNA levels for notochord sheath components

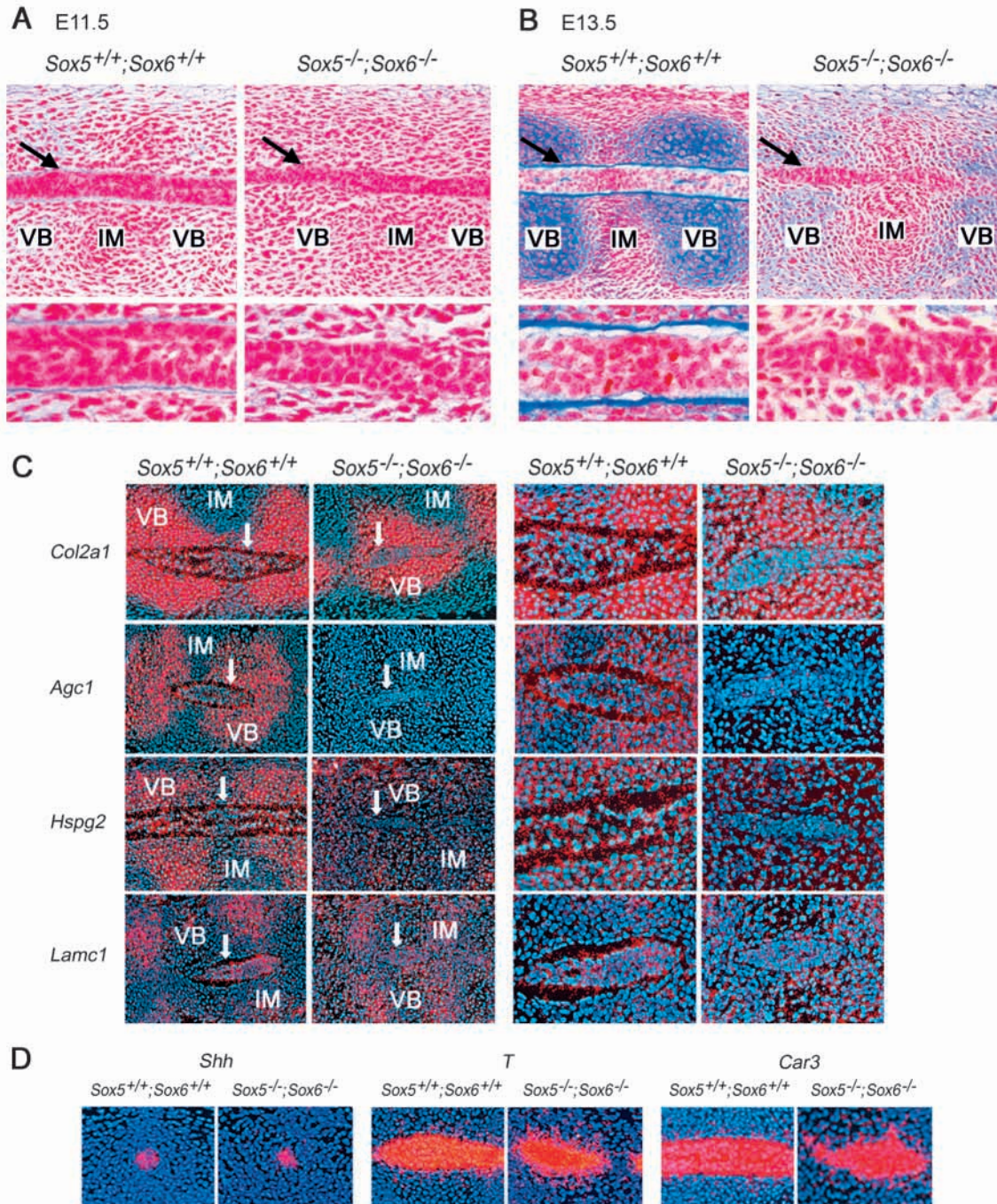


Fig. 3. Lack of notochord sheath in *Sox5*^{-/-}/*Sox6*^{-/-} embryos. (A,B) Histological analysis of mid-sagittal sections of control (left) and *Sox5*^{-/-}/*Sox6*^{-/-} (right) embryos at E11.5 (A) and E13.5 (B). Top panels show the notochord surrounded with vertebral bodies (VB) and intervertebral mesenchyme (IM). Bottom panels show the notochord at higher magnification. The control notochord is surrounded with extracellular matrix (arrows) that stains with Alcian Blue. The mutant notochord has fewer cells and lacks the matrix sheath. (C) In situ hybridization of RNAs for extracellular matrix proteins in mid-sagittal sections of control and *Sox5*^{-/-}/*Sox6*^{-/-} E13.5 embryos. The left and right panel pairs are low- and high-magnification pictures, respectively, of the same areas. Arrows indicate the notochord. *Col2a1* expression is partially downregulated in mutant notochord cells and chondroblasts. *Agc1* and *Hspg2* expression is severely downregulated in both mutant cell types. *Lamc1* expression is not significantly altered or is only slightly downregulated in mutant notochord and intervertebral cells. (D) In situ hybridization of RNAs for notochord-specific non-matrix proteins in control and *Sox5*^{-/-}/*Sox6*^{-/-} E13.5 embryos. *Shh* RNA was tested in cross-sections, and *T* and *Car3* RNA in mid-sagittal sections. The *Sox5*^{-/-}/*Sox6*^{-/-} mutation does not affect expression of these RNAs.

in E11.5 to E14.5 embryos (E13.5, Fig. 3C). As previously described (Smits et al., 2001), *Sox5*^{-/-}/*Sox6*^{-/-} chondroblasts expressed *Col2a1* at a partially reduced level, and the genes for

aggrecan (*Agc1*) and perlecan (*Hspg2*) at virtually undetectable levels. Expression of these genes was similarly reduced in *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells. Control embryos expressed

the gene for laminin $\gamma 1$ (*Lamc1*) in notochord cells and intervertebral mesenchyme, but not in chondroblasts. Mutant notochord cells expressed *Lamc1* at a similar or slightly reduced level. We also tested the expression level of genes for other basement membrane proteins, including collagen 4 (*Col4a1*) and nidogen, and for other mesenchymal and fibrous tissue proteins, including fibronectin, collagen 1 (*Col1a1*) and collagen 3 (*Col3a1*), which were detected by immunolocalization in the notochord sheath of mouse or rat embryos (Gotz et al., 1995; Hayes et al., 2001). However, these genes were not expressed in control notochord cells, and their expression in other cells was not affected by the *Sox5*^{-/-}/*Sox6*^{-/-} mutation (data not shown).

We also tested whether the *Sox5*^{-/-}/*Sox6*^{-/-} mutation affected expression of non-matrix genes in the notochord. We chose genes highly expressed in notochord cells, but not expressed in chondrocytes: *Shh* (Echelard et al., 1993), *T* (Wilkinson et al., 1990) and *Car3* (carbonic anhydrase 3) (Lyons et al., 1991). Their RNA levels were normal in the mutant notochord (Fig. 3D).

Hence, *Sox5* and *Sox6* are not needed for the initial formation of the notochord, but are required for notochord sheath formation. They are required in notochord cells and chondrocytes to express genes for common components of the cartilage matrix and notochord sheath.

***Sox5*^{-/-}/*Sox6*^{-/-} notochord cells are removed from intervertebral spaces and vertebral bodies**

From E13.5 to E15.5, the control notochord developed into nuclei pulposi, and sclerotome cells into vertebral body and inner annulus chondroblasts. Because these transformations started earlier in the thoracolumbar region than in the cervical and caudal regions, all steps could be visualized simultaneously in E14.5 embryos (Fig. 4A). In the distal tail, the notochord was still a continuous rod, intervertebral regions highly condensed, and vertebral bodies starting to accumulate some cartilage matrix. In the proximal tail, the notochord was starting to bulge in intervertebral regions. In the thoracic region and in the E15.5 lumbar region, notochord cells were completely removed from the vertebral bodies, but an acellular notochord sheath was still visible. Nuclei pulposi, vertebral body and inner annulus cartilage were fully developing.

