Immunolocalization of Proteoglycans in Meckel’s Cartilage of the Rat

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Abstract: The aim of this study was to investigate the presence and distribution of proteoglycans within Meckel’s cartilage of rat embryos. A standard indirect immunoperoxidase technique was used on paraffin sections of rat heads. Sections were incubated with monoclonal antibodies recognising core protein epitopes in the proteoglycans versican and CD44. Polyclonal antibodies localized the proteoglycans decorin, biglycan and lumican. Versican was expressed by chondrocytes, but very weakly by the extracellular matrix. Decorin was strongly expressed by both of chondrocytes and the ECM. Both of biglycan and lumican were moderately expressed by chondrocytes, but weakly by the extracellular matrix. CD44 was weakly expressed by chondrocytes only, without staining of the ECM. It is concluded that Meckel’s cartilage chondrocytes express the proteoglycans versican, decorin, biglycan, lumican and CD44 at variable levels during development in the rat. Such data are important for a greater understanding of the changes that take place during mandibular development. Further studies are needed to elucidate the exact role of proteoglycans during Meckel’s cartilage and mandibular organogenesis.

Key Words: Immunohistochemistry, Meckel’s cartilage, proteoglycans, rat.

INTRODUCTION

Proteoglycans (PG) are a group of macromolecules each composed of a core protein to which one or more highly anionic glycosaminoglycan (GAG) chains are attached through a specific sequence of trisaccharides: galactose-galactose-xylose. Proteoglycans play essential roles in the development, maintenance of function and integrity of virtually all tissues [1]. They are pivotal in cell-cell, and cell-extracellular matrix (ECM) interactions, e.g. control of mineralization, control of collagen fibrillogenesis, in addition to acting as cell-surface receptors [2]. Furthermore, PG are believed to govern the process of cell movement in numerous physiological and pathological conditions [3].

During normal embryonic development in mammals, the mandible is formed from the mandibular process of the first branchial arch. The mesenchyme of the mandibular process is derived from cranial neural crest cells, and paraxial mesoderm [4]. While the mesoderm gives rise to muscular and vascular tissues, the neural crest cells give rise to the mandibular arch skeleton, including Meckel’s cartilage [4]. The mandibular bone develops by intramembranous osteogenesis lateral to Meckel’s cartilage. Meckel’s cartilage is a pair of cylindrical rods of hyaline cartilage that give rise at their proximal ends to the malleus and incus of the middle ear [5]. The rostral (anterior to the mental foramen) and auricular (extending to the malleus) ends undergo endochondral type osteogenesis; the entire middle portion gets resorbed in the absence of any obvious formation of a calcified matrix and these cells undergo terminal cellular hypertrophy. However, the mid portion of the soft tissue of Meckel’s cartilage, at the degenerating stage, gives rise to the sphenomandibular ligament [6]. It has been demonstrated that chondroitin sulphate proteoglycans (CSPG), namely the hyaluronan (HA)-binding PG aggrecan, are the main PG of hyaline cartilage [7]. Decorin, biglycan, fibromodulin and epiphycan have also been localised to hyaline cartilage [8]. These PG have been ascribed several important roles such as ECM assembly, control of mineralisation and acting as cell surface receptors. Temporop-spatial variation of PG expression and composition during embryogenesis regulate tissue formation and integrity. For example, quail embryonic aggrecan has been shown to curtail the ability of neural crest cells to migrate within tissues [9]. Similar roles may be played by Meckel’s cartilage PG. It is still unclear how Meckel’s cartilage influences the surrounding developing mandibular bone. However, PG in Meckel’s cartilage appear to be essential in maintaining the cartilage in an un-mineralized state, for certain periods of time. PG may also be involved in communication with the surrounding mesenchymal tissues.

