



(86) Date de dépôt PCT/PCT Filing Date: 1997/10/10

(87) Date publication PCT/PCT Publication Date: 1998/05/07

(45) Date de délivrance/Issue Date: 2008/04/15

(85) Entrée phase nationale/National Entry: 1999/04/27

(86) N° demande PCT/PCT Application No.: US 1997/018342

(87) N° publication PCT/PCT Publication No.: 1998/018816

(30) Priorité/Priority: 1996/10/31 (US08/742,256)

(51) Cl.Int./Int.Cl. *C07K 7/06* (2006.01),
A61K 38/08 (2006.01), *A61K 38/18* (2006.01),
A61K 47/00 (2006.01), *C07K 14/495* (2006.01)

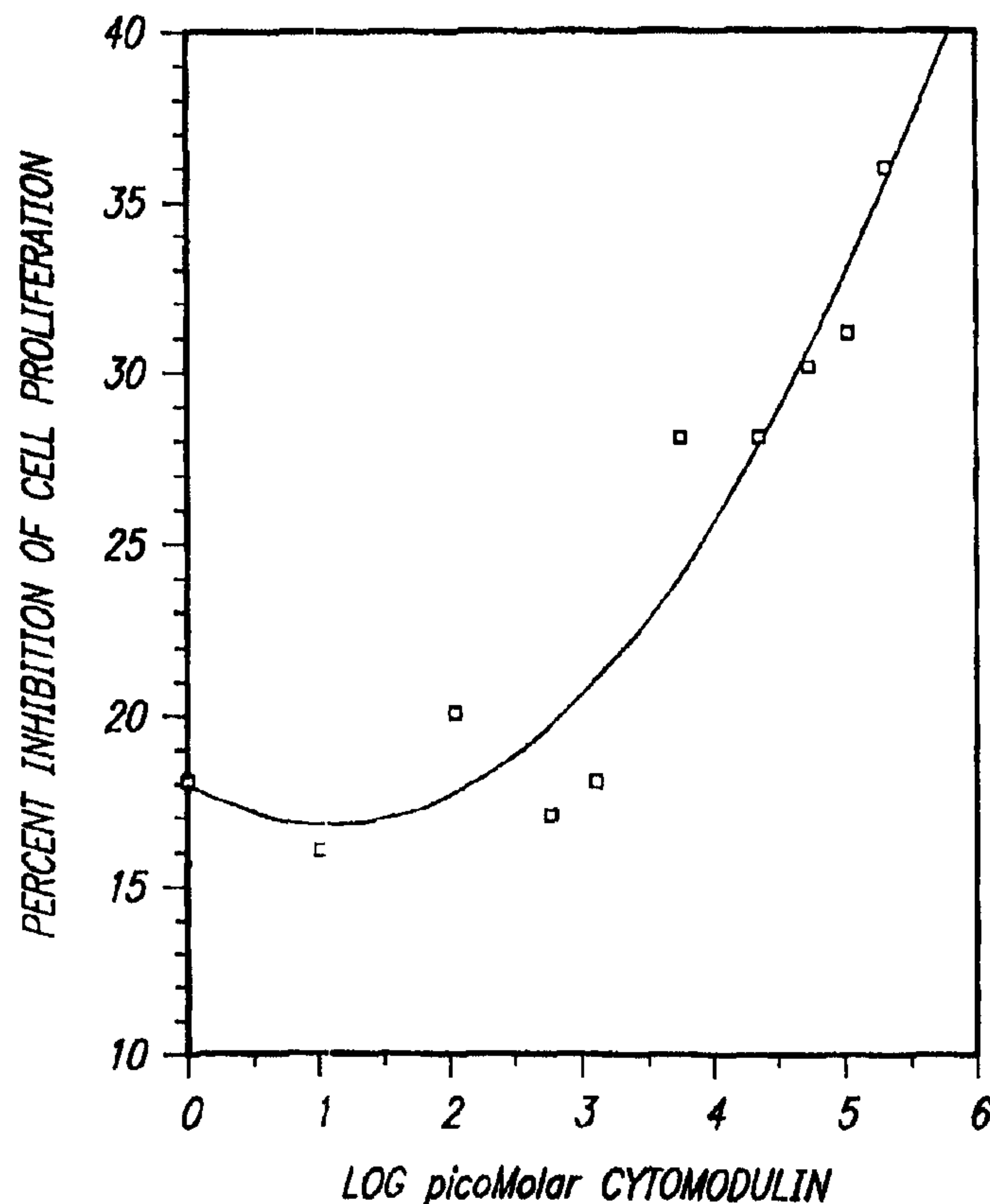
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(54) Titre : COMPOSITIONS DE PEPTIDES DOTEES D'UNE ACTIVITE RESSEMBLANT A CELLE DES FACTEURS DE CROISSANCE

(54) Title: PEPTIDE COMPOSITIONS WITH GROWTH FACTOR-LIKE ACTIVITY



(57) Abrégé/Abstract:

Small peptide mimics of TGF- β , having the general sequence $AA_j-AA_{j+1}-AA_{j+2}$ followed by AA_{i+n} shortly thereafter, have been prepared. In this sequence, AA_j is alanine, asparagine, or leucine, AA_{j+1} is valine or isoleucine, AA_{j+2} is alanine, n is 3, 4 or 5 such

(57) **Abrégé(suite)/Abstract(continued):**

that there are $n-3$ amino acid residues in between AA_{j+2} and AA_{i+n} , and AA_{j+n} is glutamic acid, aspartic acid, glutamine or asparagine. Because the essential requirement for TGF- β activity is the peptide's ability to form a stable β -bend structure under physiologic conditions, the inventive peptides are collectively referred to as β -bend peptides. Compositions for applications such as tissue repair are also provided that comprise a biocompatible matrix having cytomodulin or a cytomodulin analog admixed with or carried by the matrix.

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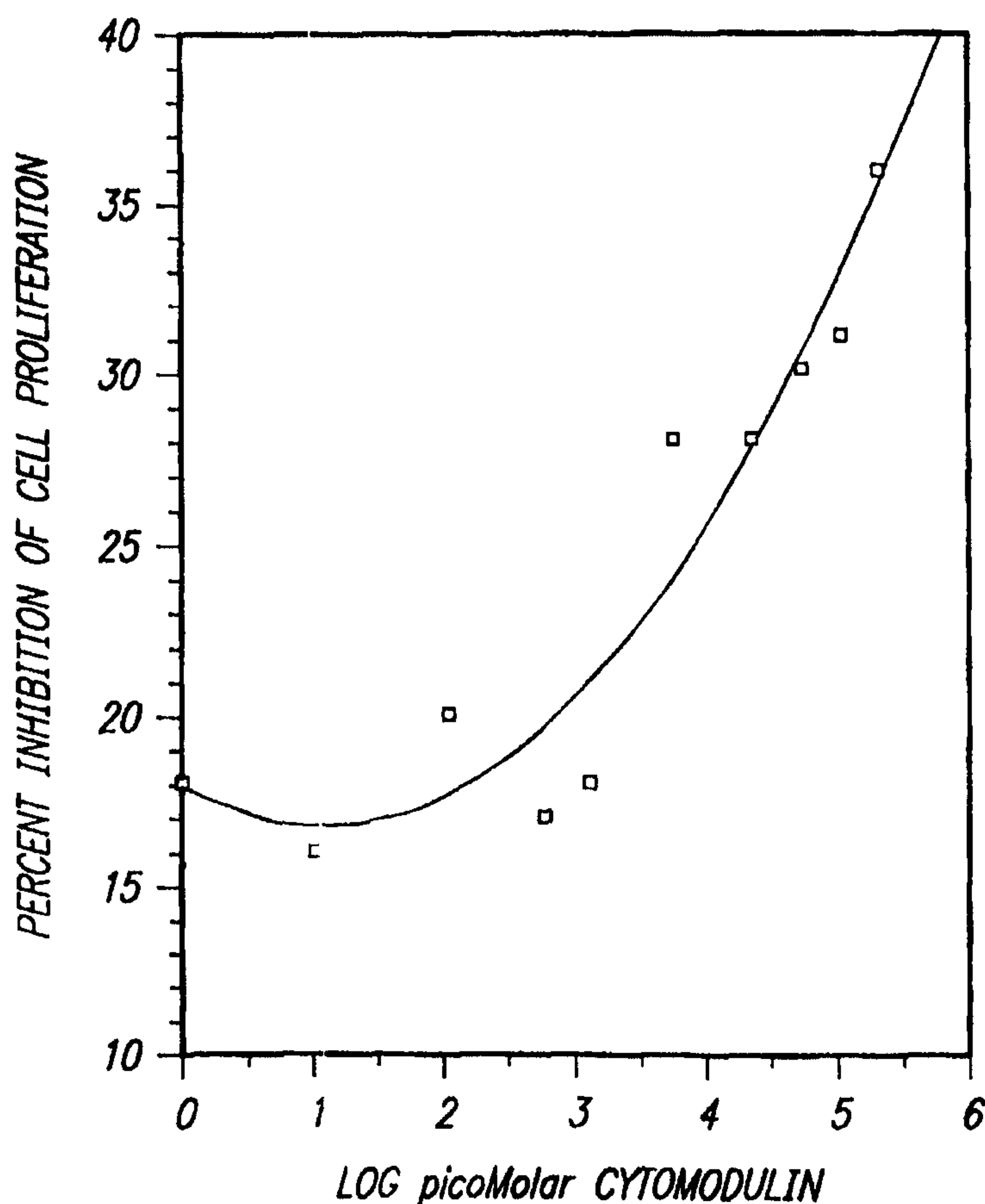
INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification ⁶ : C07K 7/06, A61K 38/08		A1	(11) International Publication Number: WO 98/18816
			(43) International Publication Date: 7 May 1998 (07.05.98)
(21) International Application Number: PCT/US97/18342 (22) International Filing Date: 10 October 1997 (10.10.97) (30) Priority Data: 08/742,256 31 October 1996 (31.10.96) US (71) Applicant: THE REGENTS OF THE UNIVERSITY OF CALIFORNIA [US/US]; 22nd floor, 300 Lakeside Drive, Oakland, CA 94612-3550 (US). (72) Inventors: BHATNAGAR, Rajendra, S.; 173 Los Robles, Burlingame, CA 94010 (US). QIAN, Jing, Jing; 3761 Elston Drive, San Bruno, CA 94066 (US). (74) Agent: BERLINER, Robert; Fulbright & Jaworski, L.L.P., 29th floor, 865 S. Figueroa Street, Los Angeles, CA 90017-2571 (US).		(81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZW, ARIPO patent (GH, KE, LS, MW, SD, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG). Published <i>With international search report.</i> <i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>	

(54) Title: PEPTIDE COMPOSITIONS WITH GROWTH FACTOR-LIKE ACTIVITY

(57) Abstract

Small peptide mimics of TGF- β , having the general sequence AA_j-AA_{j+1}-AA_{j+2} followed by AA_{i+n} shortly thereafter, have been prepared. In this sequence, AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine or isoleucine, AA_{i+2} is alanine, n is 3, 4 or 5 such that there are n-3 amino acid residues in between AA_{j+2} and AA_{i+n}, and AA_{j+n} is glutamic acid, aspartic acid, glutamine or asparagine. Because the essential requirement for TGF- β activity is the peptide's ability to form a stable β -bend structure under physiologic conditions, the inventive peptides are collectively referred to as β -bend peptides. Compositions for applications such as tissue repair are also provided that comprise a biocompatible matrix having cytomodulin or a cytomodulin analog admixed with or carried by the matrix.



PEPTIDE COMPOSITIONS WITH
GROWTH FACTOR-LIKE ACTIVITY

5 Field of the Invention

The invention generally relates to growth factors and neurotrophic factors, and more particularly to small, synthetic peptides having (or mimicking) TGF- β activity, and to matrices and compositions including
10 these small peptide mimics.

15 Background of the Invention

Growth factors are substances, such as polypeptide hormones, which affect the growth of defined populations of animal cells *in vivo* or *in vitro*, but which are not nutrient substances. Proteins involved in
20 the growth and differentiation of tissues may promote or inhibit growth, and promote or inhibit differentiation, and thus the general term "growth factor" includes cytokines and trophic factors.

Growth factors typically are polypeptides
25 ranging in molecular weights from 5000 to 50,000 daltons. Based on structural similarities, growth factors are categorized into families which include: insulin-like growth factors (IGFs), platelet-derived growth factors (PDGFs), fibroblast growth factors (FGFs), epidermal
30 growth factors (EGFs), nerve growth factors (NGFs), and transforming growth factors type-beta (TGF- β s).

Transforming growth factor- β s were originally named for their ability to transform normal fibroblasts to cells capable of anchorage-independent growth.
35 However, despite the name, TGF- β s are multifunctional growth factors that are required for the normal development, growth, and differentiation of various

epithelial, endothelial, and mesenchymal cells. As with other cytokines, the specific effect of TGF- β s depend on the particular cell type and its surrounding environment.