During the same time period, the notochord of *Sox5*^{-/-}/*Sox6*^{-/-} embryos underwent an aberrant dismantlement. First, notochord cells disappeared from presumptive intervertebral areas, but not from vertebral bodies (Fig. 4A, top two panels). The vertebral bodies were underdeveloped, but the intervertebral mesenchyme was normally condensed. Next, notochord cells formed trains that seemed to be moving out of the vertebral column between vertebral and intervertebral spaces (Fig. 4A, bottom two panels). The vertebral bodies featured some cartilage matrix, but the intervertebral mesenchyme was still condensed. The hybridization of *Sox5*^{-/-}/*Sox6*^{-/-} embryo sections with a *Car3* RNA probe convincingly illustrated the disruption of the notochord and diversion of its cells (Fig. 4B). As shown earlier (Fig. 1B), *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells were removed from vertebral bodies by E15.5, thus with a delay of approximately 2 days compared with control cells, and never developed nuclei pulposi.

We also analyzed transverse sections to ascertain that the

notochord interruptions seen in sagittal sections were not due to waviness of the notochord (Fig. 4C,D). At E12.5, the notochord was still visible and expressing *Shh* in the cervical and caudal intervertebral spaces in both control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos. In the thoracolumbar intervertebral spaces, notochord cells were identifiable in control embryos, but not in *Sox5*^{-/-}/*Sox6*^{-/-} embryos. *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells were therefore vanishing from intervertebral regions.

Hence, *Sox5* and *Sox6* are needed to maintain the notochord in intervertebral regions and to remove the notochord from vertebral bodies in a timely manner.

***Sox5* and *Sox6* are needed for notochord cell survival**

We tested whether the reduced numbers of *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells in E11.5 embryos, and the removal of notochord cells from intervertebral spaces in E12.5-E14.5 embryos resulted from abnormal cell proliferation or from cell death. As we have shown previously that *Sox5* and *Sox6* are required to promote chondroblast proliferation, we first tested notochord cell proliferation (Fig. 5A). In both control and mutant embryos, notochord cells were proliferating actively at E11.5, and less actively at E13.5. At both stages, mutant notochord cells were proliferating at a similar or higher rate than control cells. Thus, the loss of mutant notochord cells was not due to a decreased rate in cell proliferation.

Cell death was assessed by the TUNEL assay (Fig. 5B). At E11.5, few apoptotic cells were detected in control notochords, but many dying cells were seen in mutant notochords. Both control and mutant intervertebral mesenchymes featured dying cells, whereas only mutant vertebral mesenchymes featured some. At E13.5, control notochord cells were not dying, whereas *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells were undergoing massive apoptosis in prevertebral and intervertebral regions. Cells were not dying in control cartilage, but a few cells were still dying in *Sox5*^{-/-}/*Sox6*^{-/-} intervertebral mesenchyme. Hence, apoptosis explains the progressive loss of all *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells, but not the selective loss in intervertebral spaces.

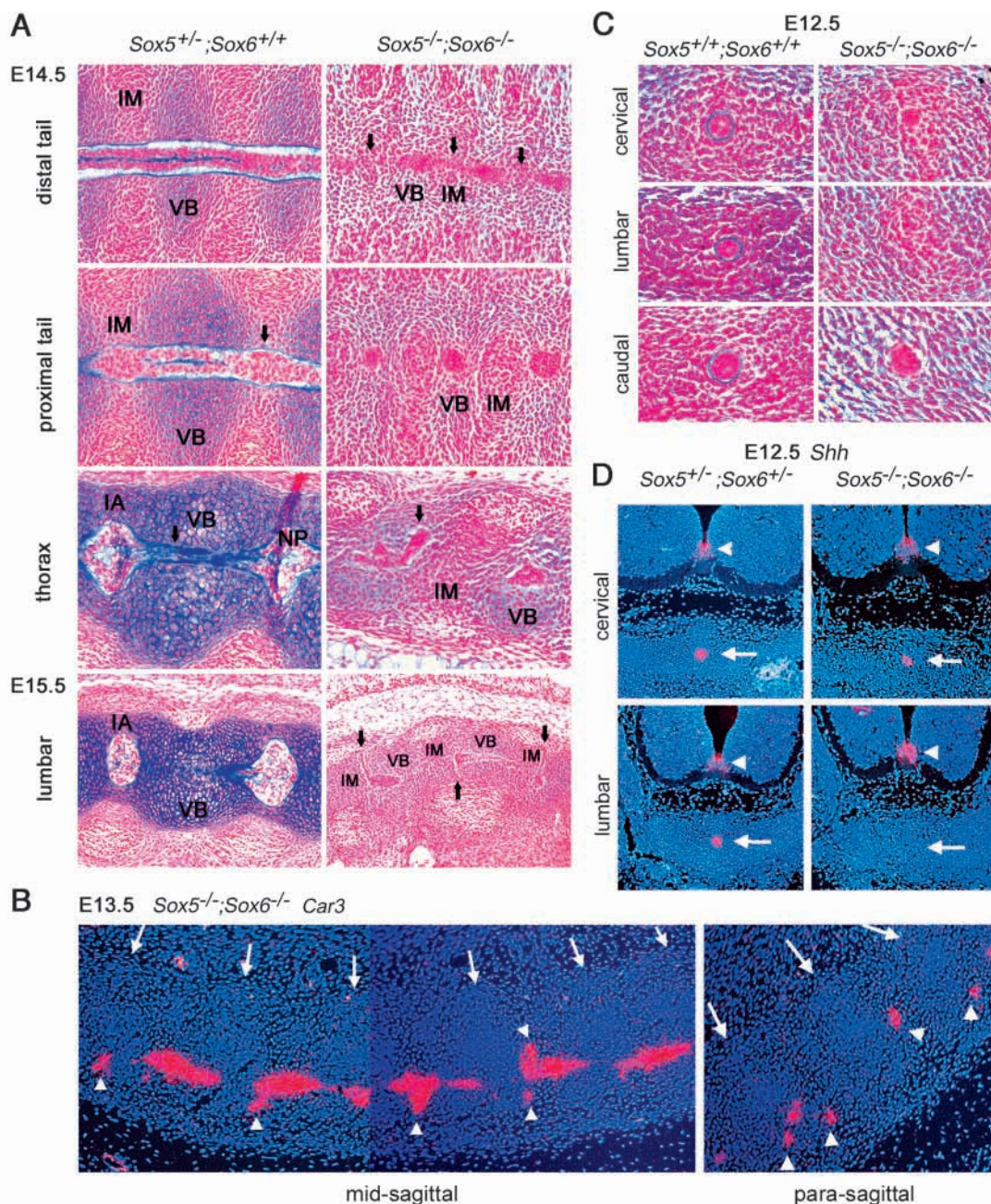
Notochord disintegration occurred in embryos lacking *T* (Herrmann and Kispert, 1994) and *Itm5a* (integrin- $\alpha 5$) (Goh et al., 1997). However, we found that *Itm5a* was expressed in intervertebral cells, but not in notochord and prevertebral cells. Moreover, its expression was not affected by the *Sox5*^{-/-}/*Sox6*^{-/-} mutation (data not shown). As shown above, *T* was expressed at normal levels in *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells (Fig. 3D). Therefore, the cause of apoptosis of *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells was not an altered expression of *T* or *Itm5a*.

***Sox5* and *Sox6* promote inner annulus chondroblast differentiation**

The exclusion of *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells from intervertebral spaces could result either from a notochord defect or from an intervertebral defect. In favor of the latter, histological analysis suggested that the *Sox5*^{-/-}/*Sox6*^{-/-} intervertebral mesenchyme was impaired in its development (Fig. 4A). To test this hypothesis further, we analyzed expression of cell-specific markers.