Epithelial-mesenchymal interactions play a pivotal role in histogenesis during mandibular development. Such interactions are mediated by the expression of various signalling molecules, such as PG and their components; the glycosaminoglycans (GAG) [10]. Animal studies, utilising 3-dimensional reconstruction, indicate that the mandible mainly grows at its periphery, and that this growth is regulated by paracrine signalling from Meckel’s cartilage and the developing tooth buds [11]. This signalling is probably mediated by a variety of macromolecules such as transforming growth factors-beta (TGF-β), bone morphogenetic proteins (BMP) and PG [11]. In an attempt to understand the role of PG in the development of the growing mandible, this study was conducted to investigate the expression and distribution of a
number of PG species at Meckel’s cartilage during in utero development in a mouse model utilising immunohistochemistry.

MATERIALS AND METHODS

Antibodies

Paraformaldehyde and sodium cacodylate (BDH Chemical Ltd Poole, England) were used for tissue fixation. A sledge base microtome (Leitz) was used to cut sections. For immunolocalization, a kit utilising a standard indirect immunoperoxidase method was used (Vectastain Quick kit, Vector Lab., California). The enzyme chondroitinase ABC (Sigma Chemical CO) was used to increase accessibility of the antibodies to their epitopes. Well-characterised antibodies were used and are listed in Table 1. These include the monoclonal (MAb) antibody 12C5 against versican, the polyclonal antibody (PAb) LF-113 against murine decorin, the (PAb) LF-106 against murine biglycan, a PAb recognising the core protein of lumican, and the MAb Hermes I against CD44.

METHODS

Tissue Preparation

This animal study was reviewed and approved by the institutional ethics committee. The specimens examined included rat heads of embryonic stages E18, E19 and E20, corresponding to post conception days 18, 19 and 20. These were fixed in 4% paraformaldehyde and 0.1 M Na cacodylate buffer for up to 48 h at 4°C. The specimens were dehydrated in 90% ethanol, embedded in paraffin wax, and 5-10 μm sections were cut. To observe normal histology, haematoxylin and eosin sections were prepared for light microscopic examination.

Immunohistochemical Staining

Prior to staining, the sections were de-paraffinized in xylene and rehydrated for 5 min in descending concentrations of alcohol (100% twice, 95% and 70% ethanol), water and PBS. To increase antibody access to epitopes, sections were pre-treated with chondroitinase ABC 0.1 IU/mL and PBS. To increase antibody access to epitopes, sections were pre-treated with chondroitinase ABC 0.1 IU/mL and PBS. To increase antibody access to epitopes, sections were pre-treated with chondroitinase ABC 0.1 IU/mL and PBS. After two 5-min rinses in PBS and Tween, the sections were incubated in streptavidin/peroxidase complex solution for 5 min (Vectastain Quick kit, Vector lab., California) and left on the sections for 60 min at room temperature. The slides were rinsed twice in PBS and Tween for 5 min and incubated with the biotinylated secondary antibody (raised in horse and recognises rabbit, mouse, goat, bovine and sheep IgG, Vectastain Quick kit, Vector lab., California) for 10 min at room temperature. After two 5-min rinses in PBS and Tween, the sections were incubated in streptavidin/peroxidase complex solution for 5 min (Vectastain Quick kit, Vector lab., California) and rinsed as above. The sections were then incubated with the peroxidase substrate solution, containing diaminobenzidine (DAB) until the brown stain developed intensely. The sections were thoroughly rinsed with double distilled water, counterstained with haematoxylin and rinsed in water, PBS and water again. They were dehydrated in ethanol, cleared in xylene and mounted for LM examination. Photomicrographs were taken with Leica microscope, using Kodak Ektachrome films.

Controls

Positive and negative control sections were used. With polyclonal (rabbit) antibodies against decorin, biglycan and lumican, some negative control sections were incubated with normal rabbit serum (concentration 1:1000, Sigma), which does not interact with the tissues studied. With monoclonal antibodies, serum or mouse immunoglobulin (MIG, Sigma, concentration 1:500) were used instead of the primary antibodies. Other sections were incubated with PBS in place of the primary antibody. The developing tooth buds in each section were used as positive/negative control tissues, based on findings of previous studies [16, 17].

