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United States Patent [19]

[11] E

Patent Number: Re. 35,694**Seyedin et al.**[45] **Reissued Date of Patent: Dec. 16, 1997**[54] **POLYPEPTIDE CARTILAGE-INDUCING FACTORS FOUND IN BONE**[75] **Inventors: Saeid Seyedin, Saratoga, Calif.; Thomas Thomas, West Roxbury, Mass.; Hanne Bentz, Newark, Calif.; Larry Ellingsworth, San Jose, Calif.; Rosa Armstrong, Palo Alto, Calif.**[73] **Assignee: Celtrix Pharmaceuticals, Inc., Santa Clara, Calif.**[21] **Appl. No.: 257,472**[22] **Filed: Jun. 9, 1994****Related U.S. Patent Documents**

Reissue of:

[64] **Patent No.: 4,843,063**
Issued: Jun. 27, 1989
Appl. No.: 204,173
Filed: Jun. 8, 1988

U.S. Applications:

[63] **Continuation of Ser. No. 664,766, Mar. 5, 1991, abandoned, which is a continuation of Ser. No. 129,864, Dec. 10, 1987, Pat. No. 4,774,322, which is a continuation of Ser. No. 767,144, Aug. 19, 1985, abandoned, which is a continuation-in-part of Ser. No. 630,938, Jul. 16, 1984, abandoned.**[51] **Int. Cl.⁶ C07G 7/00; A61K 9/00**[52] **U.S. Cl. 514/2; 514/12; 514/21; 530/324; 530/841; 424/85.1**[58] **Field of Search 514/12, 21, 2; 424/85.1; 530/324, 841**[56] **References Cited****U.S. PATENT DOCUMENTS**3,458,397 7/1969 Meyers et al. .
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Primary Examiner—Nathan M. Nutter*Attorney, Agent, or Firm*—Morrison & Foerster LLP[57] **ABSTRACT**Two proteins that are found in bone and that have in vivo chondrogenic/osteogenic activity in combination with a co-factor are described. Both proteins also were active in combination with EGF in the in vitro TGF- β assay. Each has a molecular weight of approximately 26,000 daltons by SDS-PAGE. Each is reduced to a single polypeptide indicating that the proteins are probably homodimers. One has an N-terminal sequence identical to that of human placenta-derived TGF- β whereas the other has an N-terminal sequence that is different from that of TGF- β derived from human placenta. The two proteins may be purified to homogeneity using RP-HPLC or acetic acid-urea gel electrophoresis.**12 Claims, 6 Drawing Sheets**

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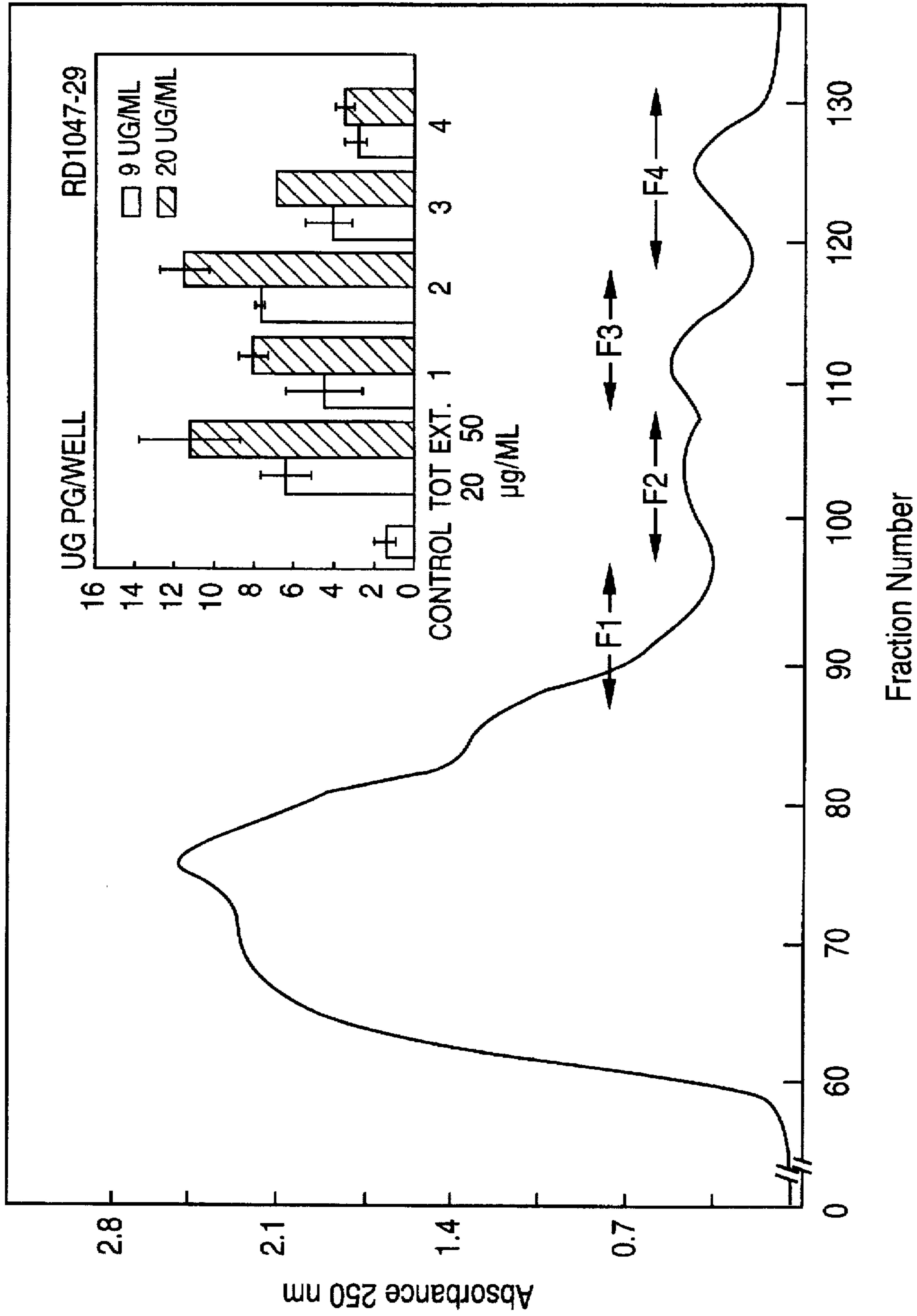


