Affinity Binding of the Cartilage Proteoglycan Protein-Keratan Sulfate Core to Immobilized Hyaluronic Acid

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The protein-keratan sulfate core of bovine nasal cartilage proteoglycan was purified by affinity chromatography on a column of immobilized hyaluronic acid. The hyaluronic acid was immobilized by reaction with a hydrazido-alkyl derivative of Sepharose in the presence of borohydride. Proteoglycan was digested with chondroitinase ABC and the entire mixture was passed over a column of the Sepharose-hyaluronic acid maintained at 4°C. After the digested chondroitin sulfate chains were washed from the column, the bound protein-keratan sulfate core was cluted with 4 M guanidinium chloride. The protein-keratan sulfate core interacts with the affinity matrix through its hyaluronic acid binding site as shown by the inhibition of binding by free hyaluronic acid and hyaluronic acid decasaccharide.

The proteoglycan molecule from bovine nasal cartilage is composed of a protein core of molecular weight 200,000 to which many chondroitin sulfate and keratan sulfate chains are covalently attached at their reducing ends (1,2). In hyaline cartilage, the extracellular proteoglycans exist mainly as large aggregates (3-7). The aggregates dissociate in concentrated solutions of electrolytes thereby allowing their ready extraction from connective tissue (4). Reassociation of the aggregate components takes place *in vitro* when the dissociative solute is removed or sufficiently diluted (4,8). The extracted proteoglycans can be isolated in aggregated or monomeric form by density gradient centrifugation under associative or dissociative conditions, respectively (8,9).

Recently, details of the aggregation process have been elucidated (5,10-13). Hardingham and Muir (10) demonstrated that cartilage proteoglycans interact *in vitro* with HA¹ producing a large increase in hydrodynamic volume as judged by gel chromatography and viscosimetry. This complex appears to consist of many proteoglycan molecules bound at

¹ Abbreviations used: HA, hyaluronic acid; Sepharose-HA, hyaluronic acid covalently attached to adipic acid dihydrazide-substituted Sepharose 2B; GuHCl, guanidinium hydrochloride; core, protein-keratan sulfate core prepared from chondroitinase ABD-digested proteoglycan.

intervals along the HA chain. Each proteoglycan molecule contains one HA-binding site located near one end of the protein core (10,11). Two low molecular weight proteins (link proteins) found in extracts of cartilage stabilize the aggregates (11,12) presumably by interaction with proteoglycan and HA. The features of these interactions are summarized in the model for the proteoglycan-HA-link protein complex proposed by Heinegård and Hascall (13) and by Rosenberg *et al.* (14).

The binding site on the proteoglycan molecule is specific for HA. Cartilage proteoglycans do not interact with polyanions such as dextran sulfate, sodium alginate, DNA, or chondroitin sulfate (10) or chondroitin (11). Furthermore, the binding site exhibits specificity for polysaccharide chain length. The HA decasaccharide, with N-acetylglucosamine at the reducing end, is much more effective than the corresponding octasaccharide in displacing the intact HA chain from the proteoglycan (11,15).

The fact that cartilage proteoglycan interacts specifically with HA makes possible its isolation and purification by affinity chromatography. In this report, we describe the preparation of an affinity matrix comprised of HA covalently attached to agarose beads. The protein-keratan sulfate core, prepared by removal of chondroitin sulfate chains from bovine nasal cartilage proteoglycan by the action of chondroitinase, binds to the immobilized HA and can be eluted with concentrated solutions of GuHCL. Also, immobilized HA can be used as a convenient binding reagent to study the interaction between core and HA. Some of these results have been reported in a preliminary communication (16).