The effects of TGF- β s on cells are generally
5 classified as proliferative and non-proliferative. As originally established with the first experiments on fibroblasts, TGF- β s are bona fide growth factors. Two important cell types in which proliferation is enhanced by TGF- β are osteoblasts and Schwann cells of the
10 peripheral nervous system. However, in many cells, TGF- β s are potent inhibitors of cell proliferation. This negative growth control may be the regulatory mechanism that checks regeneration of certain tissues and may play a role in the initiation of carcinogenesis.

15 The most important non-proliferative function of TGF- β s are in enhancing the formation of extracellular matrices. Although this is achieved primarily through the increased transcription of both collagen and fibronectin, the inhibition of the proteases from
20 degrading the matrix also contributes to its stability. Degradation of the extracellular matrix is inhibited by the decrease in the secretion of the proteases themselves and the simultaneous increase in the levels of protease inhibitors. The marked and generalized effect of TGF- β
25 on extracellular matrices is likely to play a major role in tissue repair processes and the pathogenesis of certain fibrotic diseases.

DNA encoding several different receptors for TGF- β has recently been described by Lin et al., PCT
30 application WO93/09228, published May 13, 1993. The availability of the TGF- β receptors will facilitate further assessments of TGF- β functions.

Many members of the TGF- β super family have been characterized. For example, Basler et al. have
35 graphically represented the sequence relationship between members of the TGF- β superfamily. *Cell*, 73, pp. 687-702 (1993). Massagué, *Annu. Rev. Cell Biol.*, 6, pp. 597-641

(1990) also reviews the transforming growth factor- β family, including a discussion of the mechanisms of TGF- β actions. An NMR characterization of the secondary structure of TGF- β 1 has been reported, and a refined 3-
5 dimensional crystal structure of TGF- β 2 described, by Daopin et al., *Proteins*, 17, pp. 176-192 (1993). The monomer of TGF- β 2 adopts a fold that resembles a slightly curled left hand with two anti-parallel β -sheets forming four fingers of the hand. These four finger regions
10 together with conserved disulfides define the fold for the TGF- β superfamily.

Also among TGF- β members are the bone morphogenetic proteins (BMP). The BMPs have been indicated as useful in wound healing, tissue repair, and to induce
15 cartilage and/or bone growth. For example, PCT Application 9309229, inventors Israel and Wolfman, published May 13, 1993, describes uses of proteins with bone stimulating activity such as bone fracture healing and possibly the treatment of periodontal disease and
20 other tooth repair processes. A recent special article by *C&EN*, Hubbell and Langer, pp. 42-54 (March 13, 1995) reports that a BMP has been incorporated into polymer particles so that as the polymer degrades, the protein is slowly released to surrounding tissues, where it
25 stimulates the migration of cells into the porous matrix and, ultimately, the synthesis of new bone. The article also notes that bone has been produced by slowly releasing TGF- β .

Because of the wide applicability of TGF- β s in
30 clinical therapies, they have been the focus of much research. Although much of the research involved in *vitro* uses, recent *in vivo* studies have confirmed some of the more promising *in vitro* effects. As a consequence, some of the possible clinical uses for TGF- β s include the
35 stimulation of angiogenesis, the formation of granulation tissue associated with wound healing, and the formation of bone and cartilage.

Nucleic acid encoding TGF- β and a variety of uses for TGF- β are described in U.S. Patent 5,284,763, issued February 8, 1994, inventors Derynk and Goeddel. U.S. Patent 5,258,029, issued November 2, 1993, inventors
5 Chu et al. describe preparations of stress-bearing prothesis with bony ingrowth occurring after implantation, which prothesis includes TGF- β carried by a collagen composition or a ceramic. U.S. Patent 5,368,858, issued November 29, 1994, inventor Hunziker
10 describes preparations of biodegradable matrices including TGF- β s as proliferation agents, chemotactic agents, and transforming factors.

U.S. Patent 5,055,447, inventors Palladino et al., issued October 8, 1991, describes methods and
15 compositions for the treatment or prophylaxis of septic shock caused by bacteremic infection. Thus, for example, this patent teaches a therapeutic method for a patient suffering from or at risk of septic shock by administering transforming growth factor- β . Recently,
20 the concept of "sepsis" has been viewed more broadly as an inflammatory condition, and a group of researchers have suggested the designation "systemic inflammatory response syndrome" to describe both "sepsis" (infection by the presence of bacteria in the blood stream) as well
25 as other (non-septic) inflammatory conditions. Chest, 101, pp. 1644-1655 (1992).

Thus, growth factors are useful in a number of therapeutic, clinical, research, diagnostic, and drug design applications. However, as previously mentioned,
30 growth factors are typically large. The natural members of the transforming growth factor- β family range upwards of 25 KDa molecular weight. Clinical uses of growth factors, including TGF- β s, may be limited because of their size, such as due to causing immune responses. For
35 example, human TGF- β 1 is a 25,000 dalton homodimeric protein. In addition to possible adverse immunological responses, large proteins are not often the best

candidates for drugs because of the difficulties in administration and delivery.

Consequently, small peptide mimics of natural growth factors which would avoid most of these problems would be desirable for applications including those to which TGF- β has been put or suggested. It would be advantageous to have small peptides mimicking the biological activity of the large, natural members since small peptides on a mole per mole basis would require much smaller net amounts for administration, and topical applications would be more feasible. Also, quite small peptides would tend to have little or no adverse immunological responses, and could be synthesized easily using simple peptide chemistry procedures.

15

Summary of the Invention

The present invention describes the characterization, properties, and uses of peptide mimics for TGF- β . Because the critical feature for TGF- β activity is the peptide's ability to maintain a stable β -bend structure under physiologic conditions, the peptides of the present invention are collectively referred to as β -bend peptides.

The β -bend peptides have the following initial amino acid sequence, $AA_i-AA_{i+1}-AA_{i+2}$ wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine or isoleucine, and AA_{i+2} is alanine. This sequence is either immediately followed by or has proximal thereto an amino acid with a carbonyl containing side-chain. Because the amino acid residue with a carbonyl containing side-chain does not necessarily have to be immediately followed by, that is adjacent to AA_{i+2} , it is referred to as AA_{i+n} where n is an integer equal to or greater than three. If n is greater than 3, then $n-3$ ("n minus 3") amino acid residues would be between AA_{i+2} and AA_{i+n} in the peptide sequence.

35

For example, the novel β -bend peptide may include the sequence $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}$ where AA_i through

AA_{i+2} are as before and AA_{i+3} is the amino acid residue with a carbonyl containing side-chain (and is glutamic acid, aspartic acid, glutamine, or asparagine). Another example is the sequence AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4} where the
5 residue with the carbonyl containing side-chain is not immediately adjacent, but instead is proximal to, the initial sequence. In this case, AA_i through AA_{i+2} are as before, AA_{i+3} is any suitable amino acid, and AA_{i+4} is glutamic acid, aspartic acid, glutamine, or asparagine.
10 Yet another example is the sequence AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4}-AA_{i+5}. Here, AA_i through AA_{i+2} are as before, AA_{i+3} and AA_{i+4} are suitable amino acids, and AA_{i+5} is glutamic acid, aspartic acid, glutamine, or asparagine.

The original peptide discovered to have TGF- β
15 activity has been named "cytomodulin." An especially preferred embodiment of the present invention, cytomodulin when added to cells in culture in the concentration range 10^{-9} to 10^{-6} M (1.4 pg/ml to 1400 pg/ml), elicits certain highly specific TGF- β effects in
20 several different cell types. For example, among the effects observed is the inhibition of DNA synthesis in Mv-1-Lu mink lung epithelial cells, the growth and colony formation by NRK-49 F fibroblasts in soft agar, and the induction of increased expression of type I collagen in
25 primary cultures of neo-natal human dermal fibroblasts. Moreover, initial results with human osteogenic sarcoma (HOS) cell line indicate that cytomodulin also may be a mimic for other members of the TGF- β superfamily, such as bone morphogenic proteins (BMPs) and osteogenic protein
30 (OPs), as evidenced by its ability to specifically stimulate markers (alkaline phosphatase and osteonectin) characteristic of the osteoblast phenotype.

The novel β -bend peptides are readily synthesized by techniques known to the art. Thus, in one
35 aspect of the present invention, biologically active peptides are provided for eliciting responses characteristic of those elicited by TGF- β .

In another aspect of the present invention, a composition for tissue repair comprises a biocompatible matrix combined with at least one novel β -bend peptide. The biocompatible matrix may be biodegradable or
5 nonbiodegradable. The peptide or peptides are admixed with or carried on the matrix in an amount effective to promote cell growth. Such matrices are useful in constructing templates for repair of soft and hard tissues, for rapid replacement of lost tissue, and for
10 reconstructive and plastic surgery. Such composites provide and sustain cellular regeneration and can be used in combination with other growth factors although surprising, cytomodulin, an especially preferred peptide embodiment of the invention, induces fibroblast colony
15 formation without the presence of additional growth factors such as epidermal growth factor and platelet-derived growth factor.

In yet another aspect of the present invention, a pharmaceutical formulation is provided comprising at
20 least one of the novel β -bend peptide, and a physiologically compatible carrier.

The novel β -bend peptides of the present invention have biological activity that mimic at least one biological activity of TGF- β s, such as inhibiting DNA
25 synthesis in Mv-1-Lu mink lung epithelial cells, promoting growth and colony formation by NRK-49 F fibroblasts, inducing increased expression of type I collagen, and/or inducing TGF- β expression.

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Various embodiments of this invention provide a peptide of a salt thereof having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further having an AA_{i+n} C-terminal of said sequence, wherein n is 3, 4, or 5 such that said peptide contains n-3 additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid, AA_{i+2} is alanine or N-methyl alanine, and AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend and elicits at least one biological activity of TGF- β , with the proviso that if n=3, the peptide does not have SEQ ID NO:1.

Other embodiments of this invention provide a peptide or peptide salt having an amino acid sequence selected from the group consisting of: Leu-Ile-Ala-Glu-Ala-Lys (SEQ. ID. NO:2), Leu-Ile-Ala-Pro-Glu-Ala (SEQ. ID. NO:3), Leu-Aib-Ala-Glu-Ala-Lys (SEQ. ID. NO:4), Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys (SEQ. ID. NO:5), Leu-Abu-Ala-Glu-Ala-Lys (SEQ. ID. NO:6), Leu-Ile-Ala-Asp-Ala-Lys (SEQ. ID. NO:9), Leu-Ile-Ala-Asn-Ala-Lys (SEQ. ID. NO:10), Leu-Ile-Ala-Glu-Ala-Ala (SEQ. ID. NO:11), Leu-Ile-Ala-Gln-Ala-Lys (SEQ. ID. NO:12), Leu-Ile-Ala-Gly-Gly-Glu (SEQ. ID. NO:13), Leu-Ile-Ala-Gly-Glu-Gly (SEQ. ID. NO:14), and Ala-Asn-Val-Ala-Glu-Lys (SEQ. ID. NO:15).

Other embodiments of this invention provide a composition useful for tissue repair comprising: a biocompatible matrix; and, a peptide or salt thereof having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further having an AA_{i+n} C-terminal of said

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sequence, wherein n is 3, 4, or 5 such that the peptide contains $n-3$ additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid, and AA_{i+2} is alanine, AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, wherein the peptide is other than SEQ ID NO: 1, and whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend, the peptide admixed with or carried on the matrix and in an amount effective to promote cell growth, the matrix optionally including Ala-Asn-Val-Ala-Glu-Asn-Ala (SEQ ID NO:1).

Other embodiments of this invention provide a composition useful for tissue repair, comprising: a biocompatible matrix; and, a biologically active peptide or peptide salt having an amino acid sequence selected from the group consisting of: Leu-Ile-Ala-Glu-Ala-Lys (SEQ. ID. NO:2), Leu-Ile-Ala-Pro-Glu-Ala (SEQ. ID. NO:3), Leu-Aib-Ala-Glu-Ala-Lys (SEQ. ID. NO:4), Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys (SEQ. ID. NO:5), Leu-Abu-Ala-Glu-Ala-Lys (SEQ. ID. NO:6), Leu-Ile-Ala-Asp-Ala-Lys (SEQ. ID. NO:9), Leu-Ile-Ala-Asn-Ala-Lys (SEQ. ID. NO:10), Leu-Ile-Ala-Glu-Ala-Ala (SEQ. ID. NO:11), Leu-Ile-Ala-Gln-Ala-Lys (SEQ. ID. NO:12), Leu-Ile-Ala-Gly-Gly-Glu (SEQ. ID. NO:13), Leu-Ile-Ala-Gly-Glu-Gly (SEQ. ID. NO:14), Ala-Asn-Val-Ala-Glu-Lys (SEQ. ID. NO:15), the biologically active peptide admixed with or carried on the matrix.