FIG. 1

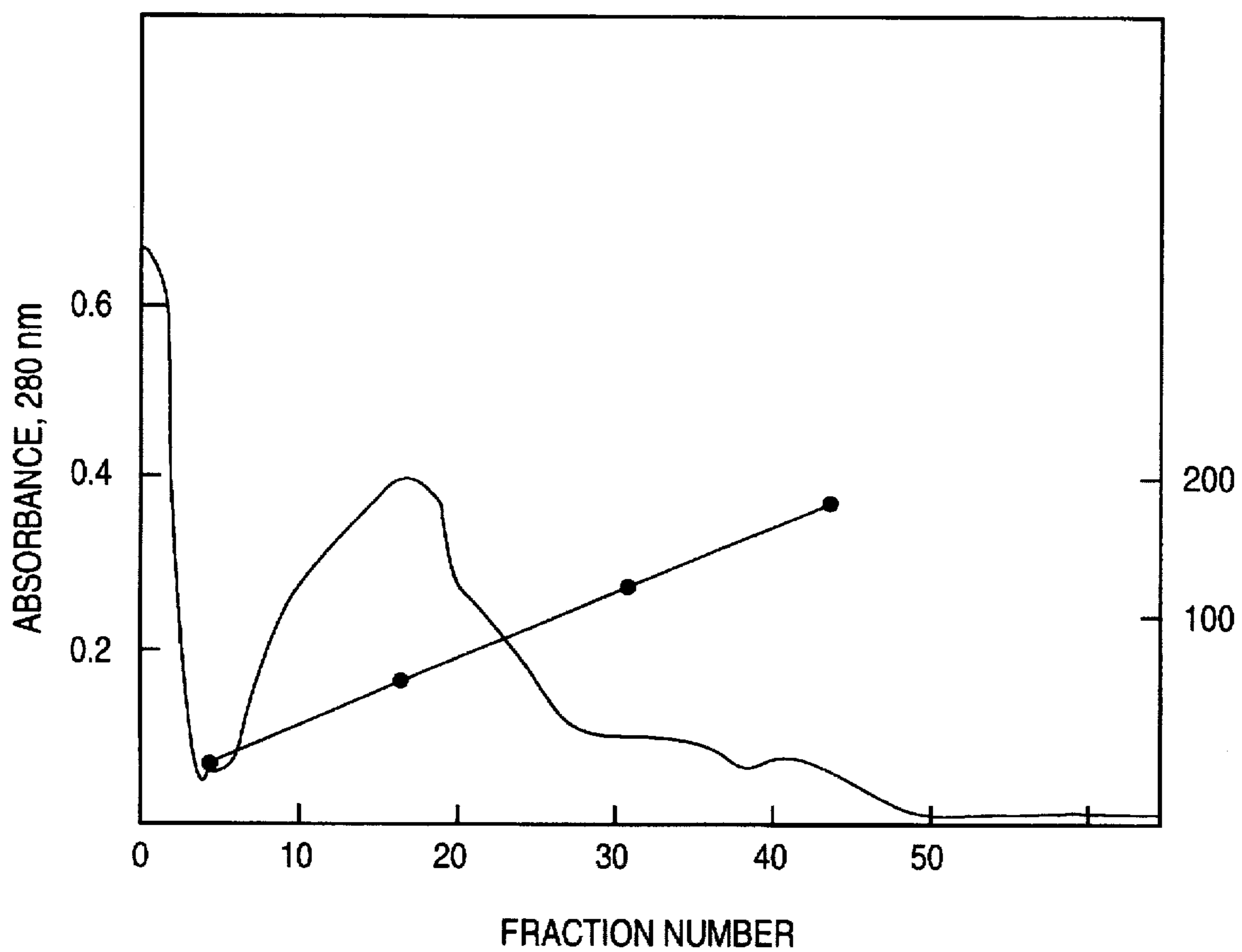


FIG. 2

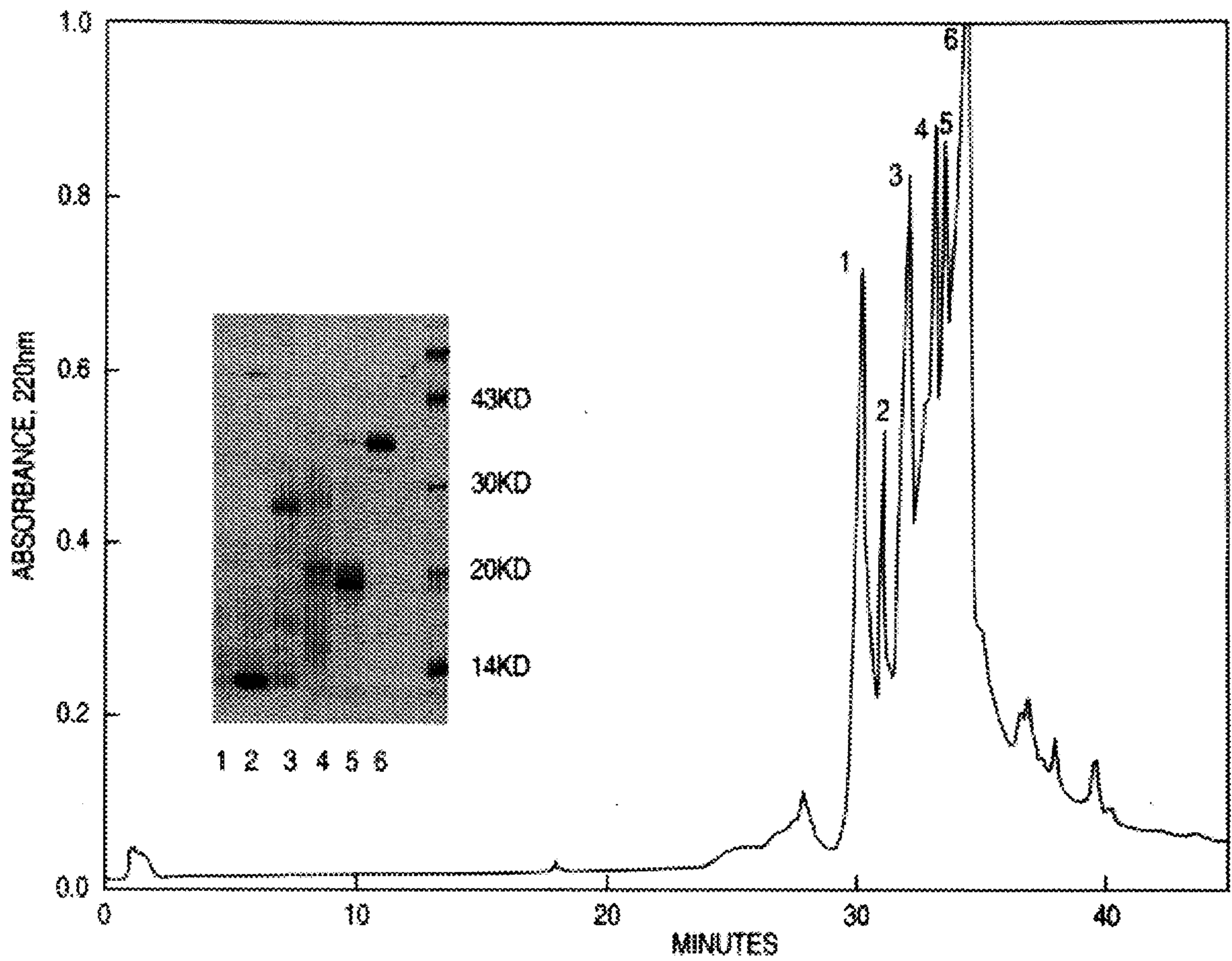


FIG. 3

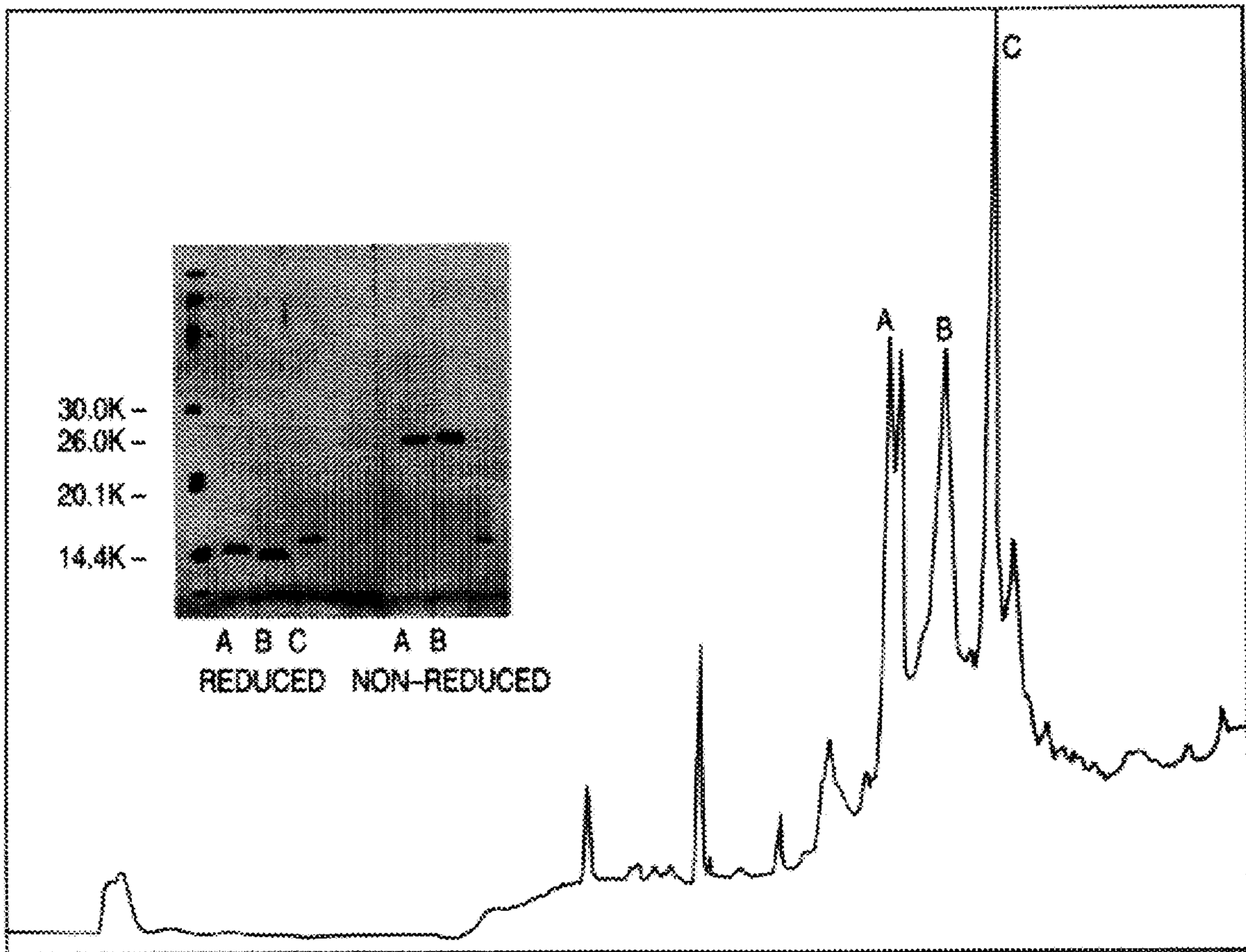


FIG. 4

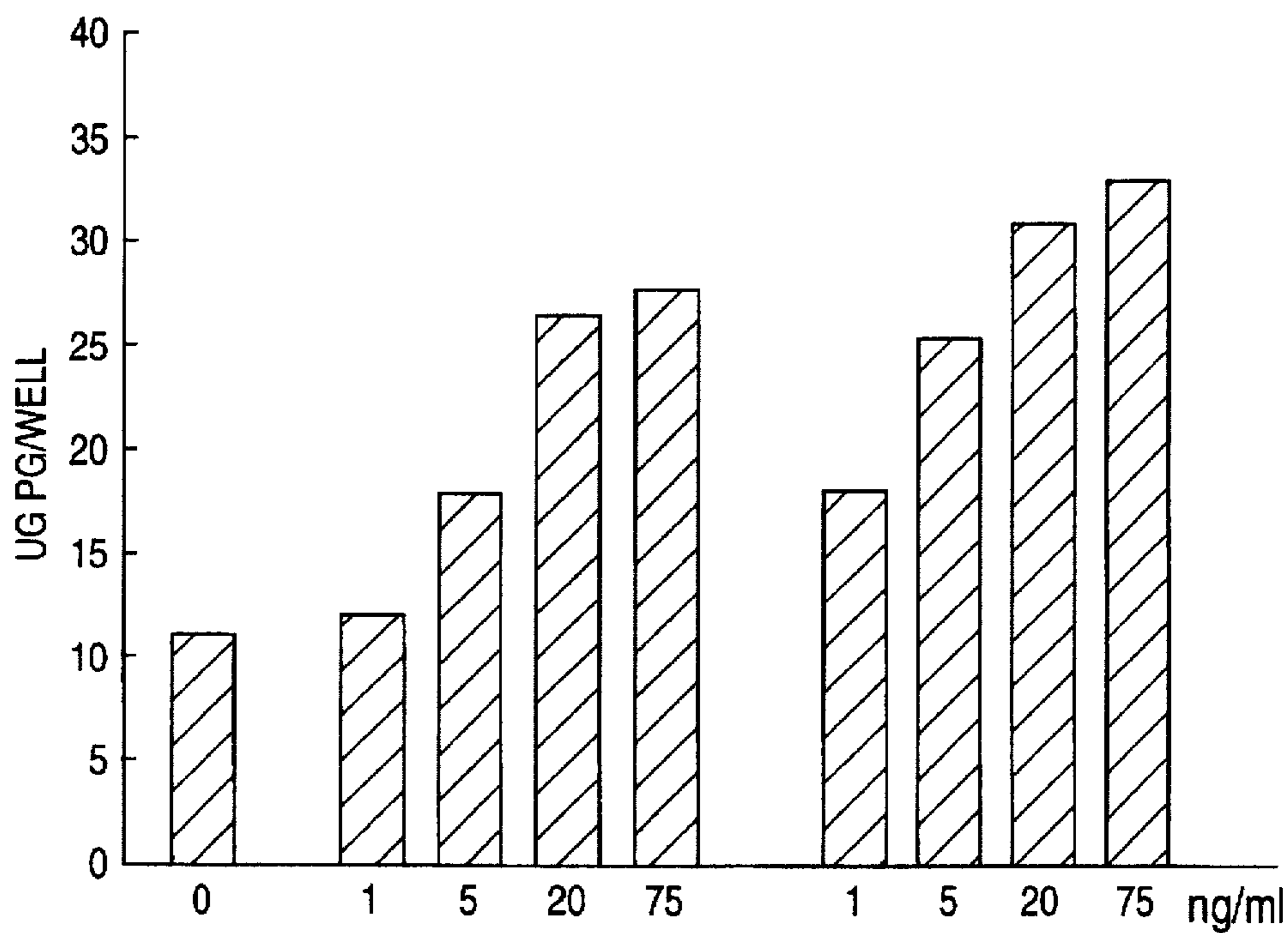


FIG. 5

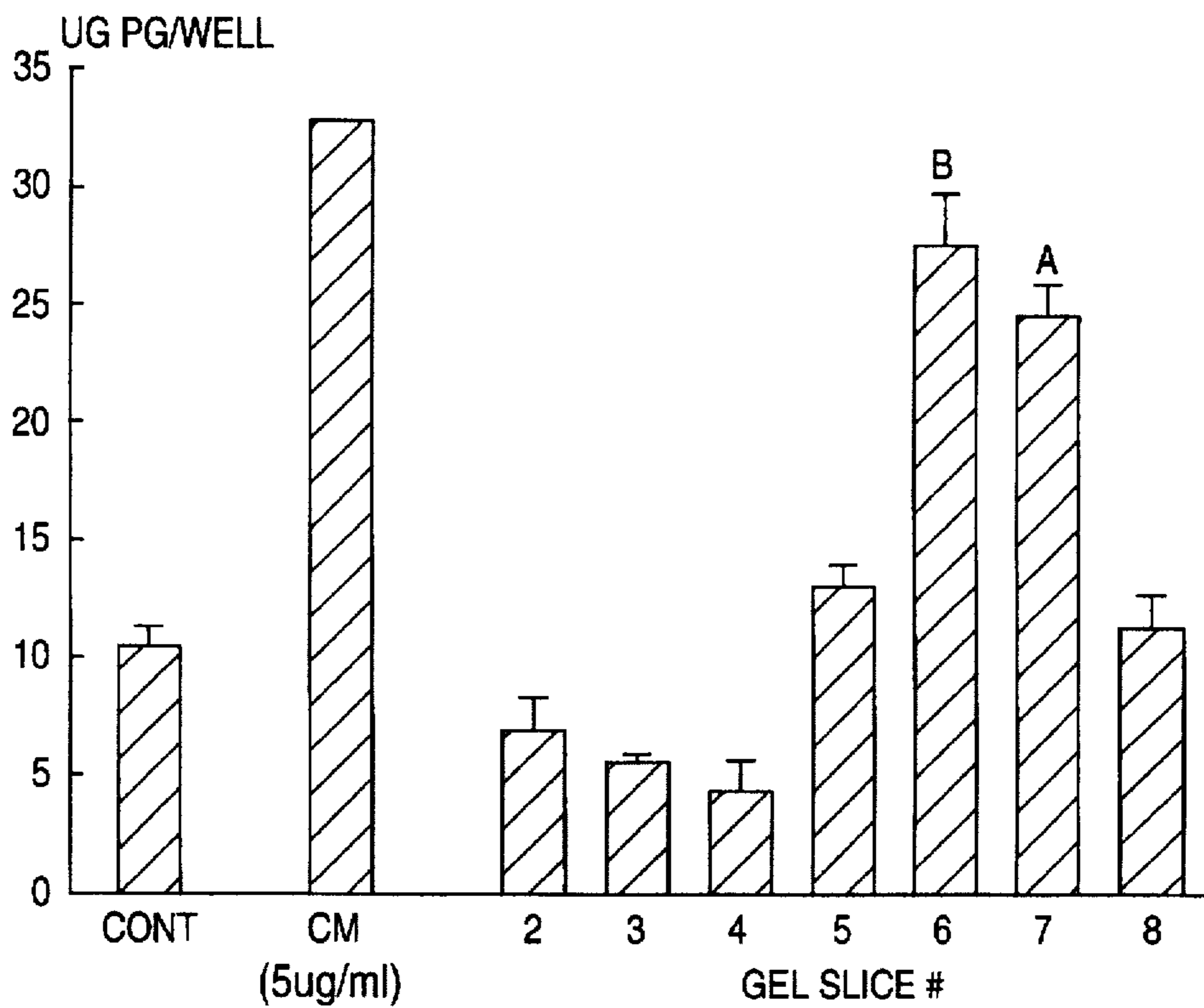


FIG. 6

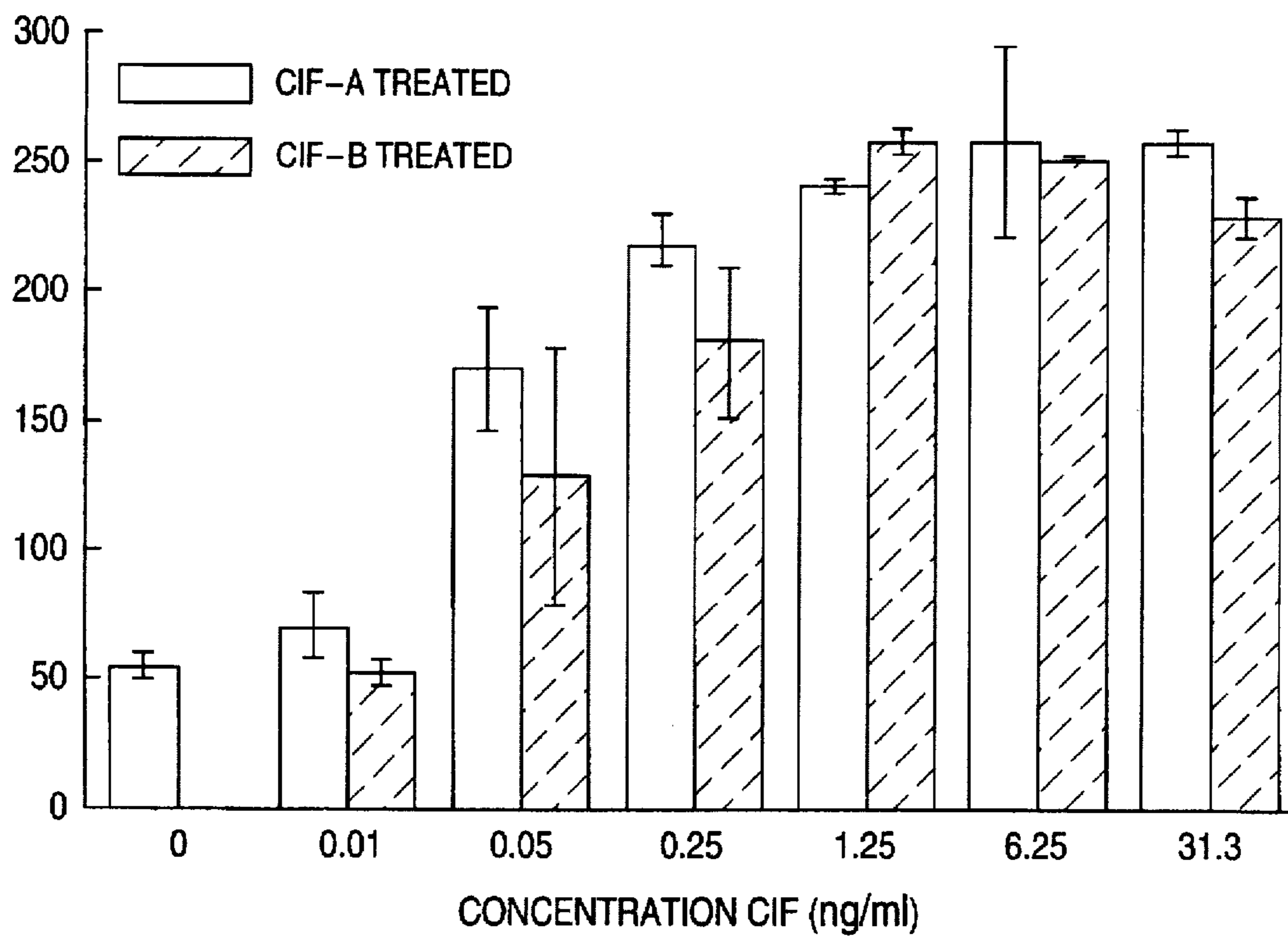


FIG. 7

POLYPEPTIDE CARTILAGE-INDUCING FACTORS FOUND IN BONE

Matter enclosed in heavy brackets [] appears in the original patent but forms no part of this reissue specification; matter printed in italics indicates the additions made by reissue.

CROSS-REFERENCE TO RELATED APPLICATION

This application is a continuation of application Ser. No. 07/664,766, filed Mar. 5, 1991 abandoned which is a RE of 07/204,173, filed Jun. 8, 1988, now U.S. Pat. No. 4,843,063, which is a continuation of copending U.S. Ser. No. 129,864, filed Dec. 10, 1987, now U.S. Pat. No. 4,774,322, issued Sept. 27, 1988, which is a continuation of U.S. Ser. No. 767,144, filed Aug. 19, 1985, now abandoned, which is a continuation-in-part of U.S. Ser. No. 630,938, filed July 16, 1984, now abandoned.