MATERIALS AND METHODS

Materials. Sepharose 4B and Sepharose 2B were obtained from Pharmacia, Inc. (Piscataway, N. J.), sodium cyanoborohydride from Aldrich Chemical Co. (Milwaukee, Wis.), adipic acid dihydrazide from Sigma Chemical Co. (St. Louis, Mo.), purified bovine testicular hyaluronidase (EC 3.2.1.35) from Worthington Biochemicals Corp. (Freehold, N. J.), chondroitinase ABC (EC 4.2.2.4) and crystalline bovine serum albumin from Miles Laboratories (Elkhart, Ind.), and cyanogen bromide from Eastman Organic Chemicals (Rochester, N. Y.). Proteoglycans were isolated from bovine nasal septal cartilage by a procedure described by Oegema et al. (17). The protein-keratan sulfate core of proteoglycan was prepared as described by Hascall and Heinegård (11). Umbilical cord HA (Grade I) was purchased from Sigma Chemical Co., and rooster comb HA was isolated by the method of Swann (18). The molecular weight of the material from umbilical cord, as determined viscosimetrically (19), was 7.8×10^5 and that from rooster comb was 1.6×10^6 . Oligosaccharides were prepared by partially digesting HA with testicular hyaluronidase and isolated by gel chromatography on a standardized column of Sephadex

G-50 (Superfine) according to the method of Hascall and Heinegård (11). Lactoperoxidase (EC 1.11.1.8) was obtained from Worthington Biochemical Corp. (Freehold, N. J.) and was insolubilized according to the method of David and Reisfeld (20). Protein-keratan sulfate core was radioiodinated with insoluble lactoperoxidase by the method described by David and Reisfeld (20) for preparing low specific activity ¹²⁵I-iodinated protein. The specific activity of the preparation was $0.7 \,\mu\text{Ci/}\mu\text{g}$ of protein. Highly purified chondroitin, which was a generous gift of Dr. J. A. Cifonelli (University of Chicago, Chicago, Ill.) was obtained from Dr. I. J. Goldstein, University of Michigan, Ann Arbor, Michigan.

Analytical procedures. Uronic acid and protein were determined by automated methods (21). The amino acid content of proteins was determined by the method of Spackman et al. (22) except that titanous chloride was used as the reducing agent in the ninhydrin reagent. The proteins were hydrolyzed at 110°C in 6 M HCl for 24 h in evacuated tubes. Hexosamines were determined with an automated amino acid analyzer after hydrolysis at 100°C in 4 M HCl for 16 h (23).

Preparation of insolubilized HA. Separose 2B was reacted with cyanogen bromide and coupled with adipic acid dihydrazide by the method described by Wilchek and Miron (24). The hydrazido-Sepharose was added to an equal volume of 0.1 m sodium acetate, pH 5, containing 0.5 mg of rooster comb HA/ml and the mixture was agitated by gentle shaking at 4°C for 16 to 20 h. The reacted Sepharose was washed on a fritted-glass filter with 10 vol of 0.2 m NaCl and 10 vol of water and then was resuspended in an equal volume of 0.1 m sodium acetate, pH 5. Sodium cyanoborohydride (19 mg/ml of Sepharose) was added and the mixture was stirred for 1 h at room temperature. The substituted gel beads were washed on a fritted-glass filter with 20 vol of water, 20 vol of 0.2 m NaCl, and again with water. The washed beads were agitated at 4°C in 4 m GuHCl/0.05 m sodium acetate, pH 5.8, for 16 h before being analyzed for coupled HA content or used for affinity chromatography. The Sepharose-HA was stored at 4°C in 4 m GuHCl/0.05 m sodium acetate, pH 5.8.

Analysis of bound HA. The concentration of the immobilized HA was estimated by determining the glucosamine or uronic acid content of the substituted beads. To determine the glucosamine, Sepharose-HA beads were washed with water on a fritted-glass filter funnel and the excess water was removed by suction. Portions (250 mg) of the moist beads were hydrolyzed in 1 ml of 4 m HCl in sealed tubes at 100° C for 16 h. The hydrolysates were filtered through glass wool and evaporated to dryness in a flash evaporator at 45°C. Water was added to the residue and removed by evaporation several times to remove HCl. The residue was dissolved in 1 ml of water and fractionated on a 0.6×5 cm column of Dowex 50 W-X8 (H⁺). After addition of the sample, the column was

washed with 10 ml of water and eluted with 10 ml of 1 m HCl. The HCl fraction was evaporated to dryness several times and the glucosamine was determined; unsubstituted hydrazido-Sepharose served as the control substance. To determine the uronic acid content, the washed beads (250 mg) were suspended in 1 ml of 0.1 m sodium acetate/0.15 m NaCl, pH 5, containing 350 turbidity-reducing units of hyaluronidase. The mixture was incubated with shaking at 37°C for 16 to 20 h. The beads were sedimented by centrifugation and the uronic acid content of the supernatant solution was determined; again unsubstituted hydrazido-Sepharose served as the control.