Other embodiments of this invention provide a pharmaceutical formulation comprising: a physiologically acceptable carrier; and, a peptide or peptide salt having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further

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having an AA_{i+n} C-terminal of said sequence and wherein n is 3, 4, or 5 such that said peptide contains n-3 additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid, AA_{i+2} is alanine or N-methyl alanine, and AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, wherein the peptide is other than SEQ ID NO: 1, and whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend and elicits at least one biological activity of TGF- β .

Other embodiments of this invention provide a pharmaceutical formulation, comprising: a physiologically acceptable carrier; and, a peptide or peptide salt having an amino acid sequence selected from the group consisting of: Leu-Ile-Ala-Glu-Ala-Lys (SEQ. ID. NO:2), Leu-Ile-Ala-Pro-Glu-Ala (SEQ. ID. NO:3), Leu-Aib-Ala-Glu-Ala-Lys (SEQ. ID. NO:4), Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys (SEQ. ID. NO:5), Leu-Abu-Ala-Glu-Ala-Lys (SEQ. ID. NO:6), Leu-Ile-Ala-Asp-Ala-Lys (SEQ. ID. NO:9), Leu-Ile-Ala-Asn-Ala-Lys (SEQ. ID. NO:10), Leu-Ile-Ala-Glu-Ala-Ala (SEQ. ID. NO:11), Leu-Ile-Ala-Gln-Ala-Lys (SEQ. ID. NO:12), Leu-Ile-Ala-Gly-Gly-Glu (SEQ. ID. NO:13), Leu-Ile-Ala-Gly-Glu-Gly (SEQ. ID. NO:14), and Ala-Asn-Val-Ala-Glu-Lys (SEQ. ID. NO:15).

Other embodiments of this invention provide a pharmaceutical formulation comprising: a TGF- β mimic or salt thereof, other than Ala-Asn-Val-Ala-Glu-Asn-Ala, the mimic having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the mimic further having an AA_{i+n} C-terminal of said sequence and wherein n is 3, 4, or 5 such that said peptide contains n-3 additional amino acid residues in between AA

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$i+2$ and AA_{i+n} , and wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid, AA_{i+2} is alanine or N-methyl alanine, and AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, whereby the mimic elicits at least one biological activity of TGF- β ; and, collagen, a collagen analog, or a collagen mimic.

Brief Description of the Drawings

10 Figure 1 graphically illustrates the inhibition of DNA synthesis of Mv-1-Lu mink lung epithelial cells by cytomodulin;

 Figure 2 are photomicrographs (magnification 500 times) wherein Panel (A) is a control, Panel (B) is
15 with 100 nM cytomodulin, and Panel (C) is 100 nM

cytomodulin plus EGF and PDGF, all five days in soft agar with NRK-49 F normal rat kidney fibroblasts;

Figure 3 graphically illustrates the modulation of gene expression in HOS cells by cytomodulin, where
5 Panels (A), (B), and (D) show increased expression while Panel (C) modulated activity depending on concentration, which is however quite characteristic of TGF- β in cells;

Figure 4 having panels (A) through (D) are Northern Blots corresponding to the data graphically
10 illustrated by Fig. 3 and its respective panels, (A) - (D);

Figure 5 gives the atomic coordinates for atom numbers 1-101 of the cytomodulin embodiment of the invention;

Figure 6 shows growth inhibition curves of
15 MV-1-Lu cells by cytomodulin analogs, Leu-Ile-Ala-Pro-Glu-Ala (SEQ. ID. NO:3 or "L1") and Leu-Ile-Ala-Glu-Ala-Lys (SEQ. ID. NO:2 or "L2"); and

Figure 7 shows the induction of collagen $\alpha 1(I)$ by cytomodulin analogs, L1 and L2, and TGF- β in neonatal
20 human dermal fibroblasts.

Description of the Preferred Embodiments

Small peptide mimics of TGF- β have been prepared. Because the critical feature for TGF- β
25 activity is a peptide's ability to maintain a stable β -bend structure under physiologic conditions, the peptides of the present invention are collectively referred to as β -bend peptides.

The inventive peptides have the following
30 initial amino acid sequence, $AA_i-AA_{i+1}-AA_{i+2}$ wherein AA_i is alanine, asparagine, or leucine, AA_{i+1} is valine or isoleucine, and AA_{i+2} is alanine. This sequence is either immediately followed by or has proximal thereto an amino acid with a carbonyl containing side-chain. Because the amino acid residue with a carbonyl containing
35 side-chain does not necessarily have to be immediately followed by, that is adjacent to AA_{i+2} , it is referred to as AA_{i+n} where n is an integer equal to or greater than three. If n is greater than 3,

then $n-3$ ("n minus 3") amino acid residues would be between AA_{i+2} and AA_{i+n} in the peptide sequence.

In one embodiment of the invention n equals three and the amino acid residue with the carbonyl containing side-chain is thus AA_{i+3} . This results in peptides having the following sequence $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}$ wherein AA_i , AA_{i+1} , and AA_{i+2} are as above and AA_{i+3} is glutamic acid, aspartic acid, glutamine, or asparagine. The original peptide with TGF- β activity and an especially preferred embodiment of the present invention, cytomodulin, Ala-Asn-Val-Ala-Glu-Asn-Ala (SEQ ID. NO:1) is of this class. Here, $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}$ corresponds to the second through fifth residues from the N-terminus of cytomodulin, -Asn-Val-Ala-Glu-. Other preferred embodiment are the peptides, Leu-Ile-Ala-Glu-Ala-Lys (SEQ ID. NO:2) and Leu-Ile-Ala-Glu-Ala-Ala (SEQ ID. NO:11). In these examples, $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}$ corresponds to the first four residues of the peptide, Leu-Ile-Ala-Glu-.

In another embodiment, n equals four and the amino acid residue with the carbonyl containing side-chain is thus AA_{i+4} . The sequence of the peptides of this class is $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4}$. In this series, AA_i , AA_{i+1} , and AA_{i+2} are as above, AA_{i+3} is either proline or glycine, and AA_{i+4} is glutamic acid, aspartic acid, glutamine, or asparagine. A preferred embodiment of this class is Leu-Ile-Ala-Gly-Glu-Gly (SEQ ID. NO:14). An especially preferred embodiment is the peptide, Leu-Ile-Ala-Pro-Glu-Ala (SEQ ID. NO:3). In both examples, the first five N-terminal amino acids correspond to $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4}$.

In yet another embodiment of the present invention, n equals five and the amino acid residue with the carbonyl containing side-chain is thus AA_{i+5} . The sequence of the peptides of this class is $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4}-AA_{i+5}$. In this series, AA_i , AA_{i+1} , and AA_{i+2} are as above, AA_{i+3} and AA_{i+4} are glycine, and AA_{i+5} is glutamic acid, aspartic acid, glutamine, or asparagine. An especially preferred member of this series is the peptide, Leu-Ile-Ala-Gly-Gly-Glu (SEQ

ID. NO:13). In this particular example there is a one to one correspondence between SEQ ID. NO:3 and the sequence $AA_i-AA_{i+1}-AA_{i+2}-AA_{i+3}-AA_{i+4}-AA_{i+5}$.

Peptides of the present invention can be synthesized by various suitable methods that are well known in the art, preferably by solid phase synthesis, manual or automated, as first developed by Merrifield and described by Stewart et al. in *Solid Phase Peptide Synthesis* (1984). Chemical synthesis joins the amino acids in the predetermined sequence starting at the C-terminus. Basic solid phase methods require coupling the C-terminal protected α -amino acid to a suitable insoluble resin support. Amino acids for synthesis require protection on the α -amino group to ensure proper peptide bond formation with the preceding residue (or resin support). Following completion of the condensation reaction at the carboxyl end, the α -amino protecting group is removed to allow the addition of the next residue. Several classes of α -protecting groups have been described, see Stewart et al. in *Solid Phase Peptide Synthesis* (1984), with the acid labile, urethan-based tertiary-butyloxycarbonyl (Boc) being the historically preferred. Other protecting groups, and the related chemical strategies, may be used, including the base labile 9-fluorenylmethyloxycarbonyl (Fmoc). Also, the reactive amino acid sidechain functional groups require blocking until the synthesis is completed. The complex array of functional blocking groups, along with strategies and limitations to their use, have been reviewed by Bodansky in *Peptide Synthesis* (1976) and, Stewart et al. in *Solid Phase Peptide Synthesis* (1984).

Solid phase synthesis is initiated by the coupling of the described C-terminal α -protected amino acid residue. Coupling requires activating agents, such as dicyclohexycarbodiimide (DCC) with or without 1-hydroxybenzo-triazole (HOBT), diisopropylcarbodiimide (DIIPC), or ethyldimethylaminopropylcarbodiimide (EDC).

After coupling the C-terminal residue, the α -amino protected group is removed by trifluoroacetic acid (25% or greater) in dichloromethane in the case of acid labile tertiary-butyloxycarbonyl (Boc) groups. A neutralizing
5 step with triethylamine (10%) in dichloromethane recovers the free amine (versus the salt). After the C-terminal residue is added to the resin, the cycle of deprotection, neutralization and coupling, with intermediate wash steps, is repeated in order to extend the protected
10 peptide chain. Each protected amino acid is introduced in excess (three to five fold) with equimolar amounts of coupling reagent in suitable solvent. Finally, after the completely blocked peptide is assembled on the resin support, reagents are applied to cleave the peptide from
15 the resin and to remove the side chain blocking groups. Anhydrous hydrogen fluoride (HF) cleaves the acid labile tertiary-butyloxycarbonyl (Boc) chemistry groups. Several nucleophilic scavengers, such as dimethylsulfide and anisole, are included to avoid side reactions
20 especially on side chain functional groups.

It is believed that slight amino acid modifications to the β -bend peptide sequences will not affect the peptides' ability to form stable β -bend structures. These modifications include techniques to
25 confer resistance to enzymatic degradation such as adding blocking groups to both the N- and C-terminal residues. Another method for preventing degradation and premature clearance by the renal system is the use of unnatural amino acid substitutes in the peptide sequence. For
30 example, N-methyl-alanine is often substituted for alanine and α -amino isobutyric acid and α -amino butyric acid are substitutes for bulky hydrophobic amino acids. Yet another technique is replacing the L-amino acid residue in the peptide sequence with a D-amino acid
35 counterpart. For example, an alanine may be replaced with D-alanine.

The novel β -bend peptides are believed to find uses as agents for enhancing the survival or inducing the growth of nerve and muscle cells. A β -bend peptide is, of course, useful as a new component of culture media for use in culturing nerve cells *in vitro*. The inventive peptides also have utility as a substitute for the natural cytokines in many fields including: in surgery as agents which promote wound healing and regeneration; in orthopedics in promoting bone repair and implant integration; in dentistry in the repair of bony defects and in implant integration; in cancer chemotherapy and in radiation treatment as cytostatic agents for protection of normal stem cells against cell-cycle specific procedures; in treatment of rheumatoid arthritis; in ophthalmology for the repair of macular injury; in ophthalmology for the treatment of uveitis; as a protective agent for splanchnic artery occlusion reperfusion injury; and, as reagents for research in the biology of growth factors.

Therapeutic compositions of this invention will include novel β -bend peptides in concentrations that depend upon the effective doses required and the modes of administration used. Various therapeutic indications for compositions containing at least one β -bend peptide will readily come to mind. One indication is topical application to incisions or exposed tissue for the promotion of wound healing. The types of wound or other traumata that can be treated include (but are not limited to): first, second, and third degree burns (especially second and third degree); epidermal and internal surgical incisions, including those of cosmetic surgery; wounds, including lacerations, incision, and penetrations; and epidermal ulcers including decubital (bed-sores), diabetic, dental, hemophiliac, and varicose.

Uses may be by a variety of methods, such as systemic administration, topical application, intravenous administration, subcutaneous administration, intra-

peritoneal injection, sub-periosteal injection, intra-tracheal administration, release from polymers or pumps, implants, or release from liposomes. Suitable implants (if using an implanted device) include, for example, gel
5 foam, wax, or microparticle-based implants. Doses used should be sufficient to achieve circulating plasma concentrations of active ingredient that are efficacious. Effective doses may be extrapolated from dose-response curves derived from in vitro or animal model test
10 systems.

The inventive peptides also are useful for inducing growth of bone. Thus, osteogenically effective amounts of at least one β -bend peptide in a pharmaceutically acceptable carrier or excipient can be
15 administered for inducing deposition and maturation of bone at the site. In addition, the peptide or peptides can be admixed with or carried by biomaterials, such as hydroxyapatite for bone generation or repair applications in a method such as is described by U.S. Patent
20 5,158,934, issued October 27, 1992, U.S. Patent 5,208,219, issued May 4, 1993, by compositions such as described in U.S. Patent 5,178,845, issued January 12, 1993.