DESCRIPTION

1. Technical Field

The present invention relates to protein chemistry. More particularly, it relates to polypeptides that are involved in cartilage induction in vivo. These polypeptides are sometimes referred to as cartilage-inducing factors (CIFs) herein.

2. Background

Parent U.S. patent application Ser. No. 630,938 describes two proteins that were isolated from bovine bone. These proteins exhibited in vitro chondrogenic activity. Further characterization and testing of these two proteins revealed that one or them may be the same polypeptide as a human placenta-derived polypeptide called "beta type transforming growth factor" (TGF- β) and described in International patent application PCT/US83/01460 published Mar. 29, 1984 under No. WO 84/01106 and EPA No. 84450016.5 published Dec. 19, 1984 under No. 0128849. The other has a different partial amino acid sequence than that reported for the human placenta-derived TGF- β , but is active in the same anchorage-independent cell growth assay as was used to characterize the biological activity of TGF- β .

There have been prior attempts to identify and isolate factors in bone that are involved in stimulating bone growth. U.S. Pat. No. 4,294,753 to Urist reports the preparation of a crude "bone morphogenic protein" by extracting demineralized bone with urea or guanidine hydrochloride and reprecipitating protein from the extract. Urist subsequently reported (Urist, M. R., Clin Orthop Rel Res (1982) 162:219) that ion exchange purification of this crude protein mixture yielded an activity that was unadsorbed to carboxymethylcellulose (CMC) at pH 4.8. Urist's most recent reports (Science (1983) 220:680-685 and Proc Natl Acad Science (USA) (1984) 81:371-375) describe BMPs having molecular weights of 17,500 and 18,500 daltons.

U.S. Pat. No. 4,434,094 reports the partial purification of a bone generation-stimulating, bone-derived protein factor by extraction with chaotropic agents, fractionation on anion and cation exchange columns, and recovery of the activity from a fraction adsorbed to CMC at pH 4.8. This new protein fraction was termed "osteogenic factor" (OF) and was characterized as having a molecular weight below about 30,000 daltons and as tracking the purification process described. The proteins of the current invention were purified to homogeneity using a purification procedure that is similar in part to that disclosed in U.S. Pat. No. 4,434,094.

DISCLOSURE OF THE INVENTION

The invention provides two CIFs found in mammalian bone that:

- (1) are co-factors for inducing cartilage formation in vivo.
- (2) promote connective tissue deposition in vivo in the absence of any added activating agent or co-factor, and
- (3) are active in the anchorage-independent cell growth assay used to characterize TGF- β (this assay is sometimes called the "TGF- β assay" herein and is described in Methods for Preparation of Media, Supplements, and Substrata for Serum-Free Animal Cell Culture (1984) pp 181-194, Alan R. Liss, Inc.). These two polypeptides are called CIF-A and CIF-B, respectively, herein.

The invention further provides a process for obtaining CIF-A and CIF-B in substantially pure form from bone. In this regard the presence in bone of proteins having activity in the TGF- β assay has not been reported previously. CIF-A has a partial (30 amino acids) N-terminal amino acid sequence reported for human placenta-derived TGF- β . But lacking total sequencing of both molecule, it is not known whether they are the same polypeptide. CIF-B has a different partial (30 amino acids) N-terminal amino acid sequence. The invention further provides compositions and methods for promoting proliferation of cells and for promoting connective tissue deposition using the CIFs.

Accordingly, one aspect of the invention is a polypeptide that: (a) is found in mammalian bone, (b) is a co-factor for inducing cartilage formation, (c) is active in the TGF- β assay, (d) is a dimer of approximately 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure.

The process for isolating the CIFs from bone involves;

- (a) extracting demineralized bone (DMB) with a chaotropic (dissociative) extractant that solubilizes nonfibrous proteins;
- (b) subjecting the extract from step (a) to gel filtration to recover a fraction containing proteins of molecular weight 10,000-40,000 daltons;
- (c) adsorbing the fraction from step (b) onto a carboxymethyl cellulose cation exchanger at approximately pH 4.5-5.5 under denaturing conditions;
- (d) eluting the adsorbed fraction from the cation exchanger with a sodium chloride gradient;
- (e) subjecting the portion of the eluate from step (d) that elutes at approximately 150 to 250 mM sodium chloride to reverse phase high performance liquid chromatography (RP-HPLC) or a nondenaturing gel electrophoresis; and
- (f) recovering the CIFs from the RP-HPLC or gel electrophoresis.

The composition for promoting proliferation of normal animal cells comprises;

- (a) at least one of the above-described polypeptides;
- (b) a TGF- β activating agent; and
- (c) a pharmaceutically acceptable carrier, the polypeptide and activating agent being present in amounts sufficient to promote cell proliferation.

The method of promoting normal cell proliferation comprises administering an effective amount of the above-described composition to a patient.

The composition for promoting connective tissue deposition comprises:

- (a) an effective amount of one or both of the above-described polypeptides substantially free of co-factor or activating agent, and
- (b) a pharmaceutically acceptable carrier.

The method of promoting connective tissue deposition comprises placing the above-described composition at a predetermined site within the body of a patient.

BRIEF DESCRIPTION OF THE DRAWINGS

In the drawings:

FIG. 1 is a graph of the optical densities (absorbances) (280 nm) and in vitro chondrogenic activities of the gel filtration fractions of the example (section C), *infra*;

FIG. 2 is a graph of the optical densities (280 nm) of eluate fractions from the preparative ion exchange chromatography of the example (section D), *infra*;

FIG. 3 is a chromatogram and autoradiograph of the SDS-PAGE analysis of the RP-HPLC resolved CM-1 fraction described in section E of the example, *infra*.

FIG. 4 is a graph of the UV absorbance and electrophoretic profiles of peaks A and B of the preparative RP-HPLC of the example (section E), *infra*;

FIG. 5 is a graph of the results of the enzyme-linked immunosorbent assays (ELISAs) for in vitro chondrogenic activity of the proteins of peaks A and B from the RP-HPLC of the example (section E), *infra*;

FIG. 6 is a graph of the results of the ELISAs of the acid-urea gel electrophoresis fractions (section F) of the example, *infra*; and

FIG. 7 is a graph of the results of the TGF- β assays described in section I of the example, *infra*.

MODES OF CARRYING OUT THE INVENTION

The polypeptides of the invention were isolated from bovine bone. The polypeptides have been only partially sequenced at this time. In view of this and since the complete amino acid sequence of TGF- β has not been reported, the primary structure relationships between the invention CIFs and TGF- β s are not known completely. However, based on comparisons of partial sequences of N-termini, molecular weight analyses, apparent secondary structures, and biological activity, CIF-A may be identical or substantially homologous to TGF- β derived from human placenta. CIF-A is, however, more prevalent in bone than TGF- β is in platelets, placenta, or kidney. Thus, if the two are the same molecule, it appears more likely that the molecule acts as a chondrogenic/osteogenic agent in its native setting rather than as a transforming growth factor.

The CIFs have in vitro chondrogenic activity by themselves, in vivo chondrogenic activity in combination with certain chondrogenic co-factors, in vivo connective tissue deposition activity by themselves, and in vitro TGF- β activity in combination with epidermal growth factor (EGF). In view of their chondrogenic activity and the mode of endochondral bone formation, they are also expected to have in vivo osteogenic activity in combination with chondrogenic co-factors.

