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Role of inter- α -inhibitor and its related proteins in urolithiasis. Purification of an inter- α -inhibitor related protein from the bovine kidney

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Abstract Urine contains several macromolecules that inhibit calcium oxalate (CaOx) crystallization. Among them is bikunin, the light chain of most of the inter- α -inhibitor (I α I) family of glycoproteins. This study aimed to verify whether bikunin and other members of the I α I family are synthesized in the kidneys or derived exclusively from the plasma. Proteins extracted from homogenized bovine kidney were applied successively to three chromatographic steps on DEAE-Sephacel, Sephacryl S-300, and Mono Q column. The inhibitory activity was assayed using a CaOx crystallization system. The presence of I α I-related proteins was determined by electrophoresis and Western blotting. The results showed that kidney extract contained a 125-kDa protein that cross-reacted with anti-I α I antibodies. This protein inhibited CaOx crystallization efficiently. According to its molecular weight and immunoreaction with anti-I α I antibody, the 125-kDa protein could be pre- α -inhibitor. The latter is known to encompass a heavy chain and bikunin, which may explain its inhibitory activity against CaOx crystallization. Consequently, we hypothesize that kidneys may produce some I α I-related proteins that are involved in the inhibition of stone formation.

Key words Nephrolithiasis · Bovine kidney · Calcium oxalate crystallization · Inter- α -inhibitor · Bikunin · Chromatography

Introduction

Urine is normally metastable with respect to calcium oxalate (CaOx), a common constituent of most human kidney stones. Thus under appropriate conditions CaOx may readily precipitate in the urine and lead to stone formation in the urinary tract. Fortunately, the presence of efficient inhibitory macromolecules, reduces the likelihood of crystallization in the urine [8, 11, 13, 16, 17, 24, 27]. The past 15 years have seen a tremendous increase in research in this area and a substantial amount of information is currently available about purification, characterization, and identification of inhibitory macromolecules [1, 2, 10, 12, 19, 27]. We identified one such molecule, uronic acid-rich protein (UAP), a glycoprotein with a molecular weight of 35 kDa in human [1, 3] and rat urine [3]. In these studies, we have shown that UAP is a potent inhibitor of CaOx crystallization. Interestingly, both human and rat UAP exhibited structural similarity to bikunin, a subunit of inter- α -inhibitor (I α I) [3,5]. Later, we confirmed that UAP is bikunin [4]. Moreover, we have demonstrated that urine contains two bikunins, one of which is a potent inhibitor of CaOx crystallization. The differences between the two urinary bikunins may be due to their origin. We hypothesized that one form of bikunin may derive from plasma and the other from the kidneys. The aim of this study was to purify inter- α -inhibitor-related proteins from bovine kidney and test their inhibitory activity on CaOx crystallization.

Materials and Methods

Materials

Bovine kidneys were obtained from the Animal Facility of the Veterinary College at the University of Florida (Gainesville, FL). During transportation to the laboratory, kidneys were kept on ice and then processed immediately on arrival as described in the Method section.

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Diethylaminoethyl (DEAE)-Sephacel, Sephacryl S-300, and Mono Q HR 5/5 column were purchased from Pharmacia Biotechnology, N.J. Before each use, gels or columns were recycled according to the manufacturer's recommendations. Salt concentration was measured by conductivity meter (Fisher Scientific, Pa). Dialysis tubing with a molecular weight cut off of 6000–8000 Da was obtained from Spectrapor (Spectrum Medical Industries Calif.). Normal human serum inter- α -inhibitor polyclonal antibody (Code AXL 424) was purchased from Accurate Chemical and Scientific, N.Y. [45 Ca] Calcium Chloride was purchased from Amersham Corporation, IL. Electrophoresis, transfer blot apparatus, low pre-stained molecular weight markers, and reagents for immunostaining were obtained from Bio-Rad Laboratories, Calif. Nitrocellulose membranes were purchased from Millipore Bedford, USA and the second antibody was from Hyclone Laboratories, Utah. All reagent grade chemicals were acquired from Fisher Scientific.