Pax1 and *Pax9* are indispensable for cell fate specification and differentiation of sclerotome cells. As expected (Deutsch

Fig. 4. Removal of *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells from intervertebral spaces. (A) Histological analysis of mid-sagittal sections of E14.5 and E15.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos. At E14.5, in the distal tail, the control notochord is continuous, but the mutant is interrupted in intervertebral regions (arrows in first row). In the intervertebral regions of the proximal tail, the control notochord is bulging (arrow in second row), whereas mutant notochord cells are missing. In both segments of the tail, control vertebral bodies are starting to accumulate cartilage matrix, but mutant vertebral bodies hardly are. In the E14.5 thoracic region and E15.5 lumbar region (third and fourth rows), control notochord cells have been removed from vertebral bodies, but the notochord sheath is still visible (arrow in left panel in third row). Vertebral body and inner annulus cartilage is developing, and nuclei pulposi expanding. In mutants, vertebral bodies are accumulating a deficient cartilage matrix, and notochord cells are forming trains (arrows in right panels in third and fourth rows) between vertebral and intervertebral regions. (B) In situ hybridization of *Car3* RNA in an E13.5 *Sox5*^{-/-}/*Sox6*^{-/-} embryo. The RNA signal identifies notochord stretches in vertebral bodies, but not in intervertebral regions (higher cell density, arrows). Arrowheads indicate notochord cells found at ectopic positions, i.e. dorsal and ventral (mid-sagittal section) or lateral (para-sagittal section) to the normal axis of the notochord, aside intervertebral regions. (C) Histological analysis of cross-sections of E12.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} presumptive intervertebral discs. The control notochord is intact in all regions. The mutant notochord is still visible in the cervical and caudal regions, but not in the lumbar region. (D) In situ hybridization of *Shh* RNA in cross-sections of E12.5 *Sox5*^{+/-}/*Sox6*^{+/-} and *Sox5*^{-/-}/*Sox6*^{-/-} embryos through intervertebral regions. *Shh* RNA is detected in the floor plate of the neural tube in both embryos (arrowheads). It is also detected in notochord cells in both lumbar and cervical control regions and in the *Sox5*^{-/-}/*Sox6*^{-/-} cervical region (arrows). By contrast, no *Shh* RNA signal is seen in the *Sox5*^{-/-}/*Sox6*^{-/-} lumbar region (arrow).



et al., 1988), *Pax1* was highly expressed in presumptive intervertebral regions in E11.5 wild-type embryos (Fig. 6A). It was correctly expressed in *Sox5*^{-/-}/*Sox6*^{-/-} sclerotome cells, indicating that these cells were correctly specified. From E11.5 to E13.5, control cells underwent chondrogenesis, and *Pax1* expression became confined to narrow zones of intervertebral mesenchyme, to perichondrium and to outer annuli. By E15.5, it remained expressed only in outer annuli, whereas, in mutant

embryos, it was still highly expressed in lower lumbar intervertebral areas.

By E15.5, the genes for the mesenchymal markers collagen 3 (Fig. 6B), collagen 1 and fibronectin (data not shown) were no longer expressed in control and mutant vertebral body chondroblasts and in control inner annulus chondroblasts. In the mutant intervertebral regions, they were being inactivated in the thoracic and higher lumbar regions (data not shown), but

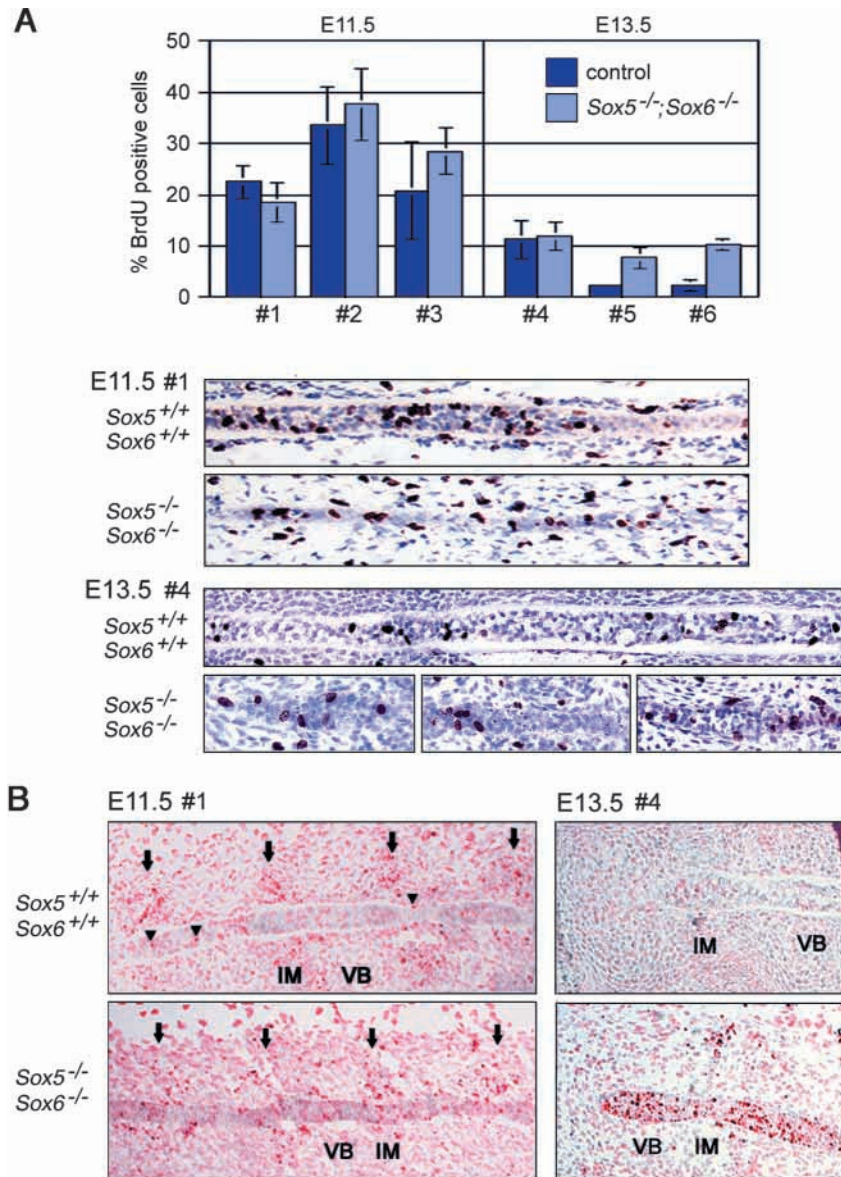


Fig. 5. Normal proliferation, but massive apoptosis of *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells. (A) BrdU labeling of proliferating cells in E11.5 and E13.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos. Three pairs of littermates were tested per age (1-6). For each embryo, notochord cells were counted in five to nine mid-sagittal sections in the thoracolumbar region, totaling 300-1000 cells. The percentage of BrdU-positive cells is presented as the mean±s.d. for all sections. Representative pictures are shown underneath. BrdU-labeled cells are brown. At both stages, mutant notochord cells proliferate at a rate that is similar to or higher than that of control cells. (B) TUNEL assay in the thoracolumbar region of control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos. At E11.5, few cells are dying in the control notochord (red, arrowheads), and many more in the mutant notochord. Cells are also dying in intervertebral mesenchyme (IM) in both embryos (arrows). At E13.5, no cells are dying in the control notochord, but massive apoptosis is occurring in the mutant notochord. No cells are dying in control cartilages, but few cells are dying in mutant intervertebral mesenchyme (arrow). VB, vertebral bodies.

were still expressed in the lower lumbar region (Fig. 6B). The differentiation process of mutant inner annulus cells was thus severely delayed, but not blocked.

The RNAs for general cartilage markers, such as aggrecan (Fig. 6B) and link protein (data not shown), were highly expressed in the cartilages of control embryos, but almost undetectable in mutant intervertebral and vertebral cells. The RNA for fibromodulin, which is expressed at high levels in inner annulus chondroblasts, was undetectable in mutant intervertebral cells (Fig. 6B). This virtual absence of expression of general and inner annulus-specific chondroblast markers revealed that the cells were not only delayed, but also impaired in their differentiation process. These defects may therefore have contributed to remove notochord cells from intervertebral regions.

Sox5 and Sox6 are both needed for notochord development

To assess the relative contribution of *Sox5* and *Sox6* to

important than *Sox5*.