Such bone repair compositions typically include
25 various calcium phosphate mineral component materials such as, for example, hydroxyapatites commercially available under the designations Synthograft, Tricalcium Phosphate, or Periogras. The hydroxyapatite (or tricalcium phosphate) may be prepared by known methods
30 rather than commercially purchased, such as those disclosed by Termine et al., *Arch. Biochem. Biophys.*, 140, TP307-325 (1970). Such a material can be supplied as a powder with preferred particle sizes typically in the range of about 100-2000 μ .

35 Another therapeutic indication for compositions containing at least one β -bend peptide is in conjunction with matrix forming materials. Preferably, the

formulations include a matrix that is capable of providing a structure for developing bone and cartilage. Potential matrices may be biodegradable or nonbiodegradable, and may be chemically or biologically
5 defined.

For one example, the matrix can be inert, solid and non-porous, such as known and presently used as vessels for cell culture. Another form that may be taken by matrices of this invention is that of soluble
10 polymers. Other suitable matrices for practice of this invention include various polymers and hydrogels. Such composites are useful in constructing templates for repair of soft tissue, for rapid replacement of lost tissue, and for reconstructive and plastic surgery.

15 Composites of the invention can thus be made with resorbable polymers of various kinds, having peptide carried by or grafted onto the lattice of the polymeric material. Of course, polymeric supports that are limited in resorbable properties such as hydroxyethyl
20 methacrylate, polymethylmethacrylate, and N-vinylpyrrolidone methylmethacrylate, as a few examples, are also feasible. The composites can then be implanted in the tissue defect.

Among the known and suitable resorbable
25 hydrogels are combinations of polylactate and polyglycollate. Compounds of the invention can be covalently bound to such materials during synthesis of the polymers themselves or the polymers can be hydrolyzed such that attachment sites are available by irradiating
30 the polymer or by chemically activating the polymer to generate free radicals. Then conventional techniques for grafting, or immobilizing, peptides onto polymer supports can be utilized to prepare inventive composites. Resorbable hydrogels or polymers so prepared are
35 particularly useful for soft tissue reconstructions. For hard tissue reconstructions or repair (e.g., bone repair) it is desirable to combine such water soluble, or

resorbable, polymer species with a bioceramic, such as for example bioglass, aluminum oxide, calcium aluminate, tricalcium phosphate, and hydroxyapatite.

When a β -bend peptide is prepared for
5 administration by mixing with physiologically acceptable carriers, i.e., carriers which are non-toxin to recipients at the dosages and concentrations employed, this will normally entail combining the inventive peptide(s) with buffers, antioxidants such as ascorbic
10 acid, low molecular weight (less than about 10 residues) polypeptides, proteins, amino acids, carbohydrates including glucose or dextrans, chelating agents such as EDTA, and other excipients. The inventive peptide for use in therapeutic administrations must be sterile. This
15 is readily accomplished by filtration through sterile filtration (0.22 micron) membranes.

The novel β -bend peptides may be administered in any pharmacologically acceptable carrier, and depending upon the desired mode of administration, may be
20 formulated along with liquid carrier into liposomes, microcapsules, polymers or wax-based and controlled release preparations, or be formulated into tablet, pill, or capsule forms.

The peptides form pharmaceutically acceptable
25 salts with organic and inorganic acids and can be administered in salt form or the novel β -bend peptide can be amidated. Examples of suitable acids for the formation of pharmaceutically acceptable salts are hydrochloric, sulfuric, phosphoric, acetic, benzoic,
30 citric, malonic, salicylic, malic, fumaric, succinic, tartaric, lactic, gluconic, ascorbic, maleic, benzene-sulfonic, methane- and ethanesulfonic, hydroxymethane- and hydroxyethane-sulfonic.

Salts may also be formed with suitable organic
35 pharmaceutically acceptable base addition salts. These organic bases form a class whose limits are readily understood by those skilled in the art. Merely for

purposes of illustration, the class may be said to include mono-, di-, and trialkylamines, such as methylamine, dimethylamine, and triethylamine; mono-, di-, or trihydroxyalkylamines such as mono-, di-, and triethanolamine; amino acids such as arginine, and lysine; guanidine; N-methyl-glucosamine; N-methyl-glucamine; L-glutamine; N-methylpiperazine; morpholine; ethylenediamine; N-benzylphenethylamine; tris(hydroxymethyl)aminomethane; and the like. (See, for example, "Pharmaceutical Salts," *J. Pharm. Sci.*, 66(1), 1-19 (1977).)

Therapeutic formulations containing at least one β -bend peptide, such as for promoting bone cell growth, may be prepared for storage by mixing at least one of the novel peptides having the desired degree of purity with optional physiologically acceptable carriers, excipients or stabilizers, in the form of lyophilized cake or aqueous solutions. Acceptable carriers, excipients or stabilizers are nontoxic to recipients at the dosages and concentrations employed when administered, and include buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid; low molecular weight (less than about 10 residues) polypeptides; proteins, such as serum albumin, gelatin or immunoglobulins.

Other components can include glycine, glutamine, asparagine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrans; chelating agents such as EDTA; sugar alcohols such as mannitol or sorbitol; salt-forming counterions such as sodium; and/or nonionic surfactants such as TWEEN, PLURONICS or PEG. Yet additional useful components desirably included in therapeutic formulations of cytomodulin are one or more other growth factors, such as, for example, epidermal growth factor (EGF) and platelet-derived growth factor (PDGF).

Initial dosing of the novel β -bend peptides for topical applications such as wound healing should be delivered to the therapeutic site in a concentration of about from 50 to 500 ng/ml and thereafter adjusted in
5 line with clinical experience. Since the inventive compositions both provide and sustain cellular regeneration, a continual application or periodic reapplication of the compositions is indicated. The clinician will be expected to modify the dosage in
10 accordance with clinical experience.

Compositions may be used in the form of a sterile irrigant, preferably in combination with a physiological saline solution, or in the form of ointments or suspensions, preferably in combination with
15 other growth factors as earlier noted, and yet further with collagen, a collagen analogue, or a collagen mimic, such as is described, for example, U.S. Patent 5,354,736, issued October 11, 1994, inventor Bhatnagar.

The compositions also may be
20 impregnated into transdermal patches, plasters, and bandages, preferably in a liquid or semi-liquid form. Automicrobial agents such as silver sulfadiazine should be included in such articles or compositions.

The β -bend peptides also may be administered
25 systemically for the treatment of wounds and similar traumata. Systemic administration is useful provided that there are no, or limited, undesirable side-effects, such as the stimulation of neoplastic cellular growth in patients with cancer. Compositions for systemic
30 administration preferably are formulated as sterile, isotonic parenteral injections or infusions.

The inventive compositions, as earlier described, either with a β -bend peptide alone or in combination with other growth factors, collagen,
35 physiologically acceptable carriers, excipients, or stabilizers as described, may be carried by (or admixed with) a biologically compatible matrix. Matrices of the

invention can be porous, and in bead, particulate, or fibrous forms. For example, calcium phosphate materials, such as apatite-based ceramics, have been suggested for producing porous tissue implants or prosthesis materials with micropores sufficient to permit tissue attachment. Thus, a therapeutic application for the compositions of the invention is where the matrix forming material is biodegradable and can be used, for example, in cartilage repair.

Matrix materials useful for filling or otherwise dressing a defect in the cartilage include, for example, fibrinogen (activated with thrombin to form fibrin in the defect or lesion), collagen, gelatin or any other biodegradable material which forms a matrix with pores sufficiently large to allow repair cells to populate and proliferate within the matrix and which can be degraded and replaced with cartilage during the repair process.

The matrices useful in the compositions and methods of this invention may be preformed or may be formed in situ, for example, by polymerizing compounds and compositions such as fibrinogen to form a fibrin matrix. Matrices that may be preformed can include collagen, collagen analogues or collagen mimics (e.g., collagen sponges and collagen fleece), chemically modified collagen, gelatin beads or sponges, gel-forming substances, any other gel forming or composite substance that is composed of a biodegradable matrix material that will fill the tissue or bone defect and allow repair cells to populate the matrix, or mixtures of the above.

Biological activities of the inventive peptides will now be further illustrated by the following examples, which are intended to be illustrative and not limiting.

EXAMPLE 1**Inhibition of DNA Synthesis of Mv-1-Lu
Mink Lung Epithelial Cells**

5 The effect of TGF- β and cytomodulin were
evaluated by determining the rate of [3 H]thymidine
incorporation into total acid-insoluble DNA and cell
10 number. See generally, Sampath et al., *Journal of
Biological Chemistry*, 267, pp. 20352-20362 (1992). DNA
synthesis rates were determined in triplicate cultures
after 24 hour treatment with various concentrations (10^{-9}
M to 10^{-6} M) of either TGF- β or cytomodulin (which was
15 synthesized by the Merrifield method) by adding
[methyl- 3 H]thymidine (2 μ Ci/ml, 80 Ci/mmol) for 6 hours
before the termination of the culture. Incorporation was
terminated by aspiration of the medium, and after washing
three times with phosphate-buffered saline, the
20 trichloroacetic acid (10%)-precipitated radioactive DNA
was extracted with 1.0% (w/v) sodium dodecyl sulfate, 0.1
M NaOH and quantitated by liquid scintillation counting.
For cell number determination, 1×10^5 cells were plated in
flasks in MEM containing 10% FBS, and after 24 hours, the
25 growth medium was replaced with serum-free medium
containing various concentrations of TGF- β and cytomodulin.
Triplicate cultures were harvested every 24 hours for the
duration of 7 days, and the cell number was determined by
counting cells released by trypsin digestion in a fixed
30 volume hemacytometer.

The growth inhibition curve for cytomodulin
were similar to that observed for TGF- β at the same
concentration range.

EXAMPLE 2**Growth and Colony Formation by NRK-49 F
Fibroblasts in Soft Agar**

5 The original assay for TGF- β , the ability to promote anchorage independent growth of normal fibroblasts is still one of the hallmarks of TGF- β activity. NRK-49 F fibroblasts were grown at 37°C in DEM supplemented with 10% fetal calf serum. The experiments were performed with culture medium, 10 ng/mg epidermal growth factor (EGF), and 10 ng/ml platelet-derived growth factor (PDGF); however, unlike TGF- β , which does not induce colony formation in the absence of these factors (see, for example, Massagu, *J. Biol. Chem.*, 259, pp. 9756-9761 (1984)), cytomodulin did induce colony formation without these two growth factors. To this, either 100 nM TGF- β (positive control) or 100 nM cytomodulin was added. NRK-49 F fibroblasts (5×10^4 cells/ml) were mixed with 0.3% agar were plated on the bottom of 35 mm culture dishes. Colony formation was observed starting on day 3 of culture.

As expected no colonies were formed in those cultures containing only the basic medium. Also, as expected, colonies with TGF- β grew colonies. Surprisingly, the cytomodulin cultures also formed colonies to approximately the same extent as the TGF- β cultures. The growth characteristics of the colonies over time were similar between TGF- β and cytomodulin cultures.

With reference to Fig. 2, photomicrographs are illustrated that were taken on day 5 of the fibroblast culturing. Colony formation was actually observed starting on day 3 of culture. As seen in Fig. 2(A), few cells survived culture in the absence of any growth factors. Fig. 2, Panel (B) and Panel (C) show the

formation of small colonies (arrows) in the presence of cytomodulin, with Panel (C) also including EGF and PDGF, which induced much larger colonies (arrows). This is analogous to the induction of colony formation by TGF- β ,
5 except that TGF- β requires the concomitant presence of epidermal growth factor (EGF) and platelet derived growth factor (PDGF); however, as seen by Panel (B), cytomodulin did induce colony formation by itself.

10

EXAMPLE 3

RNA Isolation and Northern Analysis

15 Total cellular RNA was isolated using essentially the method described by Maniatis. Sambrook et al., *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratory Press, Second Edition (1989). Cells were lysed with 0.5% SDS and 0.1 potassium acetate.
20 The lysate was extracted with phenol and centrifuged at 5000 rpm for 15 minutes. The aqueous phase was precipitated with 2 volumes ethanol in 0.1 M Tris, pH 8.0 and 0.2 M NaCl. The pellet was resuspended and quantitated by measuring the ultraviolet absorbance at
25 260 nm. RNA purity was assessed by comparing the ultraviolet absorbance at 260 nm with that at 280 nm.

RNA (10 μ g/lane) was electrophoresed at 3 to 4 v/cm through a 0.7% agarose, 2.2 M formaldehyde denaturing gel. RNA was transferred by capillary
30 transfer to nylon membranes. RNA integrity, gel loading, and transfer efficiency were assessed by methylene blue stained 28S and 18 S bands. The filters were baked at 80°C for 2 hours to immobilize the RNA. After baking, the filters were hybridized at 65°C in 0.5 M NaPO₄ buffer,
35 pH 7.0, containing 1 mM EDTA, 7% sodium dodecyl sulfate, and 1% bovine serum albumin. cDNA probes were labeled with dDIG (fluorescent probe) by the random primer

method, using Klenow enzyme. Hybridization for 18 hours at 65°C followed by washing, was performed.