Determination of the relative capacity of Sepharose-HA to bind the protein-keratan sulfate core of proteoglycans. Core (1 mg) in 1 ml of 0.5 m GuHCl/0.05 m sodium acetate, pH 5.8, was applied to a 0.6 × 4 cm column of Sepharose-HA previously equilibrated with the same buffer. After washing the column with 4 ml of the equilibration buffer, it was eluted with 5 ml of 4 m GuHCl/0.05 m sodium acetate, pH 5.8. The core content of the eluate was estimated by measuring the uronic acid by a manual carbazole method (25) since solutions containing high salt concentrations cannot be analyzed by the automated procedure. The protein-keratan sulfate core gives a positive carbazole reaction since digestion of proteoglycan with chondroitinase ABC leaves a residual glucuronic acid-containing disaccharide at the nonreducing end of the linkage region saccharides (23).

Preparation of the protein-keratan sulfate core of proteoglycan and isolation by affinity chromatography. Proteoglycan from bovine nasal cartilage (250 mg) contained in 10 ml of 0.1 M Tris-HCl/0.1 M sodium acetate, pH 7.3, was incubated with 5 units of chondroitinase ABC at 37°C for 3 h. The digest was then cooled in ice and placed on a 1.5 × 25 cm column of Sepharose-HA at 4°C. The column was washed with at least 5 column vol of 0.5 M GuHCl/0.05 M sodium acetate, pH 5.8, and then with 5 column vol of 4 M GuHCl/0.05 M sodium acetate, pH 5.8. The effluent was monitored for uronic acid by a manual carbazole method (25). The uronic acid-positive material eluting in the 4 M GuHCl was dialyzed exhaustively against glass-distilled water and freeze-dried.

Binding of the ¹²⁵I-labeled protein –keratan sulfate core to Sepharose – HA. The reaction mixtures contained 40 mg (moist weight) of Sepharose – HA and ¹²⁵I-radioiodinated core (2 × 10⁴ cpm) in 0.5 ml of 0.05 m sodium/phosphate/0.15 m NaCl, pH 7.0, containing 0.01% bovine serum albumin. To determine their capability to displace labeled core from Sepharose–HA, varying amounts of competitors of HA were added in 50 μ l of water. The mixtures were agitated by shaking for varying periods of time. The beads were then sedimented by centrifugation and the radioactivities were determined in aliquots of the supernatant solutions.

RESULTS AND DISCUSSION

Coupling HA to Hydrazido-Sepharose

The effect of reaction conditions on the yield of immobilized HA is summarized in Table 1. The use of sodium cyanoborohydride for the reduction step increased the yield about 70% over that obtained with sodium borohydride. This result was expected since, at pH 5, cyanoborohydride is more stable than borohydride (26). When reacted under the same conditions, about 84% more rooster comb HA than umbilical cord HA was attached to hydrazido-Sepharose 4B but the core-binding capacities of the two materials were similar. The affinity matrix with the highest concentration of bound HA and core-binding capacity was obtained by coupling rooster comb HA to hydrazido-Sepharose 2B using sodium cyanoborohydride as the reducing agent.

The results in Table 1 also show that to bind the HA in a stable manner, it is necessary to add borohydride, but only after the HA and the hydrazido-Sepharose have reacted for a suitable period of time. Evidently, during the initial reaction period, reactive groups on the HA react slowly with the hydrazido-Sepharose to form a product which is stable to washing

TABLE 1

Effects of Reaction Conditions on the Yield of Immobilized Hyaluronic Acid and the Capacity to Bind Protein–Keratan Sulfate Core^a