Notochord cells started to bulge in intervertebral spaces from E13.5 in *Sox5*^{+/-}/*Sox6*^{-/-} embryos, as well as in control littermates, but mutant cells started to be removed from vertebral bodies between E14.5 and E16.5 with a short delay relative to control cells (Fig. 7B). This delay coincided with a delay in cartilage development. By E16.5, control nuclei pulposi were fully developing, and its cells were swollen, whereas nuclei pulposi were underdeveloped in *Sox5*^{+/-}/*Sox6*^{-/-} embryos, and its cells were hardly swollen (Fig. 7B,C).

At birth, the nuclei pulposi of control mice appeared as swollen, translucent structures in skeletal preparations (Fig. 7D). Those of *Sox5*^{+/-}/*Sox6*^{+/-}, *Sox5*^{-/-}/*Sox6*^{+/-} and *Sox5*^{-/-}/*Sox6*^{+/-} mice were less expanded in width, but swollen and translucent. The nuclei pulposi of *Sox5*^{+/-}/*Sox6*^{-/-} and *Sox5*^{+/-}/*Sox6*^{-/-} mice were small, often fragmented and opaque. The cells in *Sox5*^{+/-}/*Sox6*^{-/-} nuclei pulposi were still incompletely swollen (Fig. 7E) and still expressing *Shh* (Fig.

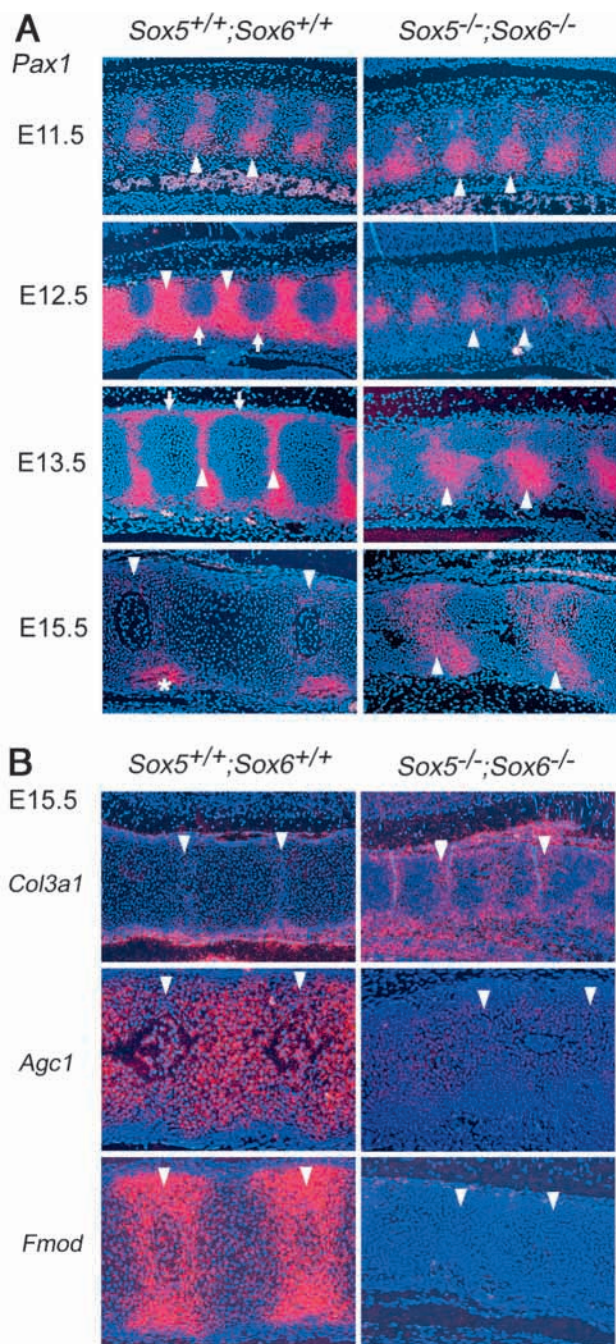


Fig. 6. Impaired development of *Sox5*^{-/-}/*Sox6*^{-/-} intervertebral mesenchyme. (A) In situ hybridization of *Pax1* RNA in mid-sagittal sections of control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos in the lower lumbar vertebral column. At E11.5, *Pax1* is expressed in pre-intervertebral cell condensations in both embryos (arrowheads). At E12.5 and E13.5, *Pax1* is expressed in control embryos in presumptive annuli (arrowheads) and perichondrium (arrows). These expression domains are wider at E12.5 than at E13.5 in the control, but similar in the mutant. At E15.5, *Pax1* expression is restricted to outer annuli in control embryos (asterisk). In *Sox5*^{-/-}/*Sox6*^{-/-} embryos, *Pax1* is still expressed in intervertebral mesenchyme (arrowheads). (B) In situ hybridization of RNAs for extracellular matrix proteins in mid-sagittal sections of the cervical region of E15.5 control and *Sox5*^{-/-}/*Sox6*^{-/-} embryos. Wild-type inner annulus cells (arrowheads) are differentiated chondroblasts: they are no longer expressing *Col3a1*, and are expressing *Agc1* and *Fmod*. *Sox5*^{-/-}/*Sox6*^{-/-} intervertebral cells (arrowheads) are still exhibiting an undifferentiated mesenchymal phenotype: they are expressing *Col3a1* at high levels, and *Agc1* and *Fmod* at low or undetectable levels.

DISCUSSION

Previous studies on the notochord mostly focused on its formation, function and control in early organogenesis. This study has mainly focused on the notochord at later stages, when it directly contributes to vertebral column formation. We have shown that *Sox5* and *Sox6* are not necessary in the early notochord for the formation of a typical rod-like structure, the expression of the important regulatory *T* and *Shh* genes, and for fulfilling inductive functions. However, they are necessary at later stages for notochord sheath formation, cell survival and development of the nuclei pulposi.

Sox5 and *Sox6* are required for notochord sheath formation

Many types of matrix molecules are present in the notochord sheath, but little is known about their function and gene regulation. The short tail (*Ts*) mouse lacks the notochord sheath (Center et al., 1988), but its genetic alteration is unknown.

The first genetic alterations causing abrogation of notochord sheath formation were identified in the laminin β 1 and laminin γ 1 genes in the zebrafish (Parsons et al., 2002). *Lamc1*-null mouse embryos die too early (E5.5) to study the role of laminin γ 1 in notochord sheath formation in this species (Smyth et al., 1999), but crucial functions of laminin 1 have likely been conserved through evolution. We found that notochord cells express *Lamc1* at the time of sheath formation, but *Lamc1* expression is not or hardly affected by the *Sox5*^{-/-}/*Sox6*^{-/-} mutation, ruling out that *Sox5* and *Sox6* mediate notochord sheath formation by controlling *Lamc1* expression.