Methods

Kidney tissue preparation

Part of bovine kidney (256 g) was sliced into small pieces after removing fatty tissue and washed several times with TRIS buffer (TRIS 0.05 M, NaCl 0.1 M, pH 7.3) to remove any blood contaminant. Then, the tissue was homogenized in a blender with 500 ml of TRIS buffer and centrifuged at 8000 rpm for 20 min. The pellet obtained was resuspended in 300 ml of TRIS buffer, homogenized again, and centrifuged. This procedure was repeated three times. All supernatants were combined and heated to 65°C in a boiling-water bath with stirring. After the solution reached 65°C, it was immediately cooled on ice and centrifuged at 8000 rpm for 20 min. The supernatant was dialyzed overnight against demineralized water at 4°C. The dialysate obtained was adjusted to pH 7.3 and NaCl 0.1 M by adding NaOH and solid NaCl, respectively and then processed for protein purification as described below.

Protein isolation

The procedure used to purify proteins from bovine kidney extract was similar to that used for human urine as previously described [1]. Briefly, the sample was incubated with DEAE-Sephacel gel pre-equilibrated with TRIS 0.05 M, NaCl 0.1 M, pH 7.3 for 30 min with stirring. The suspension was filtered, washed with equilibrated buffer, and eluted successively with 300 and 200 ml of TRIS buffer. The combined eluents were dialyzed overnight against deionized water. Then, the sample was adjusted to pH 7.3 and NaCl 0.1 M and injected to the DEAE-Sephacel column. Proteins were eluted with a linear gradient of NaCl solution from 0.1 to 0.5 M in TRIS buffer. The fractions containing I α I-related proteins were dialyzed, lyophilized, and injected into the Sephacryl S-300 column. The final purification step is the use of fast protein liquid chromatography (FPLC) chromatography on Mono Q HR 5/5 column using a linear gradient of NaCl solution from 0.1 to 0.5 M buffered with TRIS 0.05 M, pH 7.3.

Protein and uronic acid assay

After each chromatographic step, protein and uronic acid concentrations were determined in each fifth tube using Lowry's method [26] and carbazole reaction [9], respectively. According to the method used, albumin and glucuronic acid were used as standards.

Inhibition assay

To determine the inhibitory activity in the fractions, a calcium oxalate crystallization system was used as described in a previous paper [3]. The inhibition assay involves using a mixture of 1 ml

solution of calcium chloride (2 mM in TRIS 0.05 M, NaCl 0.15 M, pH 7.3) in the presence of [45 Ca] calcium chloride and 1 ml ammonium oxalate (2 mM in TRIS 0.05 M, NaCl 0.15 M, pH 7.3). The tubes containing proteins, lyophilized from 100- μ l samples, were compared with those without proteins. At the end of the assay, tubes were centrifuged at 2000 g for 5 min and 0.5 ml of supernatant was withdrawn for radioactivity determination in an LS 7500 liquid scintillation counter (Beckman Instruments, Calif.).

Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and immunoblotting

Electrophoresis was performed on 1 mm thick 10% polyacrylamide gel by using Bio Rad Mini Proteinase II apparatus. Proteins and standards were electrophoresed after they were subjected to the reduction in the presence of SDS and β -mercaptoethanol. The protein bands were revealed by staining with Coomassie blue R-250 or transferred onto nitrocellulose membrane for immunological identification.

Gels and nitrocellulose membranes were pre-equilibrated with transfer buffer (TRIS 25 mM, glycine 192 mM, methanol 200 ml). Proteins were transferred by using the same buffer at 100 volts for 1 h at 4°C. The immunochemical staining steps were performed at room temperature by using polyclonal I α I antisera at the dilution of 1:1000 as primary antibody according to the procedure described by Towbin et al. [34]. The second antibody conjugated to alkaline phosphatase was used at the dilution of 1:5000. The staining was developed by using an alkaline phosphatase kit.

