Differential Relative Sulfation of Keratan Sulfate Glycosaminoglycan in the Chick Cornea during Embryonic Development

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Purpose. To investigate structural remodeling of the developing corneal stroma concomitant with changing sulfation patterns of keratan sulfate (KS) glycosaminoglycan (GAG) epitopes during embryogenesis and the onset of corneal transparency.

METHODS. Developing chick corneas were obtained from embryonic day (E)12 to E18 of incubation. Extracellular matrix composition and collagen fibril spacing were evaluated by synchrotron x-ray diffraction, hydroxyproline assay, ELISA (with antibodies against lesser and more highly sulfated KS), and transmission electron microscopy with specific proteoglycan staining.

RESULTS. A significant relative increase in highly sulfated KS epitope labeling occurred with respect to hydroxyproline content in the final week of chick development, as mean collagen interfibrillar distance decreased. Small KS PG filaments increased in frequency with development and were predominantly fibril associated.

CONCLUSIONS. The accumulation of highly sulfated KS during the E12 to E18 timeframe could serve to fine tune local matrix hydration and collagen fibril spacing during corneal growth, as gross dehydration and compaction of the stroma progress through the action of the nascent endothelial pump. (*Invest Ophthalmol Vis Sci.* 2010;51:1365–1372) DOI:10.1167/iovs.09-4004

The cornea is a connective tissue with remarkable transparency that forms the primary refractive surface of the eye; yet, how this property arises in development remains little understood. The most widely studied model of corneal development is that of the embryonic chicken. In this tissue, uniformly thin collagen fibrils are laid down by stromal fibroblasts in a secondary stroma and consolidate as a series of orthogonal fibril bundles that later form into lamellae. At

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embryonic day (E)14, the chick cornea transmits only approximately 40% of white light, but then begins to increase in transparency so that, at E19, transmission is more than 95%, similar to the adult condition. The transparency increase after E14 is accompanied by significant dehydration of the cornea, flattening of keratocytes, and compaction of stromal collagen fibrils into a spatially ordered array. This restructuring of the stroma is key for the acquisition of corneal transparency. The stroma is key for the acquisition of corneal transparency.

Small leucine-rich proteoglycans (PGs) interact with collagen fibrils in the corneal stroma and are thought to help control fibril size and spatial organization. PGs are composed of a core protein covalently bound to sulfated glycosaminoglycan (GAG) side chains. The major GAG in the cornea is keratan sulfate (KS), and three types of PG bear these side chains: lumican, Reratocan, and mimecan (or osteoglycin). Pg E18, in the chick stroma KS PGs bear two to three GAG chains of approximately 15 kDa each, and the PGs lumican and keratocan have three of five potential linkage sites variably substituted with KS GAGs. Collagen-PG interactions are thought to occur via the PG core protein, with sulfated GAG chains extending into the interfibrillar space, where they confer stromal water-binding and sorptive tendencies that help define the hydrophilicity and swelling properties, and thus the ultrastructure, of the stromal matrix.

Corneal PGs, especially those carrying KS chains, fulfill an important role in the establishment of a properly formed stroma during corneal embryogenesis. 16 The importance of these molecules is illustrated by the fact that the corruption of sulfate motifs on KS is associated with structural matrix alterations in the corneas of humans with the inherited disease macular corneal dystrophy 17,18 and in the corneas of mice with gene-targeted deletions in lumican 19,20 or Chst5, a KS sulfotransferase enzyme.²¹ Some investigators have reported the accumulation of KS GAGs and KS PGs during embryonic development. 22-25 Molecular studies have shown that levels of mRNA for keratocan, first expressed at E6,26 exhibit a steady decline from E9 to E18, whereas, over the same time period, lumican mRNA levels remain several-fold higher.²⁷ Moreover, an approximate threefold increase in lumican core protein occurs between E7 and E9,²⁸ with a two- to threefold increase in mRNA for β -1,4-galactosyltransferase, an enzyme involved in sulfated KS synthesis, reported between E8 and E13.²⁹ Biochemical analyses have also shown that KS is synthesized by the chick cornea between E5 and E7, but that it only becomes highly sulfated by E14,30 with another study indicating a switch in the production of unsulfated to sulfated KS between E12 and E15.²⁸

The current investigation was designed to test the hypothesis that sulfation of KS provides an environment of hydration conducive to the deposition of collagen fibrils and the establishment of a highly ordered stromal matrix in the developing chick. To this end, experiments were conducted to quantify

collagen biosynthesis, KS sulfation status, collagen fibril spacing, and collagen-KS PG associations during the latter stages of chick corneal morphogenesis.