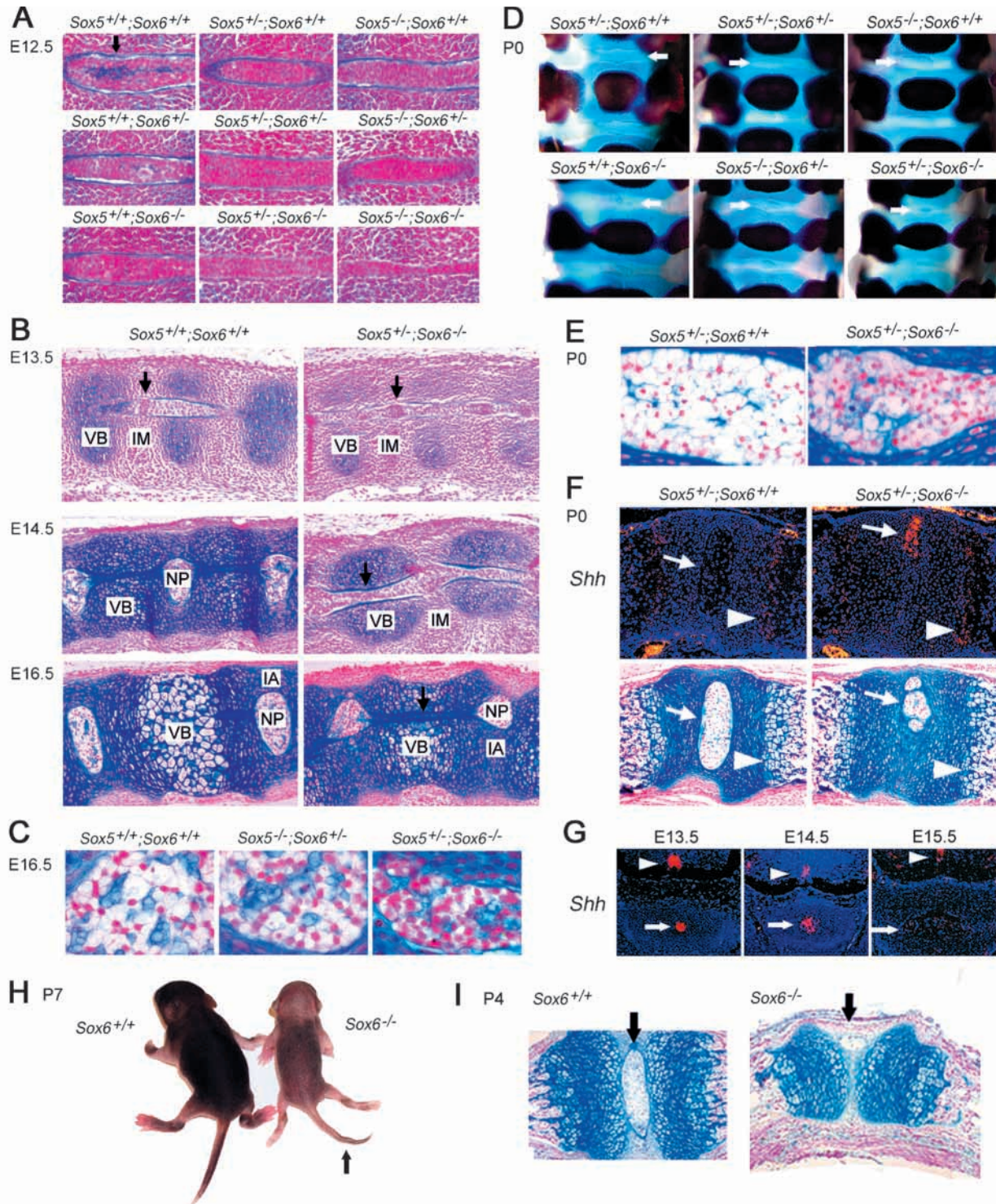
Although the notochord sheath may contain mesenchymal, fibrous tissue and basement membranes components, the genes for such components were not expressed in wild-type notochord cells at the time of sheath formation, and those that were expressed in surrounding tissues were not affected by the *Sox5*^{-/-}/*Sox6*^{-/-} mutation. Hence, *Sox5* and *Sox6* must mediate notochord sheath formation by controlling other genes.

The notochord sheath also contains cartilage matrix components. The expression of genes for these components is partially or severely downregulated in *Sox5*^{-/-}/*Sox6*^{-/-}

7F), whereas wild-type notochord cells had deactivated *Shh* expression by E15.5 upon transformation into nucleus pulposus cells (Fig. 7G).

The few *Sox6*^{-/-} mice that survived postnatally featured a kinked tail in addition to an overall growth delay (Fig. 7H). Histological analysis revealed that the tail kinks were always associated with an eccentric, small nucleus pulposus (Fig. 7I). *Sox5*^{-/-} mice died at birth, but *Sox5*^{+/-}/*Sox6*^{+/-} mice survived postnatally and all had a straight tail (data not shown).

Thus, *Sox5* and *Sox6* have mostly redundant functions in notochord development, but *Sox6* is slightly more important than *Sox5*.



chondroblasts (Smits et al., 2001) and we found here that their expression is similarly affected in *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells. *Col2a1* expression is partially downregulated. However, as *Col2a1*^{-/-} mouse embryos have a thick notochord sheath (Aszódi et al., 1998), the notochord phenotype of *Sox5*^{-/-}/*Sox6*^{-/-} embryos must be caused by the altered expression of more genes. The wild-type notochord sheath intensely stains with Alcian Blue, indicating that it contains sulfated glucosaminoglycans. Accordingly, expression of the genes for the sulfated proteoglycans aggrecan and perlecan is

high in notochord and surrounding chondrocytic cells. It is virtually abrogated in *Sox5*^{-/-}/*Sox6*^{-/-} embryos. Mouse embryos that lack either gene develop a chondrodysplasia similar to that of *Sox5*^{-/-}/*Sox6*^{-/-} embryos, but it has not been described whether they have a similar notochord phenotype (Rittenhouse et al., 1978; Watanabe and Yamada, 1999; Arikawa-Hirasawa et al., 1999; Costell et al., 1999). Nevertheless, it is reasonable to propose that these proteoglycans have crucial roles both in cartilage and the notochord sheath, and that *Sox5* and *Sox6* mediate notochord

Fig. 7. Deficient development of the notochord in *Sox5/Sox6* compound mutants. (A) Histological analysis of notochord sheath formation in E12.5 embryos with all combinations of *Sox5* and *Sox6* wild-type and mutant alleles. The notochord sheath (arrow) becomes thinner as the number of mutant alleles increases. The mutation of *Sox6* is more detrimental than that of *Sox5*. (B) Histological analysis of nucleus pulposus formation in control and *Sox5^{+/-}/Sox6^{-/-}* littermates. Mid-sagittal sections were photographed in the thoracic region. At E13.5, the notochord is starting to bulge in intervertebral spaces in both embryos (arrows). By E14.5, notochord cells have been completely removed from vertebral bodies (VB) in the control embryo, but incompletely removed in the mutant (arrow). The formation of cartilage matrix is delayed in E13.5 and E14.5 mutant embryos. By E16.5, the notochord has completed its transformation into nuclei pulposi in both embryos. The notochord sheath (arrow) is still visible in the mutant vertebral bodies. The mutant nuclei pulposi are smaller, eccentric and often fragmented. IA, inner annulus; IM, intervertebral mesenchyme; NP, nucleus pulposus. (C) High-magnification pictures of nuclei pulposi in E16.5 embryos. Control cells are highly swollen, as indicated by a large volume of cytoplasm (white) per cell nucleus (red). *Sox5^{+/-}/Sox6^{+/-}* cells are incompletely swollen and *Sox5^{+/-}/Sox6^{-/-}* cells hardly swollen. (D) Skeletal preparations of the vertebral column of newborn mice (P0) with various combinations of *Sox5* and *Sox6* null alleles. Non-mineralized cartilages and nuclei pulposi are stained with Alcian Blue, and mineralized cartilages and bones with Alizarin Red. Pictures show the L1 lumbar vertebra and flanking intervertebral discs. The nuclei pulposi (arrows) of *Sox5^{+/-}/Sox6^{+/-}* and *Sox5^{-/-}/Sox6^{+/-}* and *Sox5^{-/-}/Sox6^{+/-}* mice are less expanded in width than those of the control (*Sox5^{+/-}/Sox6^{+/-}*) mouse. The nuclei pulposi of *Sox5^{+/-}/Sox6^{-/-}* and *Sox5^{+/-}/Sox6^{-/-}* mice are severely reduced in both width and height, and opaque instead of translucent. (E) Histological analysis of nuclei pulposi in control (*Sox5^{+/-}/Sox6^{+/-}*) and *Sox5^{+/-}/Sox6^{-/-}* newborn littermates. Mutant cells are less swollen than control cells. (F) In situ hybridization of *Shh* RNA (top) and histological analysis (bottom) of sagittal sections of control (*Sox5^{+/-}/Sox6^{+/-}*) and *Sox5^{+/-}/Sox6^{-/-}* newborn littermates. Pictures show a lumbar intervertebral disc and segments of flanking vertebrae. Nucleus pulposus cells (arrows) are still expressing *Shh* in the mutant, but not in the control. Hypertrophic chondrocytes (arrowheads) are expressing *Shh* in both mice. (G) In situ hybridization of *Shh* RNA in cross-sections of wild-type embryos. Pictures show a presumptive intervertebral disc region and the ventral part of the neural tube. *Shh* is expressed in E13.5 and E14.5 notochord cells (arrows), but no longer in E15.5 nucleus pulposus cells (arrow). At all three stages, *Shh* is expressed in the neural tube floor plate (arrowheads). (H) Picture of 1-week-old (P7) control and *Sox6^{-/-}* littermates. The *Sox6^{-/-}* mouse is developmentally delayed and exhibits a kinked tail (arrow). (I) Histological analysis of longitudinal sections through the tail of 4-day-old (P4) control and *Sox6^{-/-}* mice. Control nuclei pulposi (arrows) occupy the core region of intervertebral spaces. Mutant nuclei pulposi are small and located eccentrically, where the tail forms kinks.

sheath formation by controlling these genes and possibly other genes in notochord and surrounding cells.