In view of the showings that bone inductive proteins from human, monkey, bovine and rat are nonspecies-specific in their abilities to produce endochondral bone in xenogeneic implants (Sampath, T. K., et al, Proc Natl Acad Sci (USA) (1983) 80:6591) and that human platelet/human placenta/bovine kidney-derived TGF- β is nonspecies-specific between rodents, cattle, and humans, it is believed that the polypeptides of this invention have been highly conserved among mammalian species (i.e., polypeptides from different mammalian species have amino acid sequences that vary, if at all, in one or more amino acid residue additions, deletions, or substitutions that do not affect that nonspecies-specific activity of the molecule adversely). In this regard the term "substantially equivalent" as used to describe a polypeptide is intended to mean polypeptides, whether native or synthetic and regardless of species or derivation, that have the

same amino acid sequence as the referenced polypeptide, and polypeptides of substantially homologous (i.e., at least 90% identity in amino acid sequence) but different amino acid sequence, which difference(s) does not affect nonspecies-specific activity adversely. Accordingly, the polypeptides of this invention may be derived from or found in cells or tissue of diverse animal origin or synthesized by recombinant DNA technology. Porcine or bovine long bone are preferred native sources of the polypeptides because of the ready availability of such bone and the abundance of the polypeptides in bone.

A variety of initial preparation procedures are possible when bone is used as a source, but basically the bone is first cleaned using mechanical or abrasive techniques, fragmented, and further washed with, for example, dilute aqueous acid preferably at low temperature, and then defatted by extraction with a lipophilic solvent such as ether or ethyl acetate. The bone is then demineralized by removal of the calcium phosphates in their various forms, usually by extraction with stronger acid. These techniques are understood in the art, and are disclosed, for example, in U.S. Pat. No. 4,434,094. The resulting preparation, a demineralized bone, is the starting material for the preparation of the polypeptides of the invention.

The initial extraction is designed to remove the non-fibrous (e.g., non-collagenous) proteins from the demineralized bone. This can be done with the use of chaotropic agents such as guanidine hydrochloride (at least about 4 molar), urea (8 molar) plus salt, or sodium dodecylsulfate (at least about 1% by volume) or such other chaotropic agents as are known in the art (Termine, et al, J Biol Chem (1980) 255:9760-9772; and Sajera and Hascall, J Biol Chem (1969) 244:77-87 and 2384-2396). The extraction is preferably carried out at reduced temperatures in the presence of a protease inhibitor to reduce the likelihood of digestion or denaturation of the extracted protein. Examples of protease inhibitors that may be included are phenylmethylsulfonyl fluoride (PMSF) sodium azide, N-ethyl maleimide (NEM), benzamidine, and 6-aminohexanoic acid. The pH of the medium depends upon the extractant used. The process of extraction generally takes on the order of about 4 hr to one day.

After extraction, the extractant may be removed by suitable means such as dialysis against water, preceded by concentration of ultrafiltration if desired. Salts can also be removed by controlled electrophoresis, or by molecular sieving, or by any other means known in the art. It is also preferred to maintain a low temperature during this process so as to minimize denaturation of the proteins. Alternatively, the extractant need not be removed, but rather the solution need only be concentrated, for example, by ultrafiltration.

The extract, dissolved or redissolved in chaotropic agent, is subjected to gel filtration to obtain fractions of molecular weight below about 40,000 daltons, thus resulting in a major enhancement of purity. Gel sizing is done using standard techniques, preferably on a Sephacryl column at room (10°-25° C.) temperature. The low molecular weight fraction is then subjected to ion exchange chromatography using carboxymethyl cellulose (CMC) at approximately pH 4.5-5.2, preferably about 4.8, in the presence of a nonionic chaotropic agent such as urea. Other cation exchangers may be used, including those derived from polyacrylamide and cross-linked dextran; however cellulosic cation exchangers are preferred. Of course, as in any ion exchange procedure, the solution must be freed of competing ions before application to the column, and is eluted in an increasing salt concentration gradient as is understood in the art. The

fraction eluting from CMC at about 150 to 250 mM NaCl contains the CIFs.

The eluate fraction from the cation exchange chromatography is then subjected to RP-HPLC or a non-denaturing gel electrophoresis for final purification. Standard RP-HPLC techniques and gel electrophoresis techniques are used. Exemplified below is a commercially available RP-HPLC column using a commercially prescribed RP-HPLC protocol. This final purification yields the two polypeptides in substantially pure form. "Substantially pure" means that a polypeptide contains less than about 5% by weight contaminants.

EXAMPLE

The following example is intended to illustrate the process for purification as applied to a particular sample. It is not intended to limit the invention.

A. Preparation of Demineralized Bone

Bovine metatarsal bone was obtained fresh from the slaughterhouse and transported on dry ice. The bones were cleaned of marrow and non-bone tissues, broken fragments smaller than 1 cm diameter, and pulverized in a mill at 4° C. The pulverized bone was washed twice with 9.4 liters of double distilled water per kg of bone for about 15 min each, and then washed overnight in 0.01N HCl at 4° C. Washed bone was defatted using 3×3 volumes ethanol, followed by 3×3 volumes diethylether, each washed for 20 min, and all at room temperature. The resulting defatted bone powder was then demineralized in 0.5N HCl (25 l/kg defatted bone) at 4° C. The acid was decanted and the resulting DMB washed until the wash pH was greater than 4, and the DMB dried on a suction filter.

B. Extraction of Noncollagenous Proteins

The DMB as prepared in paragraph A was extracted with 3.3 l of 4M guanidine-HCl, 10 mM ethylenediaminetetraacetic acid (EDTA), pH 6.8, 1 mM PMSF, 10 mM NEM per kg for 16 hrs, the suspension suction filtered and the non-soluble fractions were combined and concentrated at least 5-fold by ultrafiltration using an Amicon ultrafiltration (10K) unit, and the concentrate dialyzed against 6 changes of 35 volumes cold deionized water over a period of 4 days, and then lyophilized. All of the procedures of this paragraph were conducted at 4° C. except the lyophilization which was conducted under standard lyophilization conditions.

C. Gel Filtration

The extract from paragraph B, redissolved in 4 M guanidine-HCl, was fractionated on a Sephacryl S-200 column equilibrated in 4M guanidine-HCl, 0.02% sodium azide, 10 mM EDTA, pH 6.8. Fractions were assayed by their absorbance at 280 nm and chondrogenic activity by ELISA and the fractions were combined as shown in FIG. 1. Fraction F2 of FIG. 1, constituting a low molecular weight (LMW, 10,000–40,000 daltons) protein fraction possessing the greatest activity was dialyzed against 6 changes of 180 volumes of deionized water and lyophilized. All operations except lyophilization and dialysis (4° C.) were conducted at room temperature.

D. Ion Exchange Chromatography

Fraction F2 from paragraph C was dissolved in 6M urea, 10 mM NaCl, 1 mM NEM, 50 mM sodium acetate, pH 4.8 and centrifuged at 10,000 rpm for 5 min. The supernatant was fractionated on a CM52 (a commercially available CMC) column equilibrated in the same buffer. Bound proteins were eluted from the column using a 10 mM to 400 mM NaCl gradient in the same buffer, and a total volume of

350 ml at a flow rate of 27 ml/hr. Three major fractions, designated CM-1, CM-2, and CM-3 were collected as shown in FIG. 2. CM-2 and CM-3 eluted at approximately 150–250 mM NaCl. Each fraction was dialyzed against 6 changes of 110 volumes of deionized water for 4 days and lyophilized. All of the foregoing operations were conducted at room temperature except dialysis (4° C.).