Table 1. Details of the Antibodies Used in this Investigation

| Antibody | Epitope | Type | Dilution | Source | Reference |
|----------|---------|------|----------|--------|-----------|
| 12C5     | Hyaluronate binding region of versican | Monoclonal mouse IgG1 | 1:200 | Purchased from Dr. R.A. Asher, Developmental Studies Hybridoma Bank, Iowa, USA | Asher et al. [12] |
| LF-113   | Core protein epitopes of murine decorin | Polyclonal rabbit IgG | 1:100 | A generous gift from Professor L. Fisher, National Institute of Health, USA | Fisher et al. [13] |
| LF-106   | Core protein epitopes of murine biglycan | Polyclonal rabbit IgG | 1:100 | A generous gift from Professor L. Fisher, National Institute of Health, USA | Fisher et al. [13] |
| Anti-lumican | Core protein epitopes | Polyclonal, rabbit IgG | 1:200 | A generous gift from Professor J R Hassel, University of Pittsburgh, Pennsylvania, USA | Chakravarti et al. [14] |
| Hermes I | The N-terminal Hyaluronate binding domain of CD44 | Monoclonal rat IgG2 | 1:10 | Developmental Studies Hybridoma Bank, Iowa, USA | Picker et al. [15] |
RESULTS

Immunolocalization of Versican

After incubation with MAb 12C5 and chondroitinase ABC pre-treatment, the following results were obtained (Fig. 1A, B, C). At E18 there was a positive staining of chondrocytes of Meckel’s cartilage. Similar distribution and intensity of staining were observed in the E19 and E20 rat embryos. In all embryonic stages, there was only very weak staining of the ECM.

Immunolocalization of Decorin

Following incubation with LF-113 (a PAb which specifically recognises murine decorin), and chondroitinase ABC pre-digestion, an intense positive staining of chondrocytes and ECM of Meckel’s cartilage was observed at the embryonic stages E18, E19 and E20 (Fig. 1D, E, F).

Immunolocalization of Biglycan

Following incubation with LF-106 (a PAb which specifically recognises murine biglycan), with chondroitinase ABC pre-digestion, positive staining of chondrocytes was observed at E18. More intense staining was evident at E19 and E20. The ECM was mildly stained at the three stages (E18 to 20) (Fig. 1G, H, I).

Immunolocalization of Lumican

Following incubation of sections with the PAb recognising core protein epitopes in lumican, at the E18 stage, weak immunoreactivity, and at E20 Meckel’s cartilage showed strong immunoreactivity. The ECM was mildly stained at the three stages (E18 to 20) (Fig. 1J, K, L).

Immunolocalization of CD44

After incubation of sections with MAb Hermes 1, which recognises CD44, very weak immunoreactivity was observed on chondrocytes at the periphery of Meckel’s cartilage at the three stages examined (E18-E20). No evidence of staining could be seen in the ECM (Fig. 1M, N, O).

Controls

Control sections where the primary antibody was substituted with PBS, mouse immunoglobulin or rabbit serum, were all negative (Fig. 1P). The developing tooth buds and dental tissues, such as the pulp and dentine in each section, which were used as positive/negative control sections (Fig. 1Q, R), exhibited findings consistent with previous studies [16, 17].

DISCUSSION

During intra uterine life, the ECM of Meckel’s cartilage remodels by the action of proteolytic enzymes secreted by its own chondrocytes, and most of the cartilage disintegrates soon after the onset of mandibular bone formation. Although it is generally accepted that Meckel’s cartilage is essential for mandibular growth and development, the exact role of this cartilage, and thus, the roles of the tissue factors expressed within it, are still unclear.