METHODS

Specimens

For x-ray diffraction and biochemical studies, a series of 69 embryonic chick corneas was obtained that comprised daily incremental stages from E12 to E18. Corneas were excised by an incision around the limbus in eyes of embryonic chicks obtained from Hamburger-Hamilton (HH)-staged, fertilized eggs collected from a commercial hatchery (Hy-Line UK, Warwickshire, UK) approximately 2 hours earlier, where E12 = HH38, E13 = HH39, E14 = HH40, E15 = HH41, E16 = HH42, E17 = HH43 and E18 = HH44. Immediately on excision, the corneas were sandwiched between layers of clingfilm to minimize dehydration and were placed on dry ice. Frozen corneas were transferred to -80°C storage until the synchrotron x-ray diffraction experiments could be conducted. For electron microscopy, a smaller group of fertilized eggs (Henry Stewart & Co. Ltd., Louth, Lincolnshire, UK) was incubated at 37°C until staging and excision. All work was conducted in accordance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research, as well as with local ethics guidelines.

Synchrotron X-Ray Diffraction

Individual corneas (n = 69), still secured in cling film, were positioned in a sealed specimen holder between sheets of x-ray-transparent mylar where they were allowed to thaw. In turn, each cornea (n = 8-11daily from E12 to E18) was then placed into the path of a focused (1 \times 1 mm) monochromatic ($\lambda = 0.154$ nm) x-ray beam on beamline 2.1 of the Synchrotron Radiation Source, Daresbury Laboratory (Cheshire, UK). All corneas were exposed for 2 minutes each, and the resulting fiber diffraction patterns were recorded on a multiwire, gas proportional area detector plate positioned 8.25 m behind the cornea. Immediately after x-ray exposure, the corneas were refrozen and transported to the home laboratories for biochemical and immunochemical analyses (described in the following sections). Diffraction patterns (512 \times 512 pixels) were analyzed with purpose-written software (Unix-based; Sun Microsystems, Mountain View, CA) and graphics and statistics packages (Unix; Optimas, Bothell, WA; Statistica; Statsoft, Tulsa, OK, Excel; Microsoft, Redmond, WA), as described previously.³¹ The position of the first-order equatorial (i.e., interfibrillar) reflection, calibrated to the 67-nm axial D-periodic repeat of collagen in wet rat tail tendon, was used to provide a measure of the mean center-to-center collagen fibril Bragg spacing as an average throughout the whole thickness of the central cornea. Statistical significance was ascertained by using statistical tests for large data groups: one-way ANOVA with post-hoc Tukev HSD.

After x-ray analysis, the corneas were refrozen for transportation to the home laboratory and digested for 17 hours at 60°C with 1 mg/mL papain (Sigma-Aldrich, Poole, UK) in 0.05 M sodium acetate buffer (pH 5.6), containing 0.025 M EDTA and 5 mM cysteine HCl. The enzyme was inactivated at 100°C , and the digests stored at -20°C until further analysis.

Hydroxyproline Assay

Papain digests from individual corneas were hydrolyzed by using equal volumes of 11.7 N concentration HCl to supernatant at 110°C overnight. Specimens were then lyophilized. Dried hydrolyzates were reconstituted in their starting volume of distilled water and centrifuged to remove particulate material. Hydroxyproline residues were assayed in triplicate as previously described, ³² against known standards and read at 540 nm after 10 to 20 minutes incubation at 70°C. Statistical significance was ascertained by using statistical tests for large data groups: one-way ANOVA with post-hoc Tukey HSD.