Data were analyzed by scanning digoxigenin-dVTP according to manufacturer's procedure (Boehringer Mannheim Biochemica, DIG DNA labelling kit, Cat. No. 1175033).

EXAMPLE 4

10 Stable β -bend required for activity

Figure 5 shows the atomic coordinates of the bioactive structure of cytomodulin (atoms 1-101). Thus, the structure represented by Fig. 5 describes the β -bend that appears to be necessary for TGF- β activity. Without being limited by theory, if the structure-activity model is correct, analogs having substantially the same structure as cytomodulin will also exhibit TGF- β like activity. Similarly, by exploiting allosteric binding mechanisms, compounds may be synthesized with increased or decreased activity with respect to cytomodulin.

EXAMPLE 5

25 Using the three dimensional structure of cytomodulin (Fig. 5) as a guide, two initial cytomodulin analogs were designed. From studying the structure, the key features appeared to be the -Val-Ala- sequence responsible for the stable β -bend and a negatively charged side-chain shortly thereafter. At this point, the working structure-activity profile was:

- (1) a hydrophobic or neutral amino acid at position i ;
- (2) a branched hydrophobic at position $i+1$;
- 35 (3) a small aliphatic at position $i+2$, where positions $i+1$ and $i+2$ together form the critical β -bend structure; and

(4) a negatively charged side-chain shortly thereafter at either $i+3$ or $i+4$ if $i+3$ is a proline.

To test this hypothesis, two cytomodulin
 5 analogs, Leu-Ile-Ala-Glu-Ala-Lys (SEQ. ID. NO: 2 or L2)
 and Leu-Ile-Ala-Pro-Glu-Ala (SEQ. ID. NO:3 or L1) were
 synthesized and tested. In both peptides, -Val-Ala- was
 replaced by -Ile-Ala- and the first two N-terminal amino
 acid sequence of cytomodulin was replaced with leucine.
 10 In SEQ. ID. NO:3, glutamic acid is at position $i+3$ since
 it is the first side-chain after the β -bend structure.
 In SEQ. ID. NO:4, proline is at position $i+3$. Since
 proline does not have a "side-chain," the glutamic acid
 was placed at position $i+4$.

15 Both cytomodulin analogs, L1 and L2, displayed
 at least as much TGF- β like activity as cytomodulin.
 They promoted the growth of NRK-49F cells in soft agar
 and inhibited the proliferation of MV-1-Lu cells (see
 Fig. 6). They increased the expression of type I
 20 collagen and TGF- β (see Fig. 7) and decreased the
 expression of collagenase in human dermal fibroblasts.
 Moreover, as with cytomodulin, L1 and L2 also increased
 the expression of type I collagen, TGF- β and alkaline
 phosphatase in HOS cells.

25

EXAMPLE 6

Analogues of SEQ. ID. NO:2 and SEQ. ID. NO:3 were
 made to further probe the sequence requirements for TGF- β
 30 like activity. To that end, the following peptides were
 made:

	Leu-Aib-Ala-Glu-Ala-Lys	(SEQ. ID. NO:4)
	Leu-Ile-(Nme-Ala)-Glu-Ala-Lys	(SEQ. ID. NO:5)
	Leu-Abu-Ala-Glu-Ala-Lys	(SEQ. ID. NO:6)
35	Gly-Gly-Gln-Ile-Ala-Asn-Ile	(SEQ. ID. NO:7)
	Glu-Gly-Ile-Ala-Gly-Lys	(SEQ. ID. NO:8)
	Leu-Ile-Ala-Asp-Ala-Lys	(SEQ. ID. NO:9)

	Leu-Ile-Ala-Asn-Ala-Lys	(SEQ. ID. NO:10)
	Leu-Ile-Ala-Glu-Ala-Ala	(SEQ. ID. NO:11)
	Leu-Ile-Ala-Gln-Ala-Lys	(SEQ. ID. NO:12)
	Leu-Ile-Ala-Gly-Gly-Glu	(SEQ. ID. NO:13)
5	Leu-Ile-Ala-Gly-Glu-Gly	(SEQ. ID. NO:14)
	Ala-Asn-Val-Ala-Glu-Lys	(SEQ. ID. NO:15)
	Leu-Ile-Ala-Lys-Gly-Lys	(SEQ. ID. NO:16)

Of the non-standard amino acids, Aib is α -amino isobutyric acid, Nme-Ala is N-methyl alanine and Abu is
10 α -amino butyric acid.

SEQ. ID. NOS:4-6, which are minor variants of SEQ. ID. NO:2, mimicked the biological activities of TGF- β and cytomodulin as shown by the inhibition of the proliferation of Mv-1-Lu epithelial cells and increased
15 expression of collagen I and TGF- β in HOS cells. Sample thymidine incorporation data for SEQ. ID. NOS:4-6 are shown in Table 1.

TABLE 1

**Inhibition of Incorporation of ^3H -thymidine in
MV-1 Lu cells in the Presence of Test Peptides**

		^3H -Radioactivity, 10^3 dpm	
		(% Inhibition)	
		3.57 (-)	
		(IV) LAibAEAK (SEQ. ID. NO:4)	
10	1 nM	2.60	(27%)
	5 nM	2.31	(35%)
	50 nM	1.94	(46%)
	100 nM	1.44	(60%)
	500 nM	1.61	(55%)
		(V) LINmeAEAK (SEQ. ID. NO:5)	
15	1 nM	1.61	(47%)
	5 nM	1.76	(42%)
	50 nM	1.60	(41%)
	100 nM	1.81	(40%)
	500 nM	1.42	(53%)
		(VI) LAbuAEAK (SEQ. ID. NO:6)	
20	1 nM	2.02	(33%)
	5 nM	1.97	(35%)

25

However, SEQ. ID. NOs:7-8 did not display significant TGF- β activity. This was not unexpected given the working model. An amino acid with a negatively charged side-chain is not present in SEQ. ID. NO:7.

30 Although a glutamic acid is present in SEQ. ID. NO:8, it is on the N-terminal side of the β -bend and not on the C-terminal side as with cytomodulin, L1, and L2.

Inhibition of ^3H -thymidine incorporation results for SEQ ID. NOs:9-16 are shown in Table 2. The numbers shown at the various concentration are the ratio of the inhibition rate of the peptide being tested over the inhibition rate of cytomodulin (SEQ. ID. NO:1) at the same concentration. Because cytomodulin inhibits the proliferation of MV-1-Lu cells at least as much as TGF- β ,
40 cytomodulin and not TGF- β was used as a control.

TABLE 2**Growth Inhibition Activity of Inventive Peptides**

		Inhibition in comparison to cytomodulin (Inhib by new peptide/Inhib by cytomodulin)				
5	Peptide Composition	Concentration (nM)	1	10	100	1000
	LIADAK	SEQ. ID. NO:9	0.45	0.87	1.14	1.25
	LIANAK	SEQ. ID. NO:10	1.70	2.00	1.43	3.05
	LIAEAA	SEQ. ID. NO:11	1.16	1.00	1.17	1.85
10	LIAQAK	SEQ. ID. NO:12	0.90	1.10	0.80	1.56
	LIAGGE	SEQ. ID. NO:13	1.10	1.33	1.45	1.90
	LIAGEG	SEQ. ID. NO:14	0.66	0.85	1.41	1.88
	ANVAEK	SEQ. ID. NO:15	--	0.80	1.00	--
	LIAKGK	SEQ. ID. NO:16	--	0.65	0.67	--

15

As illustrated by Table 2, all peptides with sequences represented by SEQ. ID. NOS:9-16 inhibited at least some amount of thymidine uptake. From these results, the general model for preparing peptides with

20 TGF- β activity appears to work surprisingly well.

One peptide in particular is especially noteworthy. SEQ. ID. NO:10, Leu-Ile-Ala-Asn-Ala-Lys, inhibited the proliferation of Mv-1-Lu epithelial cells even more than cytomodulin at every concentration tested.

25 Because SEQ. ID. NO:10 does not contain a negatively charged side-chain, the working model clearly had to be redefined. Based upon the inhibition activities of both SEQ. ID. NOS:10 and 12 where asparagine and glutamine replaced glutamic acid, the carbonyl group (C=O) appears

30 to be the critical feature and not necessarily the entire carboxylic acid group (COO-).

Another interesting set of peptides is SEQ. ID. NOS:13-14 which explore the positional requirements of the carbonyl group. The activities of these peptides

appear to be the result of the unusual flexibility of the glycine backbone. Because glycine's backbone may sample virtually all allowable torsional angles, a carbonyl containing side-chain placed proximal to the β -bend structure could attain its necessary conformation without incurring a significant energy cost. Since SEQ. ID. NO:13 would be more flexible than SEQ. ID. NO:14, it is not surprising that SEQ. ID. NO:13 inhibits thymidine uptake more than SEQ. ID. NO:14.

Another noteworthy feature includes SEQ. ID. NO:11. Since this peptide displays at least as much activity as cytomodulin, the C-terminal lysine of L1 and L2 are clearly not important. As a result, the structure activity relationship required for TFG- β activity appears to be:

- (1) a hydrophobic or neutral amino acid at position i ;
- (2) a branched hydrophobic at position $i+1$ (i.e. Val, Ile);
- (3) a small aliphatic at position $i+2$ (i.e. Ala), where positions $i+1$ and $i+2$ together form the critical β -bend structure; and
- (4) a carbonyl containing side-chain shortly thereafter.

25

It is to be understood that while the invention has been described above in conjunction with preferred specific embodiments, the description and examples are intended to illustrate and not limit the scope of the invention, which is defined by the scope of the appended claims.

30

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: The Regents of the University of California
- (ii) TITLE OF INVENTION: Peptide Compositions With Growth Factor-like Activity
- (iii) NUMBER OF SEQUENCES: 16
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- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/US97/18342
 - (B) FILING DATE: 10 October 1997
 - (C) CLASSIFICATION:
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 - (B) TELEFAX: (604) 682-7295

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Ala	Asn	Val	Ala	Glu	Asn	Ala
1				5		

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Leu Ile Ala Glu Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 6 amino acids
 (B) TYPE: amino acid
 (C) STRANDEDNESS:
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Leu Ile Ala Pro Glu Ala
1 5

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 6 amino acids
 (B) TYPE: amino acid
 (C) STRANDEDNESS:
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Leu Xaa Ala Glu Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 6 amino acids
 (B) TYPE: amino acid
 (C) STRANDEDNESS:
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Leu Ile Xaa Glu Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 6 amino acids
 (B) TYPE: amino acid
 (C) STRANDEDNESS:
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Leu Xaa Ala Glu Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Gly Gly Gln Ile Ala Asn Ile
1 5

(2) INFORMATION FOR SEQ ID NO:8:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Glu Gly Ile Ala Gly Lys
1 5

(2) INFORMATION FOR SEQ ID NO:9:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Leu Ile Ala Asp Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:10:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Leu Ile Ala Asn Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:11:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Leu Ile Ala Glu Ala Ala
1 5

(2) INFORMATION FOR SEQ ID NO:12:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Leu Ile Ala Gln Ala Lys
1 5

(2) INFORMATION FOR SEQ ID NO:13:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Leu Ile Ala Gly Gly Glu
1 5

(2) INFORMATION FOR SEQ ID NO:14:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Leu Ile Ala Gly Glu Gly
1 5

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Ala Asn Val Ala Glu Lys
1 5

(2) INFORMATION FOR SEQ ID NO:16:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 6 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS:
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Leu Ile Ala Lys Gly Lys
1 5

It is Claimed:

1. A peptide of a salt thereof having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further having an AA_{i+n} C-terminal of said sequence, wherein n is 3, 4, or 5 such that said peptide contains $n-3$ additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein

AA_i is alanine, asparagine, or leucine,

AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid,

AA_{i+2} is alanine or N-methyl alanine, and

AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend and elicits at least one biological activity of TGF- β , with the proviso that if $n=3$, the peptide does not have SEQ ID NO:1.

2. The peptide as in claim 1, wherein n is 3 and the peptide sequence further comprises $AA_{i+4}-AA_{i+5}$ wherein AA_{i+4} and AA_{i+5} are alanine.

3. The peptide as in claim 1, wherein n is 4 and the amino acid residue in between AA_{i+2} and AA_{i+n} is proline or glycine.

4. The peptide as in claim 1, wherein n is 5 and the two amino acid residues in between AA_{i+2} and AA_{i+n} are glycine.