The midportion of Meckel’s cartilage in rats and mice disappears soon after birth, probably by cellular resorption of the chondrocytes in the anterior part of the midportion, and by differentiating into fibroblastic cells in the posterior part of the midportion [18]. In the present study, we have selected rat embryonic stages E18-E20 (usually, the last gestational day) before any disintegration of the cartilage rods, or transformation of the chondrocytes into fibroblasts within the midportion of Meckel’s cartilage has taken place. It is expected that in newborn rats the anterior part of the midportion of Meckel’s cartilage may be completely missing, and changes in the posterior part of the midportion may have started [18]. Therefore, in the present work only embryonic Meckel’s cartilage was studied. Most previous studies have investigated earlier stages Meckel’s cartilage; for example, Lee et al. [19] investigated Meckel’s cartilage at embryonic days 15-18. Moreover, some PG like versican could not be detected at earlier embryonic stages such as E15 [19]. Our aim was to investigate PG in Meckel’s cartilage at later embryonic stages just before chondrocyte transformation, or disappearance, have occurred.

Meckel’s cartilage chondrocytes share many phenotypic characteristics with other cartilaginous cells. However, they show some differences from trunk cartilage chondrocytes. For example, Meckel’s cartilage chondrocytes are the only cartilage cells that are capable of producing type I collagen in tissue cultures [6]. This probably reflects the different origins of this cartilage as, unlike other cartilage, which is derived from mesoderm, Meckel’s cartilage originates from neural crest ectomesenchyme [6]. We have shown that the PG species versican, the small leucine-rich proteoglycans (SLRPs) decorin, biglycan and lumican, and the cell surface HA-binding PG CD44 are present in Meckel’s cartilage of the rat during in utero mandibular development with variable degrees. Although CSPG have previously been shown to be present in Meckel’s cartilage [2], to date, this is first study to describe the expression of a wide range of PG in Meckel’s cartilage.

Versican (also known as PG-M) is a large CSPG of the ECM. In various developing embryonic tissues, versican is transiently expressed and plays important roles in cell adhesion [20], migration [21], proliferation, and differentiation [22]. It localizes in various tissues both in fetal and adult animals [21, 23-25]. In particular, it is expressed in prechondrogenic mesenchyme, and during the transition to cartilage its expression becomes restricted to the periphery of the newly formed cartilage [21]. In addition, versican has been isolated from rat dental pulp [26], periodontium [27], and developing mandibular condylar cartilage [28]. It has been suggested that versican is involved in positively regulating the formation of the mesenchymal matrix and the onset of chondrocyte differentiation through the attached chondroitin sulfate chains [29]. Another study [30] strongly suggests that mature versican is essential for precartilage aggregation and subsequent cartilage differentiation.

In developing cartilage, versican is transiently expressed at a high level in the mesenchymal condensation area and rapidly disappears during cartilage development [31]. Immunohistochemical studies on developing limb bud cartilage revealed that an area positive for versican gradually shifts out of the diaphysis and is replaced by an area positive for
Fig. (1). Immunoperoxidase staining for Versican showing positive staining of moderate intensity on Meckel’s cartilage chondrocytes and weak staining within the ECM at all embryonic stages (A, B, C). Decorin showed an intense positive staining on chondrocytes and within the ECM at all embryonic stages (D, E, F). Biglycan was moderately expressed at E18 (G), and more intense staining at later stages (H, I). Staining for Lumican on the chondrocytes and ECM was weak at E18 (J), moderate at E19 (K), and intense at E20 (L). Immunoperoxidase staining for CD44 showed very weak staining on chondrocytes at all stages, without any staining within the ECM (M, N, O). P, Q and R represent the control sections.