KS Quantification by Competitive ELISA

Optimized, competitive ELISAs were developed to quantify pentasulfated hexasaccharides and tetrasulfated hexasaccharides as the smallest linear structures in KS chains, as recognized by the monoclonal antibodies 5D4 and 1B4, respectively. ³³⁻³⁵ Ninety-six-well EIA microtiter plates (MP Biomedicals, Cambridge, UK) were coated by passive adsorption with a 250 ng/mL chondroitinase ABC-digested bovine articular cartilage aggrecan (BAC ABC core) antigen in a 20-mM sodium carbonate coating buffer (pH 9.6), for 14 hours at 37°C. Native aggrecan core protein is substituted with both chondroitin sulfate (CS) and KS chains, and CS was selectively removed by chondroitinase digestion to leave a KS-linked coating protein. The plates were washed with Tris saline azide (TSA) and the unreacted sites blocked with the addition of 1% (wt/vol) bovine serum albumin (BSA) in TSA. All incubations were performed for 1 hour at 37°C.

Papain digests from single corneas were serially diluted and allowed to bind with an equal volume of 5D4 (1:8000 dilution in 1% BSA/TSA) and incubated to compete against the BAC ABC core. A standard curve was generated from serial dilutions of BAC ABC core/5D4. The plates were then washed with TSA before incubation with alkaline phosphatase-conjugated goat anti-mouse antibody (1:5000 dilution; Promega, Madison, WI). The plates were again washed before alkaline phosphatase substrate (p-nitrophenyl phosphate, 1 mg/mL) was applied in DEA buffer (0.126 mM MgCl₂, 1 M diethanolamine, pH corrected to 9.8). Color development was quantified on a plate reader (Multiskan MS; Labsystems, Helsinki, Finland) at 405 nm, to determine the inhibition of binding.

The same stock corneal extracts were then assayed for lesser sulfated KS by using the 1-B-4 monoclonal antibody. Optimization of conditions established that microtiter plates were coated with 125 ng/mL BAC ABC core and ELISAs performed as described above, with a 1:4000 1-B-4 dilution. Statistical significance was ascertained with statistical tests for large data groups: one-way ANOVA with post-hoc Tukey HSD.

Electron Microscopy

On excision, corneas from E12 to E18 chicks (n = 21) were prepared for electron microscopy, as described previously. 21,36 A series of E12 to E18 corneas was fixed in 2.5% glutaraldehyde in 25 mM sodium acetate buffer with 0.05% cuprolinic blue (0.1 M MgCl₂; pH 5.7), to stain the sulfated PGs. Another series was fixed for 10 minutes in 4% paraformaldehyde (pH 7.2). These corneas were then washed twice in chondroitinase buffer (50 mM Tris, [pH 8], 60 mM sodium acetate, and 0.02% BSA) and guartered. Corneal pieces were incubated for 4 hours at 37°C in chondroitinase buffer containing 2.5 U/mL chondroitinase ABC (from Proteus vulgaris; Sigma-Aldrich) to remove the chondroitin- and dermatan-sulfate GAG chains from the tissue. Commercially prepared protease inhibitor cocktail for general use (Sigma-Aldrich) was also added to 1% total volume to inhibit endogenous PG-degradative activity. The control corneas were similarly processed in buffer minus enzyme. The samples were washed in chondroitinase buffer (minus enzyme) before fixation and staining overnight, as before. All corneas were then contrast enhanced with sodium tungstate and dehydrated and embedded in Araldite, according to standard protocols. 36 Ultrathin sections (~90-nm thick) were cut on glass knives, collected on copper grids, and stained with 1% aqueous phosphotungstic acid and saturated aqueous uranyl acetate before examination by transmission electron microscopy (JEM-1010; JEOL, Tokyo, Japan) equipped with an 11-megapixel CCD camera (Orius SC1000 CCD camera; Gatan Ltd., UK, Cambridge, UK).