E. RP-HPLC

The lyophilized fractions CM-1 and combined lyophilized fractions CM-2 and CM-3 from ¶D were each dissolved in 0.1% trifluoroacetic acid (TFA) and aliquots of the solutions loaded onto a Vydac C18 RP-HPLC column (4.6 mm ID×25 cm) and washed with 0.1% TFA for 5 min at 1 ml/min. The eluting solvent was a 0%–60% acetonitrile gradient in 0.1% TFA at a rate of 2%/min.

FIG. 3 shows the chromatogram of the RP-HPLC of CM-1. Peak 3 contains predominantly 28 Kd chondrogenic/osteogenic co-factor protein and peak 6 contains predominantly 36 Kd chondrogenic/osteogenic co-factor protein. Upon rechromatography using the same RP-HPLC system, the two polypeptides were purified to give single bands on 15% SDS-PAGE (analysis also shown in FIG. 3). These protein co-factors are not part of the present invention and are the subject of a separate, commonly owned U.S. patent application.

Two peaks were obtained from the RP-HPLC of combined CM-2 and CM-3--peak A at about 29.5 min and peak B at about 31.2 min. FIG. 4 shows the absorbance and electrophoretic profiles (reduced and nonreduced) of peaks A and B. The proteins of these peaks were designated CIF-A and CIF-B, respectively.

The proteins were stored in 0.1% TFA/acetonitrile eluting solution at -20° C. until used.

F. Alternate Purification by Gel Electrophoresis

The combined lyophilized fractions CM-2 and CM-3 were fractionated by electrophoresis on an acetic acid-urea gel using the general procedure of Paynim, S. and Chalkley, R., Arch Bioch Biophys (1969) 130:337–346.

G. Assay for In Vitro Chondrogenic Activity

The presence of the desired protein in fractions during purification was confirmed using an in vitro assay for the production of cartilage-specific proteoglycans (PG), the identity of which was confirmed by ELISA. This assay is an agarose gel culture model using mesenchymal cells isolated from rat fetal muscle. It assesses the ability of the samples to induce the production of PG. The correlation between in vitro cartilage induction and in vivo bone formation has been shown by Seyedin, S., et al. J Cell Biol (1983) 97:1950–1953.

The cell culture was prepared by removing muscle tissue aseptically from the upper limbs of nineteen-day-old Sprague Dawley rat fetuses, mincing the tissue and culturing it in Eagle's Minimum Essential Medium (MEM) with 10% fetal bovine serum (FBS) and 50 units penicillin, 50 µg streptomycin per ml. Cellular outgrowth usually reached confluency within one week, whereupon cells were trypsinized, split 1:2 and used for experimentation within the first three passages.

The cells were placed in agarose gel cultures either with control medium or with samples to be tested. The procedure was basically that of Benya, et al, Cell (1982) 30:215. Briefly, the cell monolayers were harvested by trypsinization, counted on a hemocytometer, and resuspended at two times the final cell concentration in the medium with or without the protein fraction to be tested. The

control medium was either Hams F-12, Dulbecco's Minimum Essential Medium (DMEM) or CMRL 1066 (Gibco) each containing 10% FBS and antibiotics. The test protein fractions in 0.01N HCl were diluted directly to the desired concentration of test protein diluted with an equal volume with 1% low melting agarose (Bio-Rad, #162-0017) in F-12, and 0.2 ml of the dilution was plated on 17 mm wells coated with 0.15 ml of 1% high melting (Bio-Rad, #162-0100) agarose. The resulting cultures were incubated at 37° C. for 5 min, chilled at 4° C. for 10 min, and then overlaid with 1 ml of the corresponding medium (control or test protein). The cells were then cultured in a humidified atmosphere of 5% CO₂, 95% air and fed every 3-4 days thereafter by a complete change with control medium. After 7 days the cultures were frozen and stored at -80° C. before assay.

The cultures were assayed by thawing at 4° C., homogenizing in 4M guanidine-HCl with 50 nM Na acetate, 13 mM EDTA, 6 mM NEM, and 3 nM PMSF at pH 5.8, and extracting by shaking overnight at 4° C. The supernatant fraction from centrifugation at 25,000xg for 40 min at 4° C. was dialyzed overnight at 4° C. against 50 volumes 0.2 M NaCl, 50 mM Tris, pH 7.4. The supernatant was assayed for PG by ELISA as described by Renard, et al, Anal Biochem (1980) 104:205, and in U.S. Pat. No. 4,434,094.

Briefly, for the ELISA, antiserum to PG was raised in rabbits using standard techniques which showed no cross-reactivity with hyaluronic acid or PG extracted from rat bone. Purified PG (Seyedin, S., et al, supra) from Swarm rat chondrosarcoma tissue was used as standard antigen. The dialyzed samples were diluted 1:1 (v/v) in phosphate-buffered saline (PBS) with 0.05% Tween 20, 1 mg/ml bovine serum albumin (BSA), pH 7.2 for assay. Horseradish peroxidase conjugated goat anti-rabbit IgG (Tago) was the second antibody with o-phenylenediamine as substrate.

The results of the ELISA of the RP-HPLC purified CIF-A and CIF-B are shown in FIG. 5. As indicated there, the sensitivity of the assay is within 1 to 5 ng/ml of culture media. The results of the ELISAs on the gel slices of section F are shown in FIG. 6. These results are comparable to the results on CIF-A and CIF-B (corresponding to gel slices 7 and 6) from the RP-HPLC.

H. Characterization of the Purified Proteins

CIF-A was shown to be a 25,800 dalton protein which on reduction, yielded a 14,800 dalton polypeptide by measurements of the mobilities of the proteins in a 15% Laemmli polyacrylamide gel in SDS (FIG. 4) as described by Laemmli, U. K., et al, Nature (1970) 227:680. It is well understood that molecular weights so determined are approximate and their values are dependent on the method used. The conformation of the protein affects its mobility in this system, and, therefore, the molecular weights obtained will be similar, but no necessarily identical when determined by other procedures. The presence of a single band in the profile of the reduced protein indicates the protein is probably a dimer composed of two polypeptide chains having substantially equivalent amine acid sequences (i.e., it is a

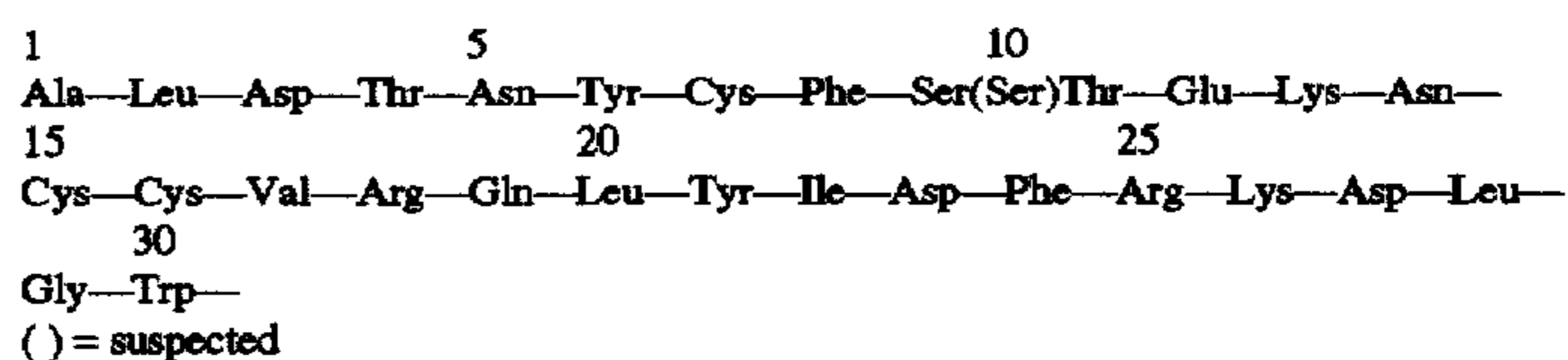
homodimer). The discrepancy between the measured weights of dimer and the individual chains is an artifact of the procedure.