Results and discussion

It is now widely recognized that urine contains several macromolecules that prevent, reduce or modulate crystallization [1, 2, 10, 12, 19, 27]. Among them, bikunin, the light chain of the I α I-related family of proteins, with a molecular weight of 35 kDa, is a potent inhibitor of CaOx crystallization [2–5]. Based on our observations that urine contains two bikunins with slightly different crystallization inhibitory activity and molecular weights, we hypothesized that kidneys may produce glycoproteins related to I α I [4]. Since then we have demonstrated that bikunin as well as the two heavy chains of I α I, namely H1 and H2, are found in the matrices of human CaOx kidney stones and are also included in CaOx crystals experimentally induced in the human urine [6]. Recent results from our laboratory have shown additionally that rat kidneys as well as LLC-PK $_1$ and MDCK cells in culture, both express bikunin m-RNA [7, 20]. It is interesting then to purify these proteins from the kidneys and test their inhibitory activity against CaOx crystal formation.

The first step in the preparation of protein extract from bovine kidney is the homogenization of the tissue. In order to avoid blood contaminants, the kidney was sliced in tiny pieces and washed several times with TRIS buffer. The protein extract obtained as described in the Materials and Methods section was processed for protein purification procedure using three chromatographic steps. The sample was subjected to a DEAE-Sephacel bath and column chromatography using a linear gradient of NaCl solution from 0.1 to 0.5 M in TRIS 0.05 M, pH 7.3. The elution profile is shown in the Fig. 1. The

first half of the fractions (1 to 60) are rich in proteins. The CaOx crystallization inhibitory activity increases rapidly in the first 10 fractions and becomes stable between 70% and 80% in the remaining fractions. The electrophoresis followed by Western blot analysis showed that fractions 30 to 70 (fraction A) contain a protein with a molecular weight of about 125 kDa which cross-reacted with anti-I α I antibodies (Fig. 2). This evidence, by itself, cannot substantiate the real presence of I α I-related proteins in the kidney. Contamination from plasma cannot be ruled out even though the precautions and controls adopted during the preparation procedure seem to exclude this event. The same conclusion has been drawn by other researchers [14].

The second chromatography step was to submit the fraction "A" to the Sephacryl S-300 column. A typical elution profile is illustrated in Fig. 3. The electrophoresis followed by Western blot analysis showed that fractions 80 to 110 contain a 125-kDa protein which immunoreacted with anti-I α I antibodies (Fig. 4). Fractions containing this protein (fraction B) were subjected to the FPLC chromatography on Mono Q column for a final purification. After elution, samples from the fractions were subjected to gel electrophoresis to evaluate their

protein content. According to the result of this analysis, the fractions were pooled into three fractions named "a", "b" and "c". These latter were then assayed on our calcium oxalate crystallization system using different concentrations from 2.5 to 20 μ g/ml. The results are summarized in Table 1. As it can be seen, fraction "c" exhibits a strong inhibitory activity compared with "a" and "b" fractions. Electrophoresis and Western blot analysis confirmed the presence of the 125-kDa protein in fraction "c" which cross-reacted with anti-I α I antibodies (not shown). According to its molecular weight and its positive immunoreaction with I α I antibodies, the 125-kDa protein could be the pre- α -inhibitor (P α I). P α I contains two chains: heavy chain H3 and bikunin [15, 28]. According to a new nomenclature proposed by Salier et al. [32], P α I is expressed by I α IH3.B. However, presence of the H3 chain has to be proven before the 125-kDa I α I-related protein obtained from bovine kidneys can be confirmed as P α I.

Figs. 1 and 2 illustrate the elution profiles, protein contents, uronic acid contents, and inhibitory activity of various fractions during chromatographic separation of the protein. It is clear that the inhibitory activity does not fully coincide with either the proteins or the uronic

Fig. 1 DEAE-Sephacel ion exchange chromatography of proteins extracted from homogenized bovine kidney. The column was eluted by a linear gradient of NaCl solution from 0.1 to 0.5 M in TRIS 0.05 M, pH 7.3. Protein and uronic acid concentration were determined according to Lowry's method and the carbazole reaction respectively. Salt concentration was measured by a conductivity meter. Inhibition assay of fractions was assayed on a calcium oxalate crystallization system. Fraction "A" was collected for further chromatographic purification