The cellular origin of the notochord sheath is unknown. Transplantation experiments in zebrafish embryos have shown that a normal notochord sheath forms when laminin 1-deficient notochord precursor cells are transplanted into a wild-type embryo, and vice versa (i.e. when wild-type notochord precursor cells are transplanted into a mutant embryo) (Parsons et al., 2002). Hence, both notochord cells and surrounding cells may contribute notochord sheath components. Other

observations, however, indicate that notochord cells must have a preponderant role in notochord sheath formation. First, we have shown that the notochord sheath starts to form in wild-type embryos before any cartilage matrix in surrounding tissues, and later, accumulates evenly in cartilaginous vertebral bodies and mesenchymal intervertebral areas. Second, embryos that lack *Bapx1* (Akazawa et al., 2000), *Pax1* and *Pax9* (Peters et al., 1999) fail to form any cartilage matrix around the notochord, but appear to form a normal notochord sheath. Finally, we have shown that *Sox6* is expressed, relative to *Sox5*, at a slightly higher level in notochord cells than in surrounding cells, and that the inactivation of *Sox6* is more detrimental for notochord sheath formation than the inactivation of *Sox5*. Based on these data, we believe that *Sox5* and *Sox6* mediate notochord sheath formation by controlling extracellular matrix genes mainly in notochord cells.

Sox5 and Sox6 are required for notochord cell survival

Notochord cell death occurs in *Sox5^{-/-}/Sox6^{-/-}* embryos later than in *T^{-/-}* (Herrmann and Kispert, 1994), *Itm5a^{-/-}* (Goh et al., 1997) and Danforth's short tail embryos (Asakura and Tapscott, 1998). However, *T* is expressed at a normal level in *Sox5^{-/-}/Sox6^{-/-}* notochord cells. *Itm5a* is not expressed in wild-type notochord cells, and not affected by the *Sox5^{-/-}/Sox6^{-/-}* mutation in surrounding cells. Therefore, *T* and *Itm5a* are not involved in the death of *Sox5^{-/-}/Sox6^{-/-}* notochord cells. The gene altered in the Danforth's short tail mouse is unknown.

Increased apoptosis of notochord cells occurs in mouse embryos with a deletion of *Jun* in the notochord and sclerotome (Behrens et al., 2003). It occurs at the same stage of development as in *Sox5^{-/-}/Sox6^{-/-}* embryos. However, *Jun*-deficient mice form a normal notochord sheath, and most notochord cells survive to form nuclei pulposi. *Jun* must therefore act independently of *Sox5* and *Sox6* in the notochord.

The phenotypic comparison of *Sox5^{-/-}/Sox6^{-/-}* embryos and laminin 1-deficient zebrafish embryos strongly suggests that *Sox5* and *Sox6* mediate notochord cell survival essentially through their roles in notochord sheath formation. The primary defect of both mutants is the lack of notochord sheath, and it is directly followed by notochord cell apoptosis upon sclerotome cell condensation. The sheath may act on notochord cells in one or several ways. It may provide vital cell-matrix interactions, store essential growth factors or establish a barrier against death signals or mechanical pressure from surrounding tissues.

Sox5 and Sox6 are required for proper relocation of notochord cells

Sox5^{-/-}/Sox6^{-/-} notochord cells are aberrantly removed from intervertebral areas, while their rates of apoptosis and proliferation are not different between vertebral and intervertebral regions, and mutant sclerotome cells are normally condensed and specified. We therefore propose that *Sox5^{-/-}/Sox6^{-/-}* notochord cells may be removed from intervertebral regions because of their specific lack of notochord sheath.

Sox5^{-/-}/Sox6^{-/-} embryos remove notochord cells from vertebral bodies later than wild-type embryos. Previous studies have suggested that the wild-type notochord is removed upon swelling pressure from the vertebral body cartilage matrix. The

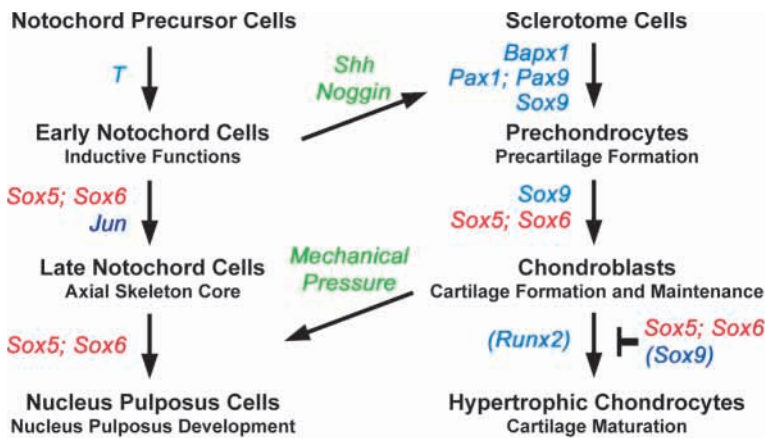


Fig. 8. *Sox5* and *Sox6* control essential steps in the notochord and chondrocyte differentiation pathways. Chondrocytes and notochord cells progress through four major differentiation steps during development. Major transcription factors known to control one or several of these steps are indicated. Biochemical (*Shh* and *noggin*) and physical (mechanical pressure) interactions occur between the two lineages at several steps. See the end of Discussion for details.

notochord persists in the cartilage-deficient *Col2a1*^{-/-} vertebral bodies (Aszódi et al., 1998), and in the undifferentiated *Bapx1*^{-/-} vertebral bodies (Lettice et al., 1999; Tribioli and Lufkin, 1999; Akazawa et al., 2000). The delay in removing *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells from vertebral bodies must result from the delay in cartilage development. It may also be facilitated by the lack of notochord sheath and apoptosis of notochord cells.

Wild-type notochord cells are believed to take refuge in intervertebral areas upon removal from vertebral bodies, but *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells remain excluded from these areas and relocate between vertebral bodies and intervertebral areas. The repositioning of the cells, although ectopic, supports the relocation model proposed for wild-type notochord cells, and indicates that *Sox5* and *Sox6* are needed to maintain and relocate notochord cells in intervertebral areas.

***Sox5* and *Sox6* promote nucleus pulposus cell differentiation**

Little is known about the genetic program of nucleus pulposus cells. They continue to express *Sox5*, *Sox6* and *T*, but not *Shh*. They undergo hypertrophy, but do not activate the hypertrophic chondrocyte marker *Col10a1* (data not shown). Their genetic program thus only partially overlaps with that of chondrocytes. *Sox5*^{-/-}/*Sox6*^{-/-} notochord cells die before developing nuclei pulposi, but the nuclei pulposi of *Sox5*^{+/-}/*Sox6*^{-/-} fetuses are severely impaired in their development. They remain rudimentary and their cells swell poorly and maintain *Shh* expression. *Sox5* and *Sox6* are the first genes for transcription factors shown to be expressed in these cells and needed for their differentiation.