RESULTS

Mesenchymally derived presumptive keratocytes that populate the developing chick cornea are responsible for establishing the secondary stroma by synthesizing and depositing hybrid collagen type I/V fibrils.^{1,2} Quantification of collagen as hy-

TABLE 1. Average Amount of Hydroxyproline and Average Collagen Interfibrillar Spacing in Embryonic Chick Corneas

| | E12 (n = 8) | E13 (n = 10) | E14 (n = 11) | E15 (n = 10) | E16 (n = 11) | E17 $(n = 10)$ | E18 (n = 9) |
|---|------------------------------------|------------------------------------|------------------------------------|----------------------------------|-------------------------------------|-------------------------------------|-------------------------------------|
| Hydroxyproline, µg/mL Collagen fibril Bragg spacing, nm | 4.38 ± 0.18 60.8 ± 0.60 | 8.06 ± 1.07 63.1 ± 1.01 | 7.43 ± 0.48 63.9 ± 0.99 | 17.48 ± 1.46 61.9 ± 1.22 | 17.23 ± 0.74 59.8 ± 1.53 | 34.52 ± 4.11 57.0 ± 0.87 | 34.69 ± 1.64 53.7 ± 0.94 |

Embryonic days E12 to E18 are equivalent to Hamburger-Hamilton stages 38 to 44 of development. Hydroxyproline measurements are also representative of total collagen content per cornea. Data are expressed as the mean \pm SE.

droxyproline (Table 1) indicated two significant increases in hydroxyproline content, found between E14 and E15 (P < 0.005) and E16 and E17 (P = 0.001). At E16 Coleman et al.³⁷ measured 20.7 μg hydroxyproline per cornea and Conrad, ³⁸ 19.5 μ g. Our value of 17.23 μ g/cornea at this developmental stage is consistent with these published values.

X-ray diffraction patterns generated by corneas at E12 gave rise to collagen interfibrillar reflections that were consistent with a rather loose, hydrated network of formative lamellae that constitutes the tissue at this time. With development, interfibrillar reflections became more intense, presumably as a consequence of growth and the added deposition of collagen as fibrils, consistent with the results of the hydroxyproline analysis shown in Table 1. Analysis of all 69 x-ray diffraction patterns showed that the average center-to-center collagen fibril spacing from E14 to E18 decreased as the cornea thinned and became more transparent (Table 1).

These collagen spacing data were highly representative of the corneal stroma as a whole, with all fibrils in the path of the x-ray beam (which measured approximately 1×1 mm) passing through the whole thickness of the cornea contributing to the x-ray pattern and thus the calculated values. X-ray analysis, however, could not identify how the stromal matrix was changing at the suprafibrillar level. Electron micrographs confirmed the overall nature of the stromal compaction and indicated, not a homogeneous coming together of widely and uniformly spaced fibrils with time, but rather a progressive coalescence of bundles of collagen fibrils in which the fibrils were already fairly well organized and packed laterally (Fig. 1). Based on the fundamental principles of x-ray fiber diffraction, the collagen-free spaces between fibril bundles, which were quite prevalent before E15, did not contribute to collagen interfibrillar reflection. Thus, the values displayed in Table 1 derive from collagen fibrils that are regularly spaced, whether they exist in bundles, as in the earlier stages that we investigated, or in stacked lamellae, as seen later in development. Our observations (Fig. 1), together with those of other investigators^{3,6} showed that stromal compaction during development entailed collagen fibril bundles coming together to form lamellae, whereas x-ray data indicated that fibrils within the bundles and lamellae are moving closer together from E14 onward.

As synchrotron x-ray diffraction is a noninvasive technique, the 69 corneas from which ultrastructural collagen fibril data were acquired (Table 1) were available for subsequent immunochemical quantification of KS epitope. Data were normalized to hydroxyproline content to reflect changes in lessersulfated (1B4-immunoreactive) and highly sulfated (5D4immunoreactive) KS epitope, relative to matrix deposition (Table 2). It is important to remember that these values represent relative epitope labeling between samples and that the nature of the antibodies used potentially allows multiple binding sites within the same GAG chain. KS values are relative to BAC-ABC core protein labeling equivalents.