Type of Sepharose	Hyaluronic acid source	Interaction time before reduction (h)	Reducing agent	Immobilized uronic acid or N-acetylglucosamine (µmol/g) ^b	Core-binding capacity ^c (absorbance at 525 nm)
4B	Umbilical cord	4	NaBH₄	0.069	9.6
4B	Umbilical cord	4	NaCNBH ₃	0.116	11.0
4B	Rooster comb	4	NaBH ₄	0.127	10.3
2B	Rooster comb	4	NaCNBH _a	0.318	16.2
4B	Rooster comb	0	NaBH ₄	0.021	0.0
4B	Rooster comb	4	None	0.005	0.2

^a A mixture containing 2.5 mg of HA and 5 ml (settled volume) of hydrazido-Sepharose in 5 ml of 0.1 M sodium acetate, pH 5.0, was mixed at 4°C for the indicated interaction time. The suspension was filtered and washed with 10 vol of 0.2 M NaCl. For reduction with NaBH₄ the derivatized Sepharose was suspended in 5 ml of water and treated with three 50-mg portions of NaBH₄ over a 1-h period at 25°C. For reduction with NaCNBH₃ the Sepharose was suspended in 5 ml of 0.1 M sodium acetate (pH 5), 50 mg of NaCNBH₃ was added, and the mixture was stirred for 1 h at 25°C. The beads were then washed and analyzed for bound HA and core-binding capacity as described in the text.

^b Micromoles of uronic acid or N-acetylglucosamine per gram of moist Sepharose-HA.

^c Total absorbance units at 525 nm eluting in the 4 м GuHCl solution.

with 0.2 M NaCl and water. After reaction with borohydride, the Sepharose-HA bond is stable in the presence of 4 M GuHCl. The labile intermediate could be a substituted hydrazone formed by the reaction of the aldehyde group at the reducing end of the HA polysaccharide chain with the hydrazide group on the Sepharose. The hydrazone would be reduced to the more stable substituted hydrazine in the presence of borohydride.

Chromatography of Protein-Keratan Sulfate Core on Sepharose-HA

The elution profile shown in Fig. 1 was obtained when 250 mg of proteoglycan was digested with chondroitinase ABC and subjected to chromatography on Sepharose-HA. Approximately 20 mg of material was eluted in the 4 m GuHCl. The yield and amino acid composition are similar to those obtained by other workers for protein-keratan sulfate core purified from chondroitinase digests by exhaustive dialysis followed by gel chromatography and/or alcohol precipitation (1). The material eluting in 4 m GuHCl was further characterized as core by its ability to bind to HA. The affinity-purified core, when subjected to gel permeation chromatography using Sepharose 2B, eluted as a broad peak in the included volume (Fig. 2A). However, if it was combined with rooster comb HA and then fractionated on Sepharose 2B, the uronic acid and protein eluted at the void volume (Fig. 2B) suggesting that the core binds to HA with

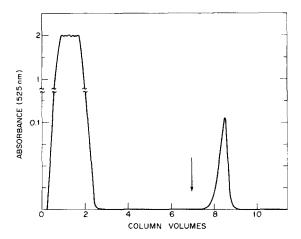


Fig. 1. Chromatography of chondroitinase-digested proteoglycan on Sepharose-HA. Proteoglycan (250 mg) from bovine nasal cartilage was digested with chondroitinase ABC and chromatographed on a 1.5×25 cm (44 ml) column of Sepharose-HA as described under Materials and Methods. The column was washed with 0.5 M GuHCl/0.05 M sodium acetate, pH 5.8. At the point indicated by the arrow, the concentration was increased to 4 M GuHCl. The elution of the column was monitored by hexuronic acid analysis.