Hypertrophic chondrocyte differentiation is also impaired in *Sox5*^{-/-}/*Sox6*^{-/-} embryos (Smits et al., 2001). Chondrocytes activate *Cbfa1* and *VEGF*, but weakly express *Col10a1*, maintain *Sox9* expression and poorly enlarge. These defects must be secondary consequences of the *Sox5*^{-/-}/*Sox6*^{-/-} mutation because *Sox5* and *Sox6* are no longer expressed in hypertrophic chondrocytes. Similarly, some of the defects of *Sox5*^{+/-}/*Sox6*^{-/-} nuclei pulposi may be secondary to defects in the earlier notochord, or in surrounding inner annuli.

***Sox5* and *Sox6* promote inner annulus cell differentiation**

The vertebral column of *Sox5*^{-/-}/*Sox6*^{-/-} fetuses is mostly

unsegmented by E15.5, even though the earlier vertebral column is properly patterned, with sclerotome cells condensing correctly and expressing *Pax1*. As previously described, the development of vertebral body cartilage is delayed and impaired (Smits et al., 2001) such that, by E15.5, it features no hypertrophic chondrocytes yet, and no growth plates. We have shown

here that the differentiation process of inner annulus cartilage is similarly delayed and impaired, such that both cartilages were indistinguishable by E15.5. Moreover, mutant inner annulus cells do not activate *Fmod*, an abundant marker of both inner annulus and periarticular chondroblasts. We previously showed that periarticular chondroblasts in *Sox5*^{-/-}/*Sox6*^{-/-} limb cartilages also fail to activate *Fmod* (Smits et al., 2001). Therefore, the impaired differentiation of *Sox5*^{-/-}/*Sox6*^{-/-} inner annulus cells is a cell-autonomous defect, not a consequence of notochord defects.

***Sox5* and *Sox6* have redundant roles in notochord development**

Sox5^{-/-} and *Sox6*^{-/-} embryos have limited notochord defects compared with *Sox5*^{-/-}/*Sox6*^{-/-} embryos, indicating that *Sox5* and *Sox6* have mostly redundant functions in notochord development, as in chondrogenesis (Smits et al., 2001). Nevertheless, the analysis of compound mutants has revealed that *Sox6* is slightly more important than *Sox5* for both notochord sheath formation and nucleus pulposus development. This functional difference between the two genes probably reflects the fact that *Sox6* is expressed, relatively to *Sox5*, at a slightly higher level in notochord cells than in surrounding chondrocytes. This conclusion is consistent with the observation that L-*Sox5* and *Sox6* have indistinguishable DNA binding and transactivation activities in vitro (Lefebvre et al., 1998).

***Sox5* and *Sox6* have critical roles in vertebral column formation**

This study and our previous study (Smits et al., 2001) have identified critical roles for *Sox5* and *Sox6* in the transcriptional control of notochord cells and chondrocytes (Fig. 8).

The differentiation of early notochord cells does not require *Sox5* and *Sox6*, but requires *T*. *Shh* and *noggin*, secreted by notochord cells, contribute to induce sclerotome cell differentiation into prechondrocytes. *Bapx1*, *Pax1*, *Pax9* and *Sox9* act downstream of *Shh* at this step in the vertebral column.

Both pathways crucially rely on *Sox5* and *Sox6* at the next step of cell differentiation. *Sox5* and *Sox6* promote expression of extracellular matrix genes, and are thereby required for notochord sheath and cartilage matrix formation. *Sox5* and *Sox6* are required for proper proliferation of chondroblasts, but

not notochord cells. By contrast, they are needed for the survival of notochord cells, but not chondroblasts. *Jun* also is needed for notochord cell survival, and *Sox9* for chondroblast differentiation.

Later, mechanical pressure from cartilage likely induces notochord cell relocation from vertebral bodies into intervertebral areas. *Sox5* and *Sox6* promote notochord cell differentiation into nucleus pulposus cells. By contrast, *Sox5* and *Sox6* prevent chondroblast differentiation into hypertrophic chondrocytes. *Runx2* may promote hypertrophic chondrocyte maturation in vertebrae (Inada et al., 1999; Kim et al., 1999) and *Sox9* may prevent chondrocyte hypertrophy (Bi et al., 2001).

In conclusion, *Sox5* and *Sox6* have essential functions in the development of cartilage, the notochord and nuclei pulposi. They are thereby critical transcriptional regulators of vertebral column scaffold formation.

We thank B. de Crombrughe, U. Deutsch, M. Hööke, A. McMahon and Y. Yamada for in situ hybridization probes; and V. Hascall, C. Kassuba, C. McDevitt and M. Warman for critical suggestions on the manuscript. P. S. was supported by a Postdoctoral Fellowship from the Arthritis Foundation. This work was supported by NIH grant AR46249 to V. L. and by the Lerner Research Institute of The Cleveland Clinic Foundation.