5. A peptide or peptide salt having an amino acid sequence selected from the group consisting of:

Leu-Ile-Ala-Glu-Ala-Lys	(SEQ. ID. NO:2),
Leu-Ile-Ala-Pro-Glu-Ala	(SEQ. ID. NO:3),
Leu-Aib-Ala-Glu-Ala-Lys	(SEQ. ID. NO:4),
Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys	(SEQ. ID. NO:5),
Leu-Abu-Ala-Glu-Ala-Lys	(SEQ. ID. NO:6),
Leu-Ile-Ala-Asp-Ala-Lys	(SEQ. ID. NO:9),
Leu-Ile-Ala-Asn-Ala-Lys	(SEQ. ID. NO:10),
Leu-Ile-Ala-Glu-Ala-Ala	(SEQ. ID. NO:11),
Leu-Ile-Ala-Gln-Ala-Lys	(SEQ. ID. NO:12),
Leu-Ile-Ala-Gly-Gly-Glu	(SEQ. ID. NO:13),
Leu-Ile-Ala-Gly-Glu-Gly	(SEQ. ID. NO:14), and
Ala-Asn-Val-Ala-Glu-Lys	(SEQ. ID. NO:15).

6. A composition useful for tissue repair comprising:
a biocompatible matrix; and,

a peptide or salt thereof having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further having an AA_{i+n} C-terminal of said sequence, wherein n is 3, 4, or 5 such that the peptide contains n-3 additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein

AA_i is alanine, asparagine, or leucine,

AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid, and

AA_{i+2} is alanine, AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, wherein the peptide is other than SEQ ID NO: 1, and whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend, the peptide admixed with or carried on the matrix and in an amount effective to promote cell growth, the matrix optionally including Ala-Asn-Val-Ala-Glu-Asn-Ala (SEQ ID NO:1).

7. A composition as in claim 6, wherein n is 3 and the peptide sequence further comprises AA_{i+4} - AA_{i+5} wherein AA_{i+4} and AA_{i+5} are alanine.

8. The composition as in claim 6, wherein n is 4 and the amino acid residue in between AA_{i+2} and AA_{i+n} is proline or glycine.

9. The composition as in claim 6, wherein n is 5 and the two amino acid residues in between AA_{i+2} and AA_{i+n} are glycine.

10. The composition as in any one of claims 6 to 9, wherein the biocompatible matrix is biodegradable.

11. The composition as in claim 10, wherein the matrix includes a resorbable polymer.

12. The composition as in any one of claims 6 to 9, wherein the biocompatible matrix is non-biodegradable.

13. The composition as in claim 12, wherein the matrix is porous.

14. A composition useful for tissue repair, comprising:
a biocompatible matrix; and,
a biologically active peptide or peptide salt having an amino acid sequence selected from the group consisting of:

Leu-Ile-Ala-Glu-Ala-Lys	(SEQ. ID. NO:2),
Leu-Ile-Ala-Pro-Glu-Ala	(SEQ. ID. NO:3),
Leu-Aib-Ala-Glu-Ala-Lys	(SEQ. ID. NO:4),
Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys	(SEQ. ID. NO:5),
Leu-Abu-Ala-Glu-Ala-Lys	(SEQ. ID. NO:6),
Leu-Ile-Ala-Asp-Ala-Lys	(SEQ. ID. NO:9),

Leu-Ile-Ala-Asn-Ala-Lys	(SEQ. ID. NO:10),
Leu-Ile-Ala-Glu-Ala-Ala	(SEQ. ID. NO:11),
Leu-Ile-Ala-Gln-Ala-Lys	(SEQ. ID. NO:12),
Leu-Ile-Ala-Gly-Gly-Glu	(SEQ. ID. NO:13),
Leu-Ile-Ala-Gly-Glu-Gly	(SEQ. ID. NO:14),
Ala-Asn-Val-Ala-Glu-Lys	(SEQ. ID. NO:15),

the biologically active peptide admixed with or carried on the matrix.

15. The composition as in claim 14, wherein the biocompatible matrix is biodegradable.

16. The composition as in claim 15, wherein the matrix includes a resorbable polymer.

17. The composition as in claim 14, wherein the biocompatible matrix is non-biodegradable.

18. The composition as in claim 17, wherein the matrix is porous.

19. A pharmaceutical formulation comprising:
 a physiologically acceptable carrier; and,
 a peptide or peptide salt having the sequence $AA_i - AA_{i+1} - AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the peptide further having an AA_{i+n} C-terminal of said sequence and wherein n is 3, 4, or 5 such that said peptide contains n-3 additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein

AA_i is alanine, asparagine, or leucine,

AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid,

AA_{i+2} is alanine or N-methyl alanine, and

AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, wherein the peptide is other than SEQ ID NO: 1, and whereby the peptide forms a stable β -bend structure in physiologic conditions under which TGF- β has a stable β -bend and elicits at least one biological activity of TGF- β .

20. The formulation as in claim 19, wherein n is 3 and the peptide sequence further comprises AA_{i+4} - AA_{i+5} wherein AA_{i+4} and AA_{i+5} are alanine.

21. The formulation as in claim 19, wherein n is 4 and the amino acid residue in between AA_{i+2} and AA_{i+n} is proline or glycine.

22. The formulation as in claim 19, wherein n is 5 and the two amino acid residues in between AA_{i+2} and AA_{i+n} are glycine.

23. The formulation as in any one of claims 19 to 22, wherein the peptide is in the form of a salt.

24. A pharmaceutical formulation, comprising:
a physiologically acceptable carrier; and,
a peptide or peptide salt having an amino acid sequence selected from the group consisting of:

Leu-Ile-Ala-Glu-Ala-Lys	(SEQ. ID. NO:2),
Leu-Ile-Ala-Pro-Glu-Ala	(SEQ. ID. NO:3),
Leu-Aib-Ala-Glu-Ala-Lys	(SEQ. ID. NO:4),
Leu-Ile-(N-methyl-Ala)-Glu-Ala-Lys	(SEQ. ID. NO:5),
Leu-Abu-Ala-Glu-Ala-Lys	(SEQ. ID. NO:6),
Leu-Ile-Ala-Asp-Ala-Lys	(SEQ. ID. NO:9),
Leu-Ile-Ala-Asn-Ala-Lys	(SEQ. ID. NO:10),
Leu-Ile-Ala-Glu-Ala-Ala	(SEQ. ID. NO:11),
Leu-Ile-Ala-Gln-Ala-Lys	(SEQ. ID. NO:12),

Leu-Ile-Ala-Gly-Gly-Glu	(SEQ. ID. NO:13),
Leu-Ile-Ala-Gly-Glu-Gly	(SEQ. ID. NO:14), and
Ala-Asn-Val-Ala-Glu-Lys	(SEQ. ID. NO:15).

25. The formulation as in claim 24, wherein the peptide is in the form of a salt.

26. The formulation of any one of claims 19 to 25, further comprising a peptide of the sequence (SEQ. ID. NO: 1).

27. A pharmaceutical formulation comprising:

a TGF- β mimic or salt thereof, other than Ala-Asn-Val-Ala-Glu-Asn-Ala, the mimic having the sequence $AA_i-AA_{i+1}-AA_{i+2}$ defining the N-terminus or following alanine as the N-terminus, the mimic further having an AA_{i+n} C-terminal of said sequence and wherein n is 3, 4, or 5 such that said peptide contains n-3 additional amino acid residues in between AA_{i+2} and AA_{i+n} , and wherein

AA_i is alanine, asparagine, or leucine,

AA_{i+1} is valine, isoleucine, α -amino isobutyric acid, or α -amino butyric acid,

AA_{i+2} is alanine or N-methyl alanine, and

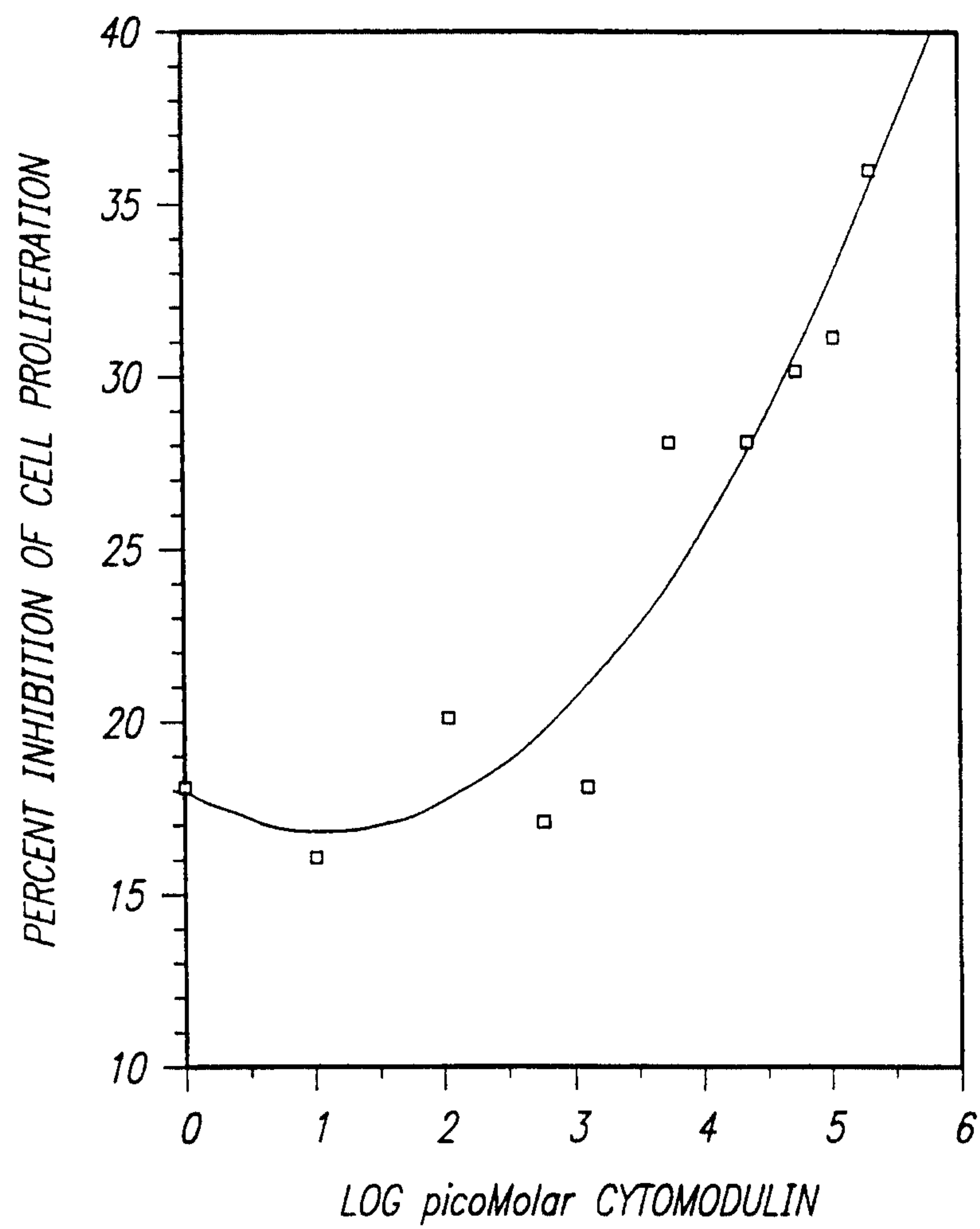
AA_{i+n} is glutamic acid, aspartic acid, glutamine, or asparagine, whereby the mimic elicits at least one biological activity of TGF- β ; and,

collagen, a collagen analog, or a collagen mimic.

28. The pharmaceutical formulation as in claim 27, wherein the collagen analog or collagen mimic has enhanced cell binding with respect to collagen.