Levels of highly sulfated KS as a proportion of hydroxyproline throughout the E12-to-E18 interval were consistently higher than corresponding amounts of the low-sulfated epitope. The amount of lesser-sulfated KS fluctuated between E12 and E18, but did not change appreciably, whereas levels of highly sulfated KS increased markedly. As can be seen in Table 2, highly sulfated KS levels remained fairly steady between E13 and E15, after which a sharp rise occurred (P = 0.001), such that by E16 the average amount had more than doubled from levels at E15. Although there was a reduction subsequently, it is not statistically significant (P = 0.528). Examination by electron microscopy of corneas in the E12-to-E18 timeframe in which chondroitin/dermatan sulfate GAG chains had been enzymatically removed from the tissue revealed cuprolinic bluecontrasted sulfated KS PGs as small collagen fibril-associated filaments that increased in abundance with time.

Discrete fibril bundles coalescing into a continuous matrix have been observed by other investigators³ and are apparent in Figures 1, 2. Measurements of (1) hydroxyproline, (2) highly and lesser sulfated KS, and (3) collagen fibril spacing were performed for each of the 69 corneas examined (Tables 1, 2). Consequently, we can seek correlations independent of developmental day. In doing so, it is evident that a positive correlation exists between hydroxyproline content and highly sulfated KS, as assessed by 5D4 binding ($R^2 = 0.682$), and between hydroxyproline content and lesser-sulfated KS, as measured by 1B4 binding ($R^2 = 0.395$), as shown in Figure 3. These parallel changes presumably reflect embryonic growth and continued matrix deposition. However, no appreciable correlation was found between the ratio of KS to hydroxyproline and collagen fibril spacing, when data from all 69 corneas were analyzed independent of developmental stage (Fig. 4).

DISCUSSION

The changing sulfation patterns of KS in the developing chick cornea raise several questions regarding the likely functional role of this molecule with respect to its sulfation status. Sulfa-

Table 2. The Average Amounts of 1B4- and 5D4-Labeled KS Epitope Relative to BAC-ABC Core after Normalization to Hydroxyproline Content in Embryonic Chick Corneas

| | E12 $(n = 8)$ | E13 $(n = 10)$ | E14 (n = 11) | E15 $(n = 10)$ | E16 $(n = 11)$ | E17 $(n = 10)$ | E18 $(n = 9)$ |
|---|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|
| Lesser sulfated KS (1B4-labeled epitope/hydroxyproline) | 0.125 ± 0.043 | 0.128 ± 0.032 | 0.153 ± 0.046 | 0.129 ± 0.024 | 0.109 ± 0.016 | 0.114 ± 0.022 | 0.118 ± 0.024 |
| Highly sulfated KS (5D4-labeled epitope/hydroxyproline) | 0.775 ± 0.261 | 0.958 ± 0.152 | 0.865 ± 0.160 | 0.840 ± 0.111 | 1.879 ± 0.167 | 1.449 ± 0.143 | 1.874 ± 0.206 |

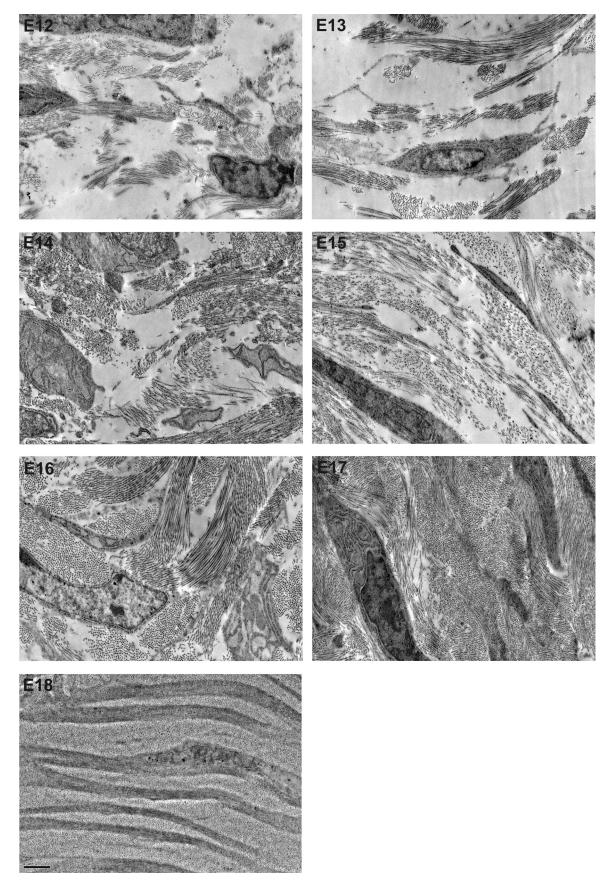


Figure 1. Transmission electron micrographs of developing chick corneal stroma. Tissue compaction progressed from loose bundles of collagen fibrils at E12 to a mature lamellar form at E18. Scale bar, $1~\mu m$.