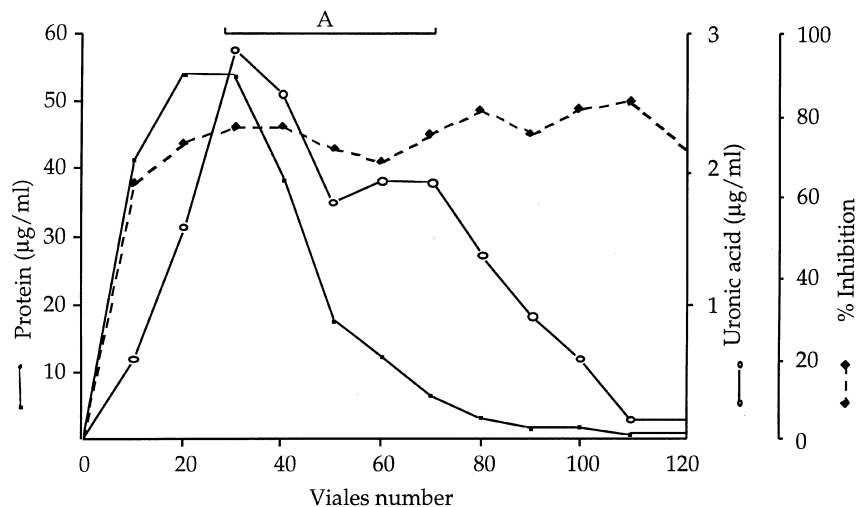


Fig. 2 Western bolt analysis using inter- α -inhibitor antibody of each tenth fraction obtained from the DEAE-Sephacel column. Molecular weight markers are shown on the left

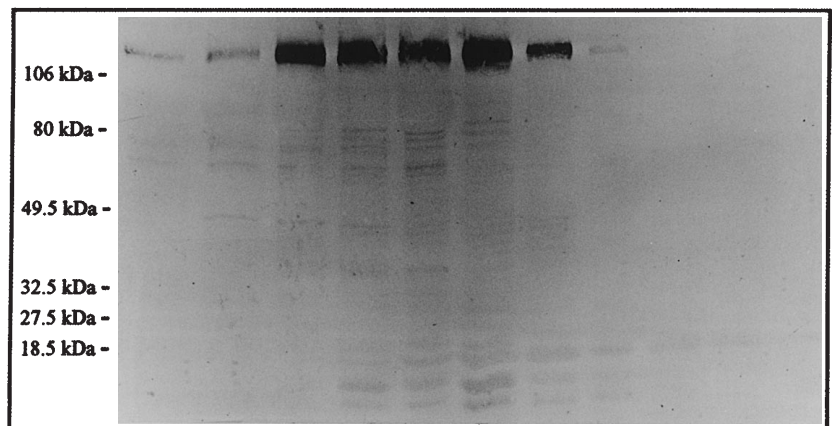


Fig. 3 Fraction "A" obtained from DEAE-Sephacel chromatography was eluted on a Sephacryl S-300 column using TRIS 0.05 M, NaCl 0.1 M, pH 7.3 as eluent. Protein concentration, uronic acid content, and inhibition activity were determined as in Fig. 1. Fraction "B" was taken for a further chromatographic step