P: negative control section; Q: positive staining for biglycan in the enamel organ and dental papilla of a developing molar tooth bud at the E19 stage; R: positive staining for decorin in the dental papilla, odontoblasts and dentin in a developing tooth bud. OOE: outer enamel epithelium; SR: stellate reticulum; IEE: inner enamel epithelium; dp: dental pulp, df: dental follicle.
aggreecan [24]. Differentiating chondrocytic cells showed similar expression patterns of these molecules [29]. Therefore, it was generally accepted that mature chondrocytes do not express versican (but express aggreecan). However, although versican has not been detected by immunohistochemical studies, constitutive low level transcription of the versican gene has been observed in cartilage and chondrocytes [31, 32]. Furthermore, biochemical studies demonstrated the presence of versican in human articular cartilage from the fetal stage to mature adult [33]. Unlike previous immunohistochemical studies, we have demonstrated positive immunostaining of versican within chondrocytes of Meckel’s cartilage. It is possible that the expression of versican is a unique feature of Meckel’s cartilage at later embryonic stages.

In the present study, versican was expressed by Meckel’s cartilage chondrocytes in all embryonic stages, with no changes detected in the intensity of staining as the embryos matured. Our findings are in general agreement with those of Ishizeki and Nawa [34], who demonstrated the expression of CSPG, using immunohistochemical methods, in addition to toluidine and Alcian blue staining. However, our findings contrast with those of Lee et al. [19], who were not able to demonstrate immunoreactivity to versican in Meckel’s cartilage. The reason for this disagreement may be that, in their study, Lee et al. [19] have included earlier stage Meckel’s cartilage (embryonic days 15 to 18), whereas in our study, later stages (E18 to E 20) were investigated. Another reason may be that Lee et al. [19] have used a different antibody. They have used the MAb (5D5) which recognises the core proteins of large PG, such as versican, neurocan and brevican. The antibody used in our study (12C5) is a more specific antibody, which specifically recognises the HA-binding region of versican core protein.

The SLRPs belong to the large family of leucine-rich repeat proteins, which are characterized by multiple adjacent domains bearing a common leucine-rich motif. They can be divided into several sub-families based on their gene organization, the number of leucine-rich repeats and the type of GAG chain substituent. Decorin and biglycan are part of one subfamily, whereas fibromodulin and lumican are part of a second subfamily. The GAG chains have been associated with the interaction with several growth factors, and enable the SLRPs to provide a sink for growth factor accumulation within the extracellular matrix. The SLRPs can therefore, help modulate chondrocyte metabolism by regulating growth factor access to the cells [35].

Decorin and biglycan can be detected in fetal cartilage and bone [36]. Although decorin is involved in the organization of collagen fibrils, in the modulation of collagen metabolism and in interactions with various glycoproteins, retinoic acid and growth factors, e.g. TGF-β [37], the functions of biglycan are not well understood. It may participate in the control of chondrocyte proliferation [31]. *In vitro* studies of human articular cartilage have shown that the expression level of decorin increases and that of biglycan decreases with age [38]. Within developing bone, biglycan is localized to the walls of the osteocyte lacunae and bone cell surface, whilst decorin is present throughout the osteoid matrix and associated with bone cells associated with osteogenic and non-osteogenic layers of the periosteum [36]. We have found that decorin is expressed by chondrocytes and surrounding ECM without any change in intensity of staining in the three embryonic stages examined. As for biglycan, staining was less intense than decorin, but gradually increased throughout development. Similarly, Kavanagh and Ashhurst [39], investigating the immunolocalization of biglycan during development and aging of articular cartilage in the rabbit knee joint, found that biglycan is present in the deeper regions of the epiphysial cartilage, which eventually ossifies, but could not be detected in the interzone, the region from which the articular cartilage develops and maintains its cartilaginous nature after birth. Decorin and biglycan have been isolated from predentine, predentine-dentine interface and the mineralised dentine ECM [40]. Milan et al. [41] have demonstrated that the composition and functions of decorin and biglycan differ with transition from the non-mineralised predentine towards the mineralised dentine matrix. Furthermore, it was found that with this transition from the non-mineralised to mineralised dentine, there was an increasing affinity to hydroxyapatite, which was predominantly facilitated through GAG chains [42]. Based on the above findings, our results of a sustained expression of decorin and biglycan in Meckel’s cartilage might suggest a role of these SLRPs in inhibiting mineralisation of this cartilage.