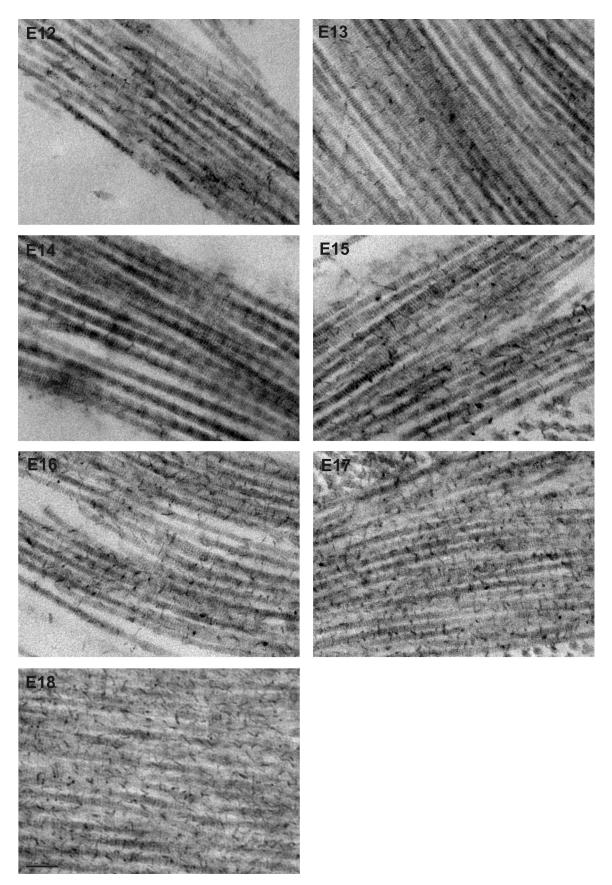


Figure 2. Transmission electron micrographs of chondroitinase digested developing chick corneal stroma. Increased deposition of cuprolinic blue-stained sulfated KS PG filaments occurs in the E12-E18 timeframe. Scale bar, $1~\mu m$.

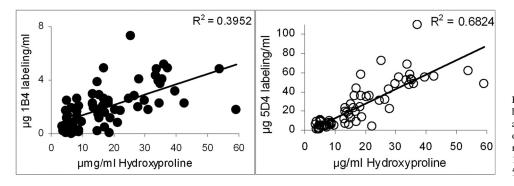


FIGURE 3. Correlation of antibody labeling in relation to BAC-ABC core against hydroxyproline content (without normalization to hydroxyproline, raw data not shown) for lesser (*left*, 184 reactive) and more highly (*right*, 5D4 reactive) sulfated KS.

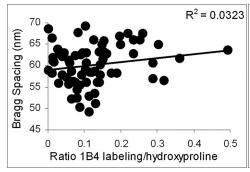
tion of individual corneal KS chains is not homogeneous, and unsulfated disaccharides, which are predominant nearer the linkage region, become increasingly sulfated toward the nonreducing terminal.³⁹ Moreover, KS chains undergo sequential elongation and sulfation during synthesis. 40 The absence of sulfotransferase enzyme in human corneas with macular corneal dystrophy, as in mutant corneas of Chst5-knockout mice, results in an immature, truncated GAG and an atypical corneal phenotype. 18,21 Thus, KS chain length and sulfation are dictated by tissue-specific factors such as the presence of processing enzymes. During chick development, galactose-transferase activity increases with KS biosynthesis and is maintained at an unusually high level in adult cells. 29 The sequential mechanism of GAG subunit sulfation becomes apparent in the absence of the sulfotransferase enzyme, when neither GlcNAc nor Gal residues are sulfated. 41 The negative charge provided by sulfation could serve to prevent KS GAG chain collapse during elongation and perhaps offer an explanation of the differential sulfation between shorter and longer KS chains. The presence of increasing amounts of highly sulfated KS during embryonic development is possibly linked to the appearance of the sulfate donor, 3'-phosphoadenosine-5'-phosphosulfate (PAPS). 42 The measured elevation of 5D4 labeling at the time of increased fibrillogenesis and matrix compaction could be a result of PAPS availability facilitating increases in KS chain length and greater sulfation. The combination of these events could enhance organized stroma formation through the binding of core proteins and the hydrophilic influence of sulfate groups.