REFERENCES

- Adams, D. S., Keller, R. and Koehl, M. A. (1990). The mechanics of notochord elongation, straightening and stiffening in the embryo of *Xenopus laevis*. *Development* **110**, 115-130.
- Akazawa, H., Komuro, I., Sugitani, Y., Yazaki, Y., Nagai, R. and Noda, T. (2000). Targeted disruption of the homeobox transcription factor *Bapx1* results in lethal skeletal dysplasia with asplenia and gastroduodenal malformation. *Genes Cells* **5**, 499-513.
- Akiyama, H., Chaboissier, M.-C., Martin, J. E., Schedl, A. and de Crombrughe, B. (2002). The transcription factor *Sox9* has essential roles in successive steps of the chondrocyte differentiation pathway and is required for expression of *Sox5* and *Sox6*. *Genes Dev.* **16**, 2813-2828.
- Arikawa-Hirasawa, E., Watanabe, H., Takami, H., Hassell, J. R. and Yamada, Y. (1999). Perlecan is essential for cartilage and cephalic development. *Nat. Genet.* **23**, 354-358.
- Asakura, A. and Tapscott, S. J. (1998). Apoptosis and epaxial myotome in Danforth's short-tail (Sd) mice in somites that form following notochord degeneration. *Dev. Biol.* **203**, 276-289.
- Aszodi, A., Chan, D., Hunziker, E., Bateman, J. F. and Fässler, R. (1998). Collagen II is essential for the removal of the notochord and the formation of intervertebral discs. *J. Cell Biol.* **143**, 1399-1412.
- Behrens, A., Haigh, J., Mechta-Grigoriou, F., Nagy, A., Yaniv, M. and Wagner, E. F. (2003). Impaired intervertebral disc formation in the absence of *Jun*. *Development* **130**, 103-109.
- Bi, W., Deng, J. M., Zhang, Z., Behringer, R. R. and de Crombrughe, B. (1999). *Sox9* is required for cartilage formation. *Nat. Genet.* **22**, 85-89.
- Bi, W., Huang, W., Whitworth, D. J., Deng, J. M., Zhang, Z., Behringer, R. R. and de Crombrughe, B. (2001). Haploinsufficiency of *Sox9* results in defective cartilage primordia and premature skeletal mineralization. *Proc. Natl. Acad. Sci. USA* **98**, 6698-6703.
- Carlson, E. C. and Kenney, M. C. (1980). Surface ultrastructure of the isolated notochord in vitro: the effect of the perinotochord sheath. *Anat. Rec.* **197**, 257-276.
- Center, E. M., Marcus, N. M. and Wilson, D. B. (1988). Abnormal development of the notochord and perinotochordal sheath in duplicatas posterior, patch, and tail-short mice. *Histol. Histopathol.* **3**, 405-412.
- Chiang, C., Litingtung, Y., Lee, E., Young, K. E., Corden, J. L., Westphal, H. and Beachy, P. A. (1996). Cyclopia and defective axial patterning in mice lacking Sonic hedgehog gene function. *Nature* **383**, 407-413.
- Christ, B., Huang, R. and Wilting, J. (2000). The development of the avian vertebral column. *Anat. Embryol.* **202**, 179-194.
- Cleaver, O. and Krieg, P. A. (2001). Notochord patterning of the endoderm. *Dev. Biol.* **234**, 1-12.
- Costell, M., Gustafsson, E., Aszodi, A., Mörgelin, M., Bloch, W., Hunziker, E., Addicks, K., Timpl, R. and Fässler, R. (1999). Perlecan maintains the integrity of cartilage and some basement membranes. *J. Cell Biol.* **147**, 1109-1122.
- Cunliffe, V. T. and Ingham, P. W. (1999). Switching on the notochord. *Genes Dev.* **13**, 1643-1646.
- de Crombrughe, B., Lefebvre, V. and Nakashima, K. (2001). Regulatory mechanisms in the pathways of cartilage and bone formation. *Curr. Opin. Cell Biol.* **13**, 721-727.
- Deutsch, U., Dressler, G. R. and Gruss, P. (1988). Pax 1, a member of a paired box homologous murine gene family, is expressed in segmented structures during development. *Cell* **53**, 617-625.
- Echelard, Y., Epstein, D. J., St-Jacques, B., Shen, L., Mohler, J., McMahon, J. A. and McMahon, A. P. (1993). Sonic hedgehog, a member of a family of putative signaling molecules, is implicated in the regulation of CNS polarity. *Cell* **75**, 1417-1430.
- Fleming, A., Keynes, R. J. and Tannahill, D. (2001). The role of the notochord in vertebral column formation. *J. Anat.* **199**, 177-180.
- Goh, K. L., Yang, J. T. and Hynes, R. O. (1997). Mesodermal defects and cranial neural crest apoptosis in alpha5 integrin-null embryos. *Development* **124**, 4309-4319.
- Gotz, W., Osmer, R. and Herken, R. (1995). Localisation of extracellular matrix components in the embryonic human notochord and axial mesenchyme. *J. Anat.* **186**, 111-121.
- Hayes, A. J., Benjamin, M. and Ralphs, J. R. (2001). Extracellular matrix in the development of the intervertebral disc. *Matrix Biol.* **20**, 107-121.
- Herrmann, B. G. and Kispert, A. (1994). The T genes in embryogenesis. *Trends Genet.* **10**, 280-286.
- Hogan, B., Beddington, R., Costantini, F., Lacy, F. (1994). Early mouse development. In *Manipulating the Mouse Embryo. A Laboratory Manual*, 2nd edn, pp. 63-81. Cold Spring Harbor, NY: Cold Spring Harbor Laboratory Press.
- Inada, M., Yasui, T., Nomura, S., Miyake, S., Deguchi, K., Himeno, M., Sato, M., Yamagiwa, H., Kimura, T., Yasui, N. et al. (1999). Maturational disturbance of chondrocytes in *Cbfa1*-deficient mice. *Dev. Dyn.* **214**, 279-290.
- Karsenty, G. and Wagner, E. F. (2002). Reaching a genetic and molecular understanding of skeletal development. *Dev. Cell* **2**, 389-406.
- Kim, I. S., Otto, F., Zabel, B. and Mundlos, S. (1999). Regulation of chondrocyte differentiation by *Cbfa1*. *Mech. Dev.* **80**, 159-170.
- Laurie, G. W., Horikoshi, S., Killen, P. D., Segui-Real, B. and Yamada, Y. (1989). In situ hybridization reveals temporal and spatial changes in cellular expression of mRNA for a laminin receptor, laminin, and basement membrane (type IV) collagen in the developing kidney. *J. Cell Biol.* **109**, 1351-1362.
- Lefebvre, V., Li, P. and de Crombrughe, B. (1998). A new long form of *Sox5* (*L-Sox5*), *Sox6* and *Sox9* are co-expressed in chondrogenesis and cooperatively activate the type II collagen gene. *EMBO J.* **17**, 5718-5733.
- Lefebvre, V. (2002). Toward understanding the functions of the two highly related *Sox5* and *Sox6* genes. *J. Bone Miner. Metab.* **20**, 121-130.
- Lettice, L. A., Purdie, L. A., Carlson, G. J., Kilanowski, F., Dorin, J. and Hill, R. E. (1999). The mouse bagpipe gene controls development of axial skeleton, skull, and spleen. *Proc. Natl. Acad. Sci. USA* **96**, 9695-9700.
- Li, S. W., Prockop, D. J., Helminen, H., Fassler, R., Lapvetelainen, T., Kiraly, K., Peltarri, A., Arokoski, J., Lui, H., Arita, M. and Khillan, J. S. (1995). Transgenic mice with targeted inactivation of the *Col2* alpha 1 gene for collagen II develop a skeleton with membranous and periosteal bone but no endochondral bone. *Genes Dev.* **9**, 2821-2830.
- Lyons, G. E., Buckingham, M. E., Tweedie, S. O. and Edwards, Y. H. (1991). Carbonic anhydrase III, an early mesodermal marker, is expressed in embryonic mouse skeletal muscle and notochord. *Development* **111**, 233-244.
- McMahon, J. A., Takada, S., Zimmerman, L. B., Fan, C.-M., Harland, R. M. and McMahon, A. P. (1998). Noggin-mediated antagonism of BMP signaling is required for growth and patterning of the neural tube and somite. *Genes Dev.* **12**, 1438-1452.
- Metsäranta, M., Toman, D., de Crombrughe, B. and Vuorio, E. (1991). Specific hybridization probes for mouse type I, II, III and IX collagen RNAs. *Biochim. Biophys. Acta.* **1089**, 241-243.
- Parsons, M. J., Pollard, S. M., Saude, L., Feldman, B., Coutinho, P., Hirst, E. M. A. and Stemple, D. L. (2002). Zebrafish mutants identify essential roles for laminins in notochord formation. *Development* **129**, 3137-3146.

- Peters, H., Wilm, B., Sakai, N., Imai, K., Maas, R. and Balling, R.** (1999). Pax1 and Pax9 synergistically regulate vertebral column development. *Development* **126**, 5399-5408.
- Pourquié, O., Coltey, M., Teillet, M. A., Ordahl, C. and le Douarin, N. M.** (1993). Control of dorsoventral patterning of somitic derivatives by notochord and floor plate. *Proc. Natl. Acad. Sci. USA* **90**, 5242-5246.
- Rittenhouse, E., Dunn, L. C., Cookingham, J., Calo, C., Spiegelman, M., Dooher, G. B. and Bennett, D.** (1978). Cartilage matrix deficiency (cmd): a new autosomal recessive lethal mutation in the mouse. *J. Embryol. Exp. Morphol.* **43**, 71-84.
- Rufai, A., Benjamin, M. and Ralphs, J. R.** (1995). The development of fibrocartilage in the rat intervertebral disc. *Anat. Embryol.* **192**, 5242-5246.
- Smits, P., Li, P., Mandel, J., Zhang, Z., Deng, J. M., Behringer, R. R., de Crombrughe, B. and Lefebvre, V.** (2001). The transcription factors L-Sox5 and Sox6 are essential for cartilage formation. *Dev. Cell* **1**, 277-290.
- Smyth, N., Vatanserver, H. S., Murray, P., Meyer, M., Frie, C., Paulsson, M. and Edgar, D.** (1999). Absence of basement membranes after targeting the *LAMC1* gene results in embryonic lethality due to failure of endoderm differentiation. *J. Cell Biol.* **144**, 151-160.
- Teillet, M., Watanabe, Y., Jeffs, P., Duprez, D., Lapointe, F. and le Douarin, N. M.** (1998). Sonic hedgehog is required for survival of both myogenic and chondrogenic somitic lineages. *Development* **125**, 2019-2030.
- Tribioli, C. and Lufkin, T.** (1999). The murine Bapx1 homeobox gene plays a critical role in embryonic development of the axial skeleton and spleen. *Development* **126**, 5699-5711.
- Wallin, J., Wilting, J., Koseki, H., Fritsch, R., Christ, B. and Balling, R.** (1994). The role of Pax1 in axial skeleton development. *Development* **120**, 1109-1121.
- Watanabe, H. and Yamada, Y.** (1999). Mice lacking link protein develop dwarfism and craniofacial abnormalities. *Nat. Genet.* **21**, 225-229.
- Wilkinson, D. J., Bhatt, S. and Herrmann, B. G.** (1990). Expression pattern of the mouse T gene and its role in mesoderm formation. *Nature* **343**, 657-659.