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FIG. 1



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FIG. 2A
CONTROL

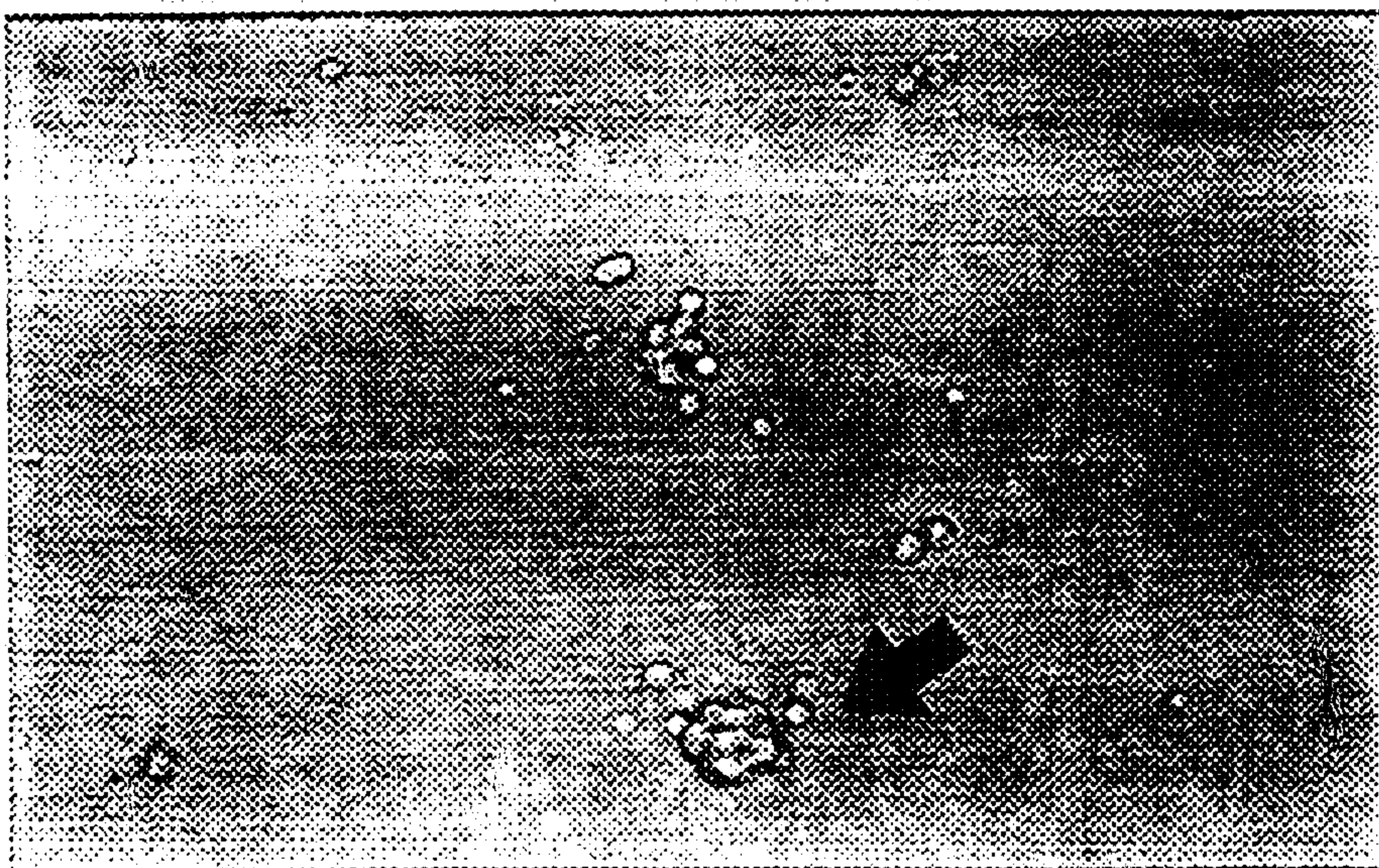
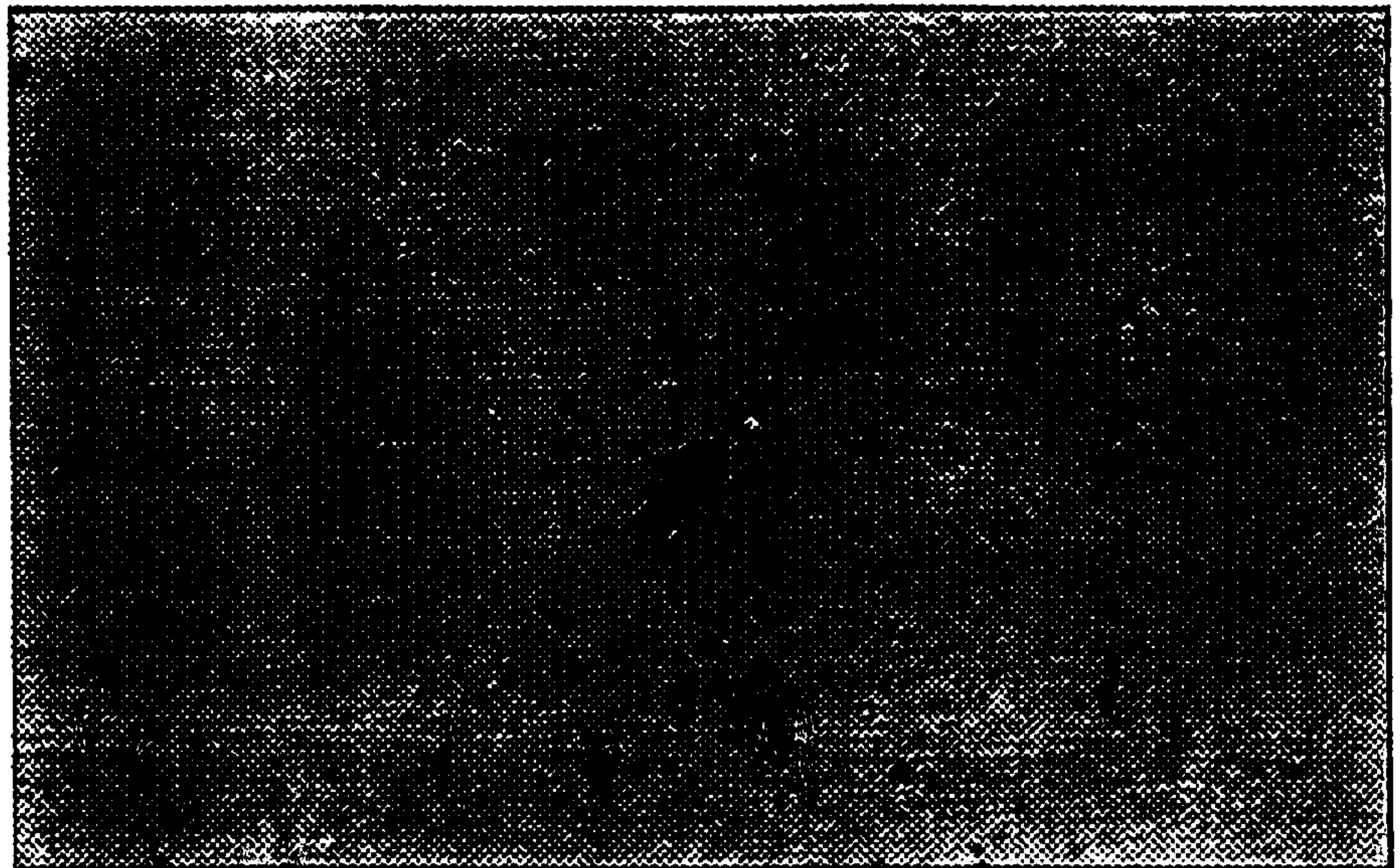
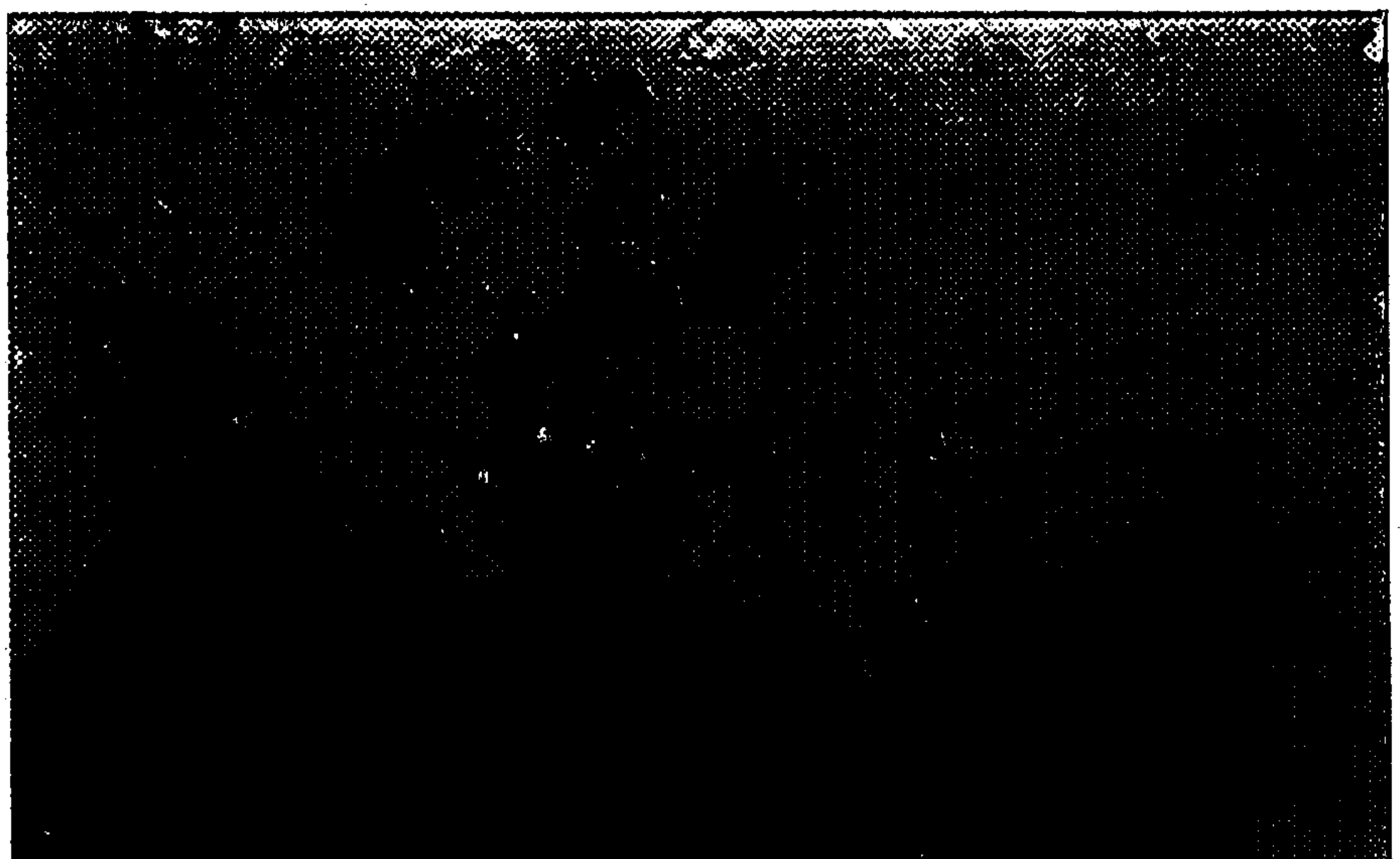