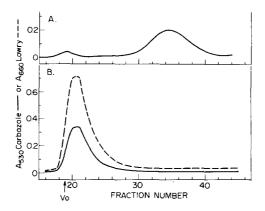


FIG. 2. Gel chromatography of protein-keratan sulfate core and protein-keratan sulfate core-HA complex on Sepharose 2B. A 0.8×135 cm column of Sepharose 2B was eluted with 0.5 M sodium acetate, pH 7.0. Fraction volumes were 1.35 ml. The elution of hexuronic acid (——) was monitored by the automated carbazole assay and protein (---) by the automated Lowry assay. (A) Affinity purified core, 3.1 mg, in 1 ml of 0.5 M sodium acetate, pH 7.0, was chromatographed. (B) Affinity purified core, 2.4 mg, and 100 μg of HA in 1 ml of 0.5 M sodium acetate pH 7.0, were incubated at 25° C for 1 h and then chromatographed.

a resultant increase in hydrodynamic volume. It should be noted that all of the affinity-purified core binds to free HA. In contrast, only 70% of the core isolated by gel chromatography interacts with HA (11). Inactive core molecules evidently do not bind to the affinity matrix. Thus, it appears that the chromatographic isolation of protein-keratan sulfate core with Sepharose-HA is dependent on the specific interaction through the HA-binding site and is not the result of nonspecific binding to the anionic groups of the HA-substituted beads.

Only a small fraction of undigested proteoglycan binds to Sepharose—HA. This material, which interacts with the affinity matrix, is believed to consist of small proteoglycans² similar to those described by Heinegård (27). These results suggest that the majority of the proteoglycan molecules are too bulky to interact with the immobilized HA.

Binding of ¹²⁵I-Labeled Protein-Keratan Sulfate Core to Sepharose-HA

To further assess the specificity of the interaction between core and immobilized HA, core was labeled with ¹²⁵I and its binding to Sepharose–HA was studied under equilibrium conditions. As shown in Fig. 3, ¹²⁵I-labeled core bound to Sepharose–HA rapidly, reaching 85% of the

² J. E. Christner, M. L. Brown, and D. D. Dziewiatkowski, unpublished observations.

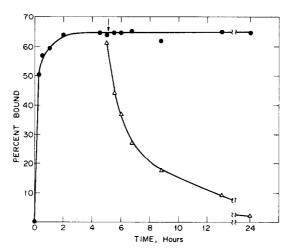


FIG. 3. Binding of the 125 I-labeled protein-keratan sulfate core to Sepharose-HA and displacement as a function of time. The reaction mixtures contained 40 mg of Sepharose-HA (containing 4 μ g of bound HA) and 125 I-labeled core (2 × 10⁴ cpm) in 0.5 ml of 0.05 m sodium phosphate/0.15 m NaCl, pH 7.0, containing 0.01% bovine serum albumin. The suspensions were agitated by shaking at 25°C for the indicated time periods and the extent of binding was determined as given in the text. At the time point indicated by the arrow, 25 μ l of H₂O (\bullet) or 22.5 μ g of HA dodecasaccharide (\triangle) dissolved in 25 μ l of H₂O was added to separate incubation mixtures. The extent of binding was determined at the indicated time intervals.

maximum level in 15 min. In contrast, the displacement of the core by HA dodecasaccharide was much slower. These results are similar to those obtained by Hascall and Heinegård (11) for the binding of proteoglycan to HA and its displacement by HA oligosaccharides as measured by the viscosimetric method.

The fraction of 125 I label bound to Sepharose–HA decreased with increasing total core concentration (Fig. 4). These results indicate that unlabeled core competes with its labeled counterpart for binding to immobilized HA. The inset of Fig. 4 displays the data in reciprocal form from which the binding capacity and binding constant can be derived. Each core molecule occupies 8 to 10 disaccharide units along the HA polysaccharide chain (11). The expected binding capacity of 1 g of substituted beads containing 0.21 μ mol of covalently coupled uronic acid is approximately 0.02 μ mol or 9 mg of core of molecular weight 4.5 × 10⁵ (1). This value is only threefold greater than that obtained experimentally. The relatively high capacity and binding constant values suggest that the immobilized HA is readily accessible to the core molecules in solution.