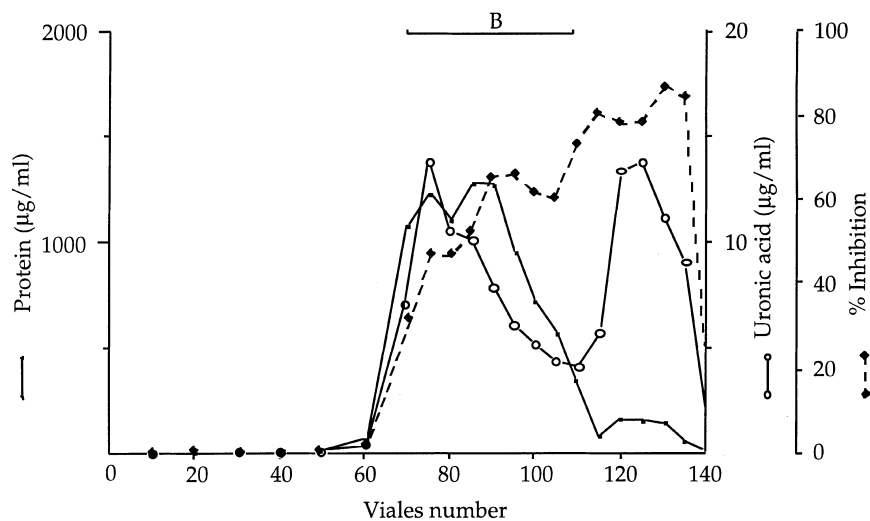


Fig. 4 Western bolt analysis by using inter- α -inhibitor antibody of each tenth fraction (from 70 to 140) obtained from Sephacryl S-300 column. Molecular weight markers are shown on the left

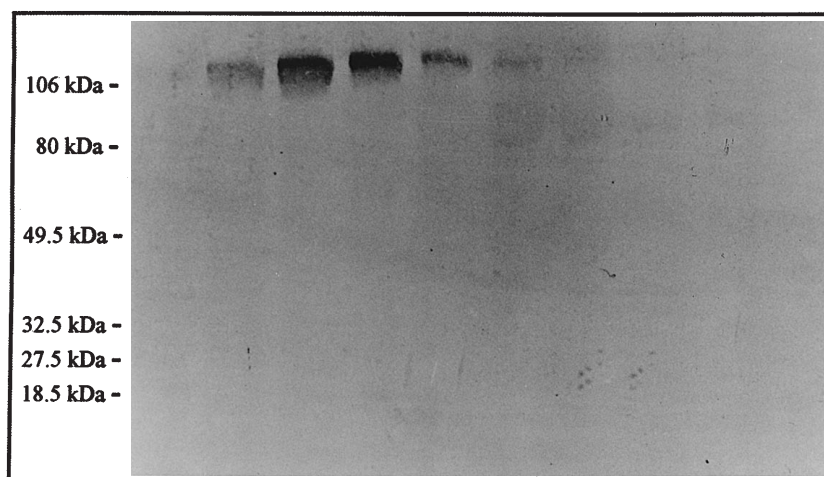


Table 1 Calcium oxalate crystal growth inhibitory activity of fractions obtained from the Mono Q column. The results are expressed as the percentage of calcium oxalate growth inhibited

Fraction	Concentration (µg/ml)			
	2.5	5	10	20
a	4.0	15.5	20.0	27.6
b	7.3	26.7	52.4	70.1
c	20.9	46.6	62.4	74.2

acid contents, indicating that substances other than proteins detectable by Lowry's method or macromolecules containing uronic acid may participate in inhibition of crystallization. It is also possible that these macromolecules are active at very low concentrations and no change in the activity is noticed after a certain level is reached. It is clear that fractions A and B contained the 125-kDa I α I-related protein which is a potent inhibitor of CaOx crystallization. The 125-kDa protein itself can be a modulator of CaOx crystallization or its light chain, bikunin may be responsible for the inhibi-

tory activity against CaOx crystallization since we have demonstrated in several studies that bikunin is a potent inhibitor of CaOx crystal formation [1–4].

It is generally recognized that I α I-related proteins are synthesized in the liver [29, 31]. In addition Kastern et al. [21] have shown that α 1-microglobulin mRNA levels, studied by an RNA dot blot assay, were high in the liver and kidney of rats. This finding leads to the hypothesis that bikunin is also produced by the kidney since α 1-microglobulin and bikunin are synthesized by a common mRNA precursor [22, 25]. Moreover, in some pathological conditions such as inflammation, an increase in bikunin synthesis occurs in the absence of proteolytically induced breakdown of I α I complexes [18, 30]. Recent results from our laboratory have shown an expression of bikunin mRNA in rat kidneys as well as LLC-PK₁ and MDCK cells in culture [7, 20]. Presence of the 125-kDa protein described here and most other I α I-related proteins, in the kidneys and urine, indicate that these proteins are also produced by the kidneys, since under normal conditions they would be unable to pass through the glomerular sieves. Results of our recent studies de-

scribed here and elsewhere suggest that various α I-related proteins can be produced by renal cells and some of them may be involved in modulation of crystallization within the nephron.

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