FIG. 2B

FIG. 2C



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FIG. 3A

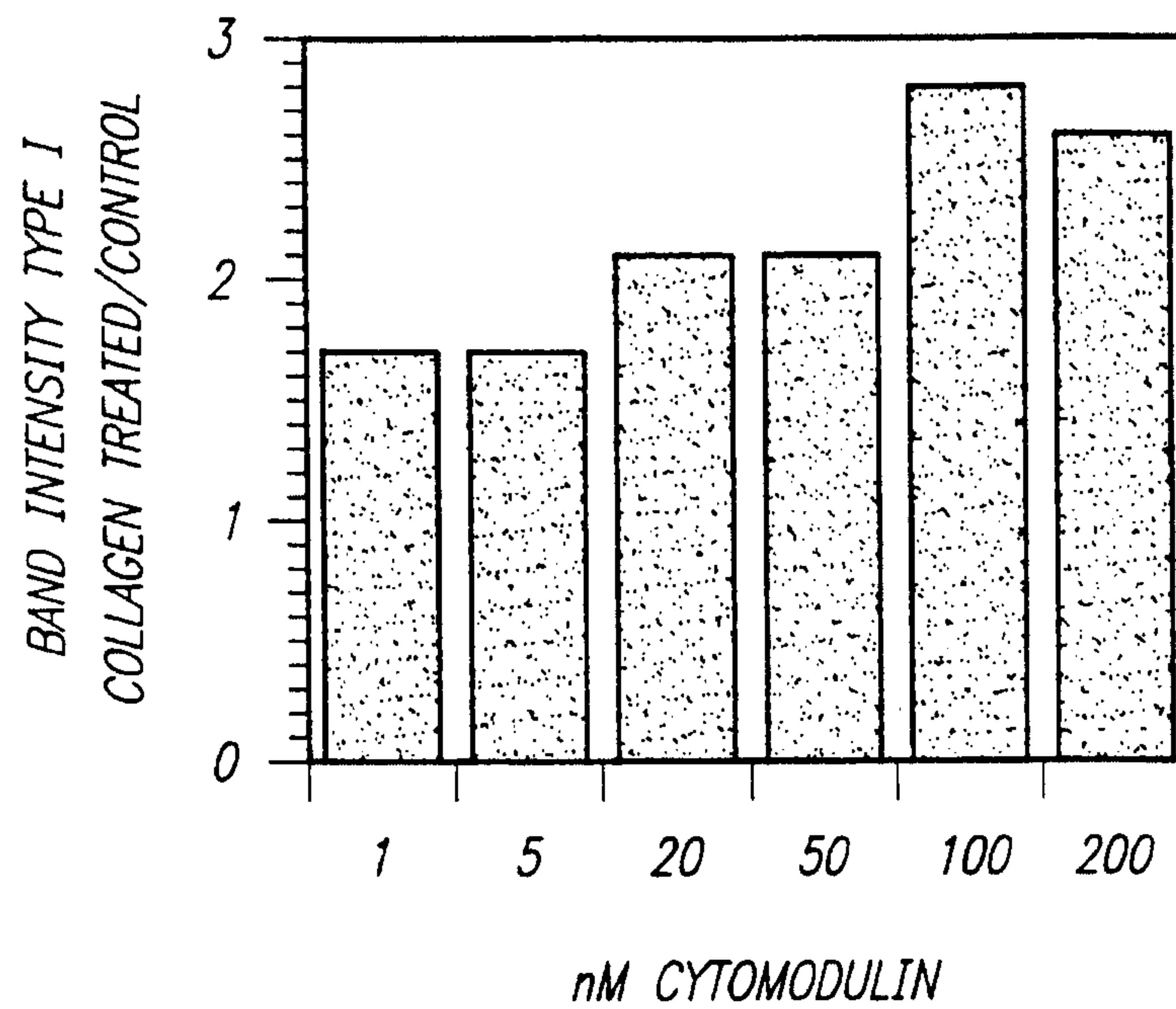
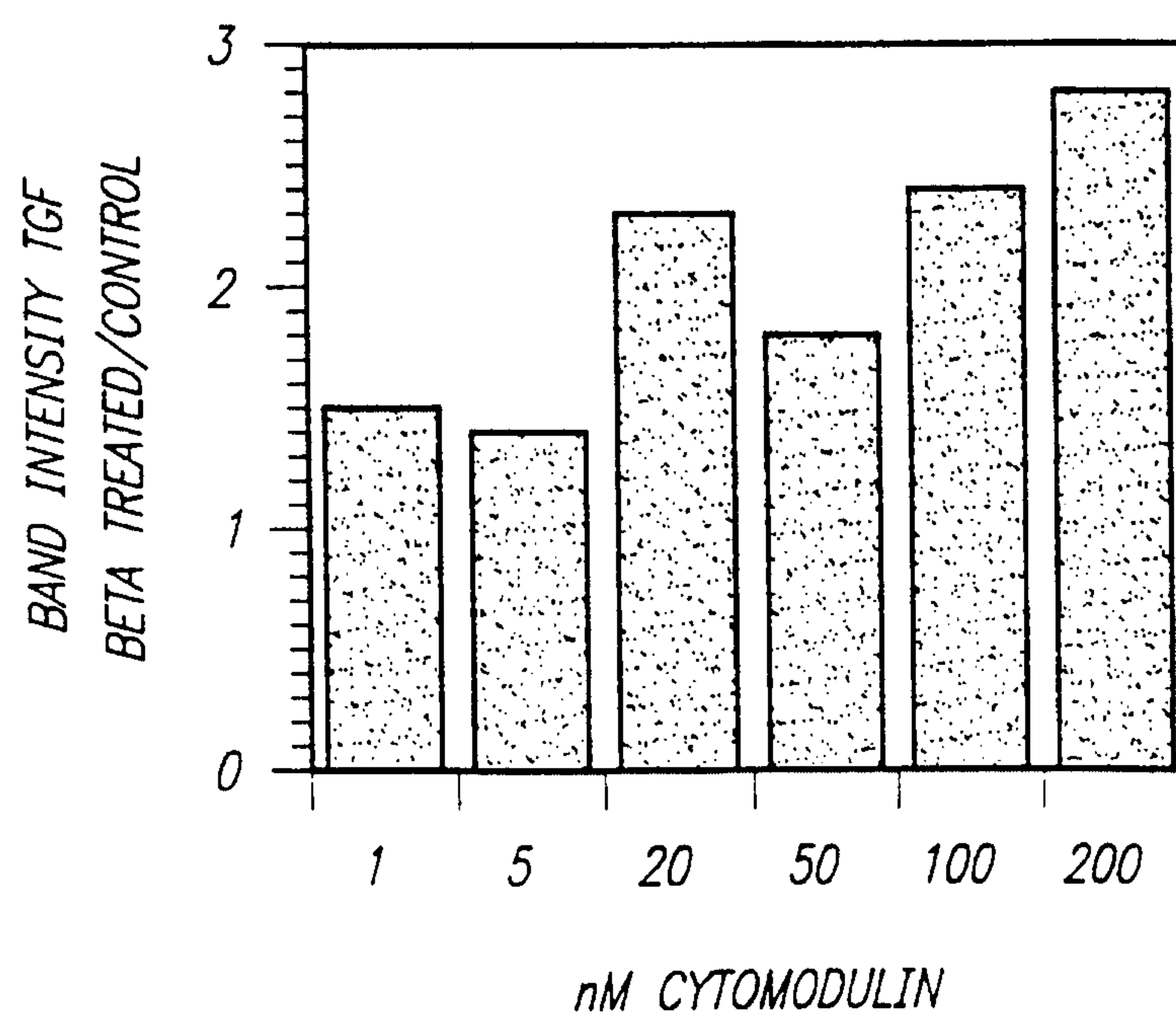


FIG. 3B



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FIG. 3C

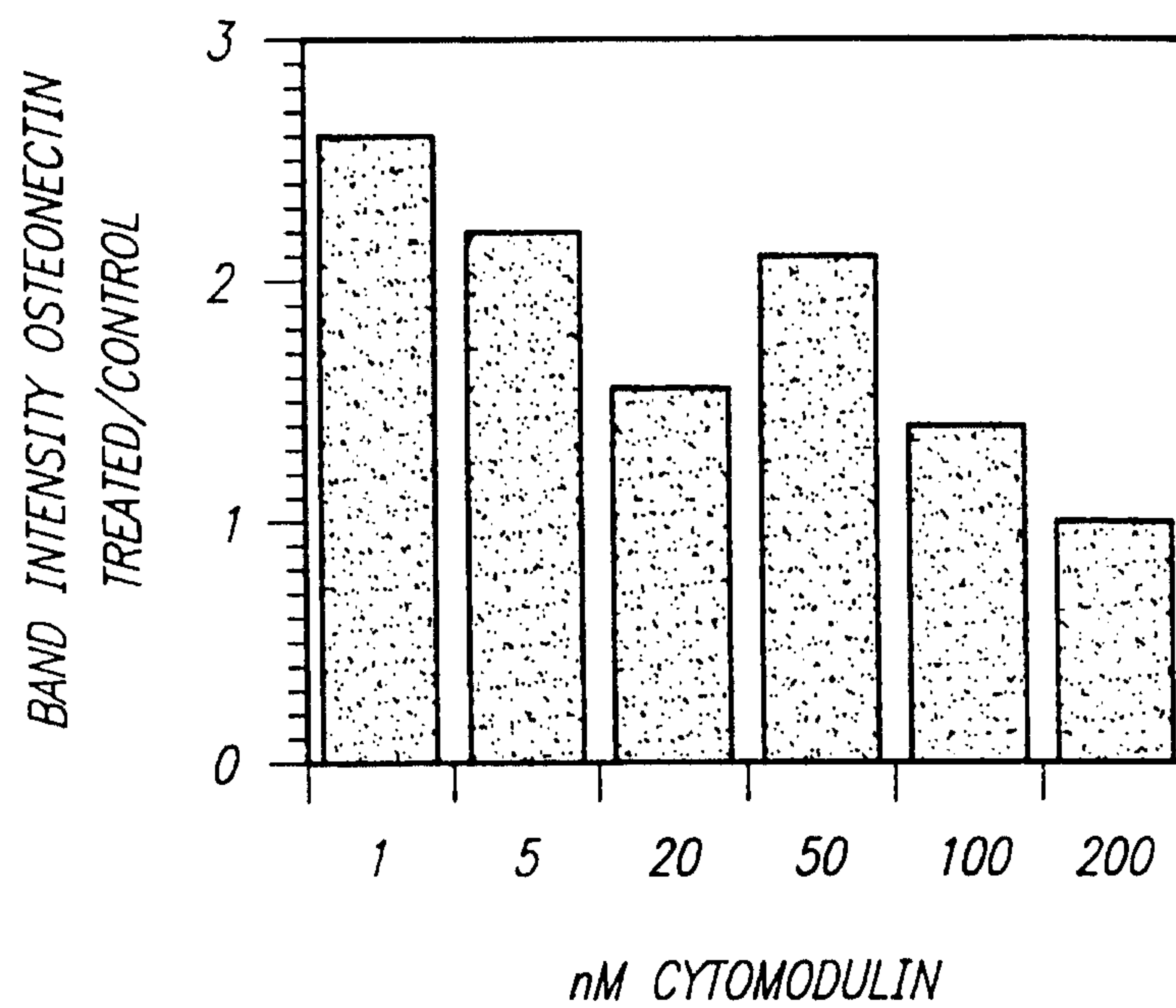


FIG. 3D

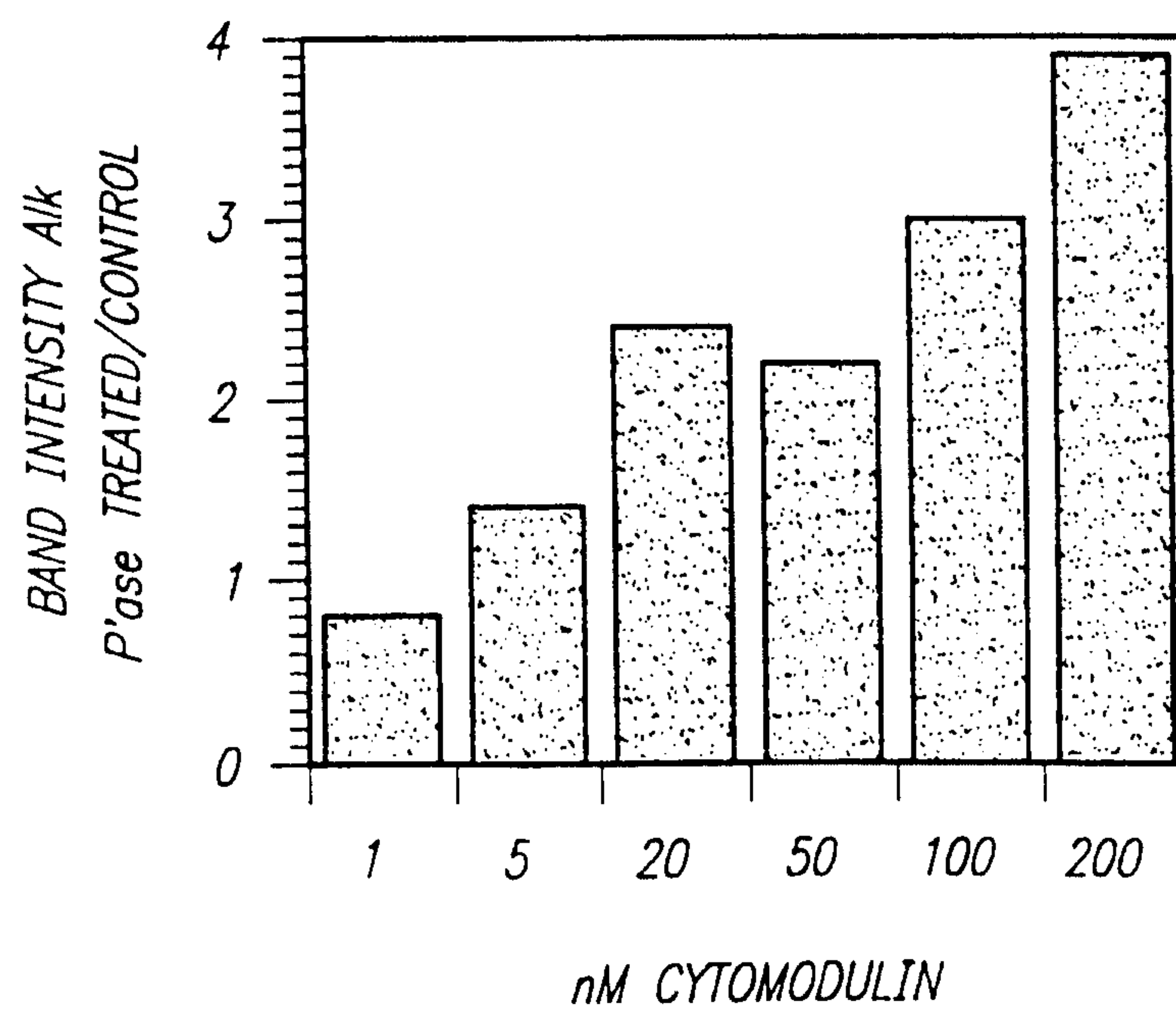


FIG. 4B