The ¹²⁵I-labeled core also could be displaced from the affinity matrix by HA deca- and dodecasaccharides (Fig. 5). When HA at a concentra-

tion equivalent to 2 μ M uronic acid was added to the reaction mixture which contained 20 µM insolubilized uronic acid, the core binding was decreased to 50% of the control value. Taking into account that about 30% of the immobilized polysaccharide is accessible to the solution, the results indicate that HA in the soluble and insoluble phases has similar affinities for the HA-binding site. Octasaccharide at a high concentration (25 µm uronic acid) inhibited the binding only 8% while chondroitin had little or no effect (Fig. 5). Hardingham and Muir (15) and Hascall and Heinegård (11) previously demonstrated that HA deca- and dodecasaccharides inhibited the interaction between proteoglycan and HA while the octasaccharide did not. Furthermore, Hascall and Heinegård (11) showed that the inhibition of core-HA interaction as detected by gel permeation chromatography exhibited a similar specificity and that chondroitin was ineffective in competing for the HA-binding site. In other experiments it was shown that HA decasaccharide, which was modified by esterification of carboxyl groups on the glucuronic acid residues, did not displace the ¹²⁵I-labeled core. The elimination of competition by blockage of the oligosaccharide carboxyl groups is consistent with pre-

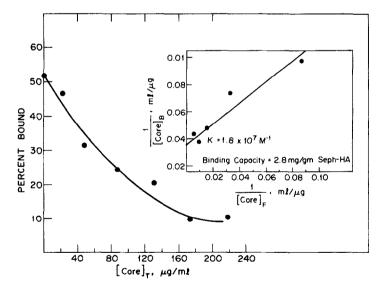


FIG. 4. Displacement of the ¹²⁵I-labeled protein-keratan sulfate core from Sepharose-HA by unlabeled protein-keratan sulfate core. The reaction mixtures contained ¹²⁵I-labeled core (1.8 × 10⁴ cpm), the indicated amounts of unlabeled core, and 5 mg of Sepharose-HA (containing 0.5 μ g of HA) in a total volume of 0.5 ml of 0.05 m sodium phosphate/0.15 m NaCl, pH 7.0, containing 0.01% bovine serum albumin. The mixtures were agitated at 25°C for 16 h and the extent of binding was determined as described in the text. [Core]_T: Total concentration of core; [Core]_B: concentration of core bound to Sepharose-HA; [Core]_F: concentration of free core.

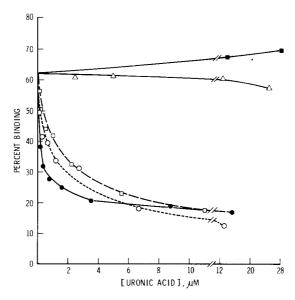


Fig. 5. Displacement of ¹²⁵I-labeled protein–keratan sulfate core from Sepharose–HA by HA, oligosaccharides of HA, and chondroitin. The reaction mixtures contained 40 mg of Sepharose–HA (containing 11.7 nmol of bound uronic acid), ¹²⁵I-labeled core (1.8 × 10⁴ cpm), and the indicated amounts of poly- and oligosaccharides in 0.55 ml of 0.05 M sodium phosphate/0.15 M NaCl, pH 7.0, containing 0.01% bovine serum albumin. The mixtures were agitated by shaking for 4 h and the extent of binding was determined as described in the text. The polysaccharides used were HA (○); HA dodecasaccharide (●), decasaccharide (□), and octasaccharide (△); and chondroitin (■).

vious results (28) which showed that the methyl ester derivative of HA decasaccharide did not inhibit proteoglycan-HA interaction. The strict dependence of the inhibition on oligosaccharide chain length and composition strongly supports the specificity of the interaction between protein-keratan sulfate core and Sepharose-HA.

The current model assumes that cartilage proteoglycan aggregates are stabilized by link proteins which interact with both the HA and the proteoglycan (11-14). Although interaction between link proteins and HA has been demonstrated (11,29), nothing is known about the requirements of polysaccharide chain size and composition for the complex formation. The competitive binding studies demonstrated that the HA-substituted beads are useful for measuring the affinity and specificity of the interaction of protein core and HA. Presumably, they also could be used to elucidate the details of the link protein-HA aggregation process.

Hardingham and Adams (30) recently developed a sensitive assay for the determination of hyaluronic acid in cartilage. This assay, which requires $1-20 \mu g$ of HA, is dependent on the formation of large aggregates between the HA and cartilage proteoglycans which are detected by gel permeation chromatography. Using the competitive binding method described in the present report, 0.01 μg of HA could be measured by a relatively simple procedure.

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