CIF-A maintained its activity in the ELISA assay of paragraph G above even after heating for 3 min at 100° C. in PBS, after treatment with collagenase for 2 hrs at 37° C. in 0.1M Tris, pH 7.4, 5 mM CaCl₂, 0.02 mM PMSF with a ratio of collagenase to protein of 400 units/mg protein, and after treatment with trypsin for 2 hrs at 37° C. in 50 mM Tris, pH 7.4, 10 mM CaCl₂ with a ratio of trypsin to protein of 100 units/mg of protein. However, the protein lost activity after treatment for 1 hr at room temperature in PBS containing 5 mM dithiothreitol (DTT), which would effect reduction of disulfide linkages. Similarly, SDS treatment or fractionation on SDS-PAGE resulted in inactivation of the protein, presumably due to denaturation or complexing by the SDS. The partial amino acid composition of the CIF-A is shown in Table 1.

TABLE 1

Amino Acid	Mols/100 Mols Recovered
Asp	9.2
Glu	9.2
Ser	7.0
His	2.7
Gly	16.5
Thr	2.7
Arg	5.9
Ala	6.6
Tyr	3.2
Met	0.0
Val	7.5
Phe	3.0
Ile	3.9
Leu	8.6
Lys	13.9
Pro	ND
Cys	ND
Trp	ND

Amino acid sequence analysis of the protein of peak A by the ninhydrin method showed that it has the following single N-terminal sequence:



This N-terminal sequence is identical to that reported for human placenta-derived TGF-β.

CIF-B had a slightly different molecular weight (26,000) as measured by the same procedure. This difference may be caused by the procedure. Accordingly, both proteins are considered to have a molecular weight of approximately 26,000 daltons as measured by SDS-PAGE. On reduction, CIF-B showed a single band at approximately 14,200 daltons indicating that it, too, is probably a homodimer. It has the amino acid composition set forth in Table 2.

with pharmaceutically acceptable carriers that are adapted for the particular mode of administration. Topical dosage forms will typically be formulated as sprays, gels, ointments, or salves. Implants will be formulated as injectables. Systemic dosage forms may be formulated for enteral administration (e.g., liquids, pills, tablets) or for parenteral injection. The dosages used in such applications cannot be specified because of the nature of all proliferation activity and the variability in wounds and other traumata.

The CIFs may also be useful for treating bone deficiencies, such as osteoporosis and osteopetrosis, systemically. For such treatment the CIFs will be formulated in therapeutically effective amounts with injectable carriers and administered parenterally to the patient.

The ability of the CIFs to withstand treatment with trypsin without losing activity may make it possible to isolate them from demineralized bone powder by means of enzymatic digestion. In such a process the demineralized bone powder is digested with an aqueous solution of trypsin and/or other proteases that do not degrade the proteins of interest under conditions at which such enzymes are active. This treatment digests the majority of other protein components in the powder. The proteins of interest may be purified from the resulting digest using one or more of the fractionation techniques described above (gel filtration, ion exchange chromatography, RP-HPLC or nondenaturing gel electrophoresis). Depending upon the extent to which the proteins of interest are released from the bone matrix and not complexed with other materials, use of solubilizing agents may be avoided. In this regard the pure CIFs are substantially soluble in water.

We claim:

[1. A method of promoting proliferation of cells in an animal comprising

administering to the animal a composition comprising a TGF-beta activating agent, a pharmaceutically acceptable carrier, and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF-beta assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide and activating agent are present in amounts sufficient to promote bone growth.]

[2. The method of claim 1 wherein each chain of said dimer has the following N-terminal sequence: Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp-.]

[3. A method for inducing cartilage and bone formation for repairing, replacing, and or augmenting cartilage and bone tissue in animals, which method comprises:

administering to a mammal in need of bone repair, replacement, or augmentation a composition comprising a TGF-beta activating agent, a pharmaceutically acceptable excipient, and a polypeptide the (a) is found in mammalian bone, (b) is active in the TGF-beta assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide and activating agent are present in amounts sufficient to promote bone growth.]

[4. The method of claim 3 wherein each chain of said dimer has the following N-terminal sequence: Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp-.]

[5. A method for treating osteoporosis or osteopetrosis, which comprises:

systemically administering an effective amount of a composition comprising a TGF-beta activating agent, a pharmaceutically acceptable excipient, and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF-beta assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide and activating agent are present in amounts sufficient to promote bone growth.]

[6. The method of claim 5 wherein each chain of said dimer has the following N-terminal sequence: Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp-.]

[7. The method of claim 3 wherein each chain of said dimer has the following N-terminal sequence: Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp.]

[8. The method of claim 5 wherein each chain of said dimer has the following N-terminal sequence: Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp.]

9. A method for inducing cartilage and bone formation for repairing, replacing, and/or augmenting cartilage and bone tissue in animals, which method comprises administering to an animal in need of bone repair, replacement, or augmentation a composition comprising a TGF-beta activating agent, a pharmaceutically acceptable excipient, and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF-beta assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide and activating agent are present in amounts sufficient to promote bone growth.

10. The method of claim 9 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp (TGF- β 1).

11. The method of claim 9 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp (TGF β -2).

12. A method for treating osteoporosis or osteopetrosis, which comprises systemically administering to an individual in need of such treatment an effective amount of a composition comprising a TGF-beta activating agent, a pharmaceutically acceptable excipient, and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF-beta assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide and activating agent are present in amounts sufficient to promote bone growth.

13. The method of claim 12 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp (TGF β -1).

14. The method of claim 12 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp (TGF β -2).

15. A method for inducing cartilage and bone formation for repairing, replacing, and/or augmenting cartilage and bone tissue in animals, which method comprises administering to an animal in need of bone repair, replacement, or augmentation a composition comprising a pharmaceutically acceptable excipient and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF- β assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide is present in an amount sufficient to promote bone growth.

16. The method of claim 15 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp (TGF β -1).

17. The method of claim 15 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp (TGF β -2).

18. A method for treating osteoporosis or osteopetrosis, which comprises systemically administering to an individual in need of such treatment an effective amount of a composition comprising a pharmaceutically acceptable excipient and a polypeptide that (a) is found in mammalian bone, (b) is active in the TGF- β assay, (c) is a co-factor for inducing cartilage formation in vivo, (d) is a dimer having an approximate molecular weight of 26,000 daltons as determined by SDS-PAGE, and (e) is substantially pure, wherein said polypeptide is present in an amount sufficient to promote bone growth.

19. The method of claim 18 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Thr-Asn-Tyr-Cys-Phe-Ser-(Ser)-Thr-Glu-Lys-Asn-Cys-Cys-Val-Arg-Gln-Leu-Tyr-Ile-Asp-Phe-Arg-Lys-Asp-Leu-Gly-Trp (TGF β -1).

20. The method of claim 18 wherein each chain of said dimer has the following N-terminal sequence:

Ala-Leu-Asp-Ala-Ala-Tyr-Cys-Phe-Arg-Asn-Val-Gln-Asp-Asn-Cys-Cys-Leu-Arg-Pro-Leu-Tyr-Ile-Asp-Phe-Lys-Arg-Asp-Leu-Gly-Trp (TGF β -2).

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