In the present study, lumican was gradually upregulated throughout embryonic stages E18 to E20. In accordance with our results, Ying et al. [43] have demonstrated, using northern plot and *in situ* hybridisation, that in earlier stages mouse embryos, there is almost no lumican expression, but that the expression of this PG increases with organ development and function.

CD44 is an adhesion molecule distinct from cadherin and integrins. This protein functions as a HA receptor and is involved in cell migration [44]. *In vitro* studies suggest that CD44 is expressed in chondrocytes [45]. In the present study, CD44 was weakly expressed by Meckel’s cartilage chondrocytes. This pattern of CD44 expression in rat Meckel’s cartilage is different from that previously reported in rat tibial cartilage [46]. However, it is consistent with its distribution in the rat mandibular condylar cartilage [47]. Cranial bone and cartilage, including Meckel’s cartilage and mandibular condylar cartilage, originate in neural crest-derived mesenchyme [48]. In contrast, bones and cartilage in limbs are derived from lateral plate mesoderm. Thus, Meckel’s cartilage has a different developmental origin from tibial cartilage. Therefore, the different pattern of CD44 expression observed in these two types of cartilage may be due to their different origins. In fact, CD44 is used as a marker of neural-crest-derived cells and embryonic stem cells [49].

The temporal expression of PG during various stages of development found herein, suggests that the expression of PG may be up- or down-regulated by chondrocytes as necessary for cell-cell or cell-ECM signalling within Meckel’s cartilage, or between chondrocytes of Meckel’s cartilage and the surrounding mesenchymal tissues and developing alveolar bone. Changes in the expression and chemical composition of various macromolecules, including PG and growth factors, are well-known events during histogenesis and morphogenesis of various tissues during orofacial development, including mineralised tissues such as tooth dentine [40-42]. The very exact role of PG in Meckel’s cartilage
cannot be readily revealed from the present work, however, we postulated, based on the functions of PG reported by other studies [50] that Meckel’s cartilage PG may inhibit mineralization of the cartilage during mandibular development, where signalling from Meckel’s cartilage chondrocytes and/or ECM is believed to be necessary for mandibular development, which may be mediated by PG. It is known that the functions of PG including SLRPs depend upon their core protein and GAG chains. SLRPs interact, via their core protein, with fibrillar collagen, thus regulating fibril diameter and fibril-fibril interactions. Attachment of core proteins to collagen also limits access of collagenase to its substrate, thereby protecting collagen from proteolytic action [50]. The function of SLRPs, such as decorin, biglycan and lumican is best illustrated by the abnormal phenotypes in knock-out mice. Inhibition of decorin expression was associated with thinning of the skin, accumulation of loose connective tissue in the subdermal layer, abrupt fracture and dissection of the skin between the deeper dermis and the fascia [51]. Biglycan knockout mice demonstrated reduced growth rate, decreased bone mass and differences in bone shape, when compared to normal mice [52]. Absence of lumican produces skin laxity and corneal opacity, with an increased proportion of abnormally thick collagen fibrils and delayed corneal epithelial wound healing [53]. Therefore, the PG of Meckel’s cartilage are probably involved in similar functions, including regulation of collagen fibrillogenesis, preservation of Meckel’s cartilage integrity, inhibition of cell proliferation, inhibition of Meckel’s cartilage mineralization and possibly influencing the developing mandibular bone surrounding Meckel’s cartilage.

The results presented here show how the distributions of the PG versican, decorin, biglycan, lumican and CD44 vary during development in rat Meckel’s cartilage. In addition, they suggest that the Meckel’s cartilage chondrocytes are a distinct population of cells. Such data are important for a greater understanding of the changes that take place during mandibular development. Further studies are needed to elucidate the exact role of PG during Meckel’s cartilage and mandibular organogenesis.

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