A major purpose of the present study was to investigate the structural remodeling of the stroma in relation to changing sulfation patterns of constitutive KS epitope during chick corneal morphogenesis. The results confirm previous synchrotron x-ray diffraction studies that documented the progressive compaction of corneal collagen fibrils with embryonic growth in the chick. ^{43,44} In the dataset examined herein, compaction occurred after E14 (Table 1), with electron microscopy indicating that the compaction happened within bundles of collagen fibrils in which fibrils are already fairly well organized spatially (Fig. 1). The x-ray diffraction and electron microscopy

findings, when considered together, imply that collagen bundles are coalescing into lamellae, whereas, at the same time, fibrils within bundles are becoming more closely spaced. Between E15 and E16, the stroma experiences a significant increase in highly sulfated KS epitope that is disproportionate to the measured increase in hydroxyproline (Table 2). This observation is illustrative of an accelerated deposition of highly sulfated KS post-E15 which exceeds the anticipated in ovo corneal growth rate, as implied by hydroxyproline deposition. Thus, as the chick cornea develops and becomes transparent and compacted, the nature of KS antigenicity changes, with the highly sulfated KS fraction accumulating preferentially over that of the lesser-sulfated isoform and from a higher starting point. It seems counterintuitive to reason that negatively charged, highly sulfated KS GAG would accumulate as the stroma condenses and act as a biological driving force for tissue dehydration. Quite the opposite: The water-retentive properties of this hydrophilic macromolecule would be expected to swell the corneal matrix, rather than to aid in its compaction. Indeed, the current data show that the stroma was already undergoing compaction when the cornea, at E15, experienced the first sizable increase in highly sulfated KS epitope.

This result suggests that the emergence of more highly sulfated KS GAG in the chick cornea as it matures in ovo is not a driving force correlating with the homogeneous compaction of collagen fibrils in the coalescing collagen bundles and developing stromal lamellae and perhaps points to the influence of other modulating factors that help drive the structural remodeling of the developing corneal stroma in the week before hatching.

Activity of the bicarbonate pump in the corneal endothelium is critical for controlling corneal hydration and thickness and thus for maintaining transparency in the adult eye. 45-47 It is also likely to be important in stromal water regulation and compaction in the developing cornea. However, the precise stage at which the endothelial pump mechanism becomes active is unknown. Histologic investigations have shown that an intact endothelial monolayer is established by E5 to E6.5 Stromal fluid at this time is mostly immobilized by hyaluronic



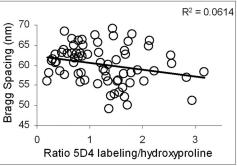


FIGURE 4. Correlation of sulfated KS antibody labeling (relative to BAC-ABC core) per hydroxyproline in relation to collagen Bragg spacing for lesser (*left*, 1B4 reactive) and more highly (*right*, 5D4 reactive) sulfated KS.

acid; later, small leucine-rich PGs, particularly KS PGs most likely influence the water-binding characteristics of the stroma. Conrad et al., 48 studying gene expression associated with chick corneal development, including those genes involved in endothelial transport, found 5- to 10-fold increases in sodiumpotassium ATPase transporter, sodium bicarbonate transporter, and carbonic anhydrase II (CA2), alongside increases in KS gene expression from E9 to E16. CA2 in particular, through increased transport of Na+/HCO3- and Cl- would be expected to drive thinning of the cornea. Perhaps the accumulation of oversulfated KS during this time serves to maintain local collagen hydration, as gross dehydration and compaction progress through the action of the nascent endothelial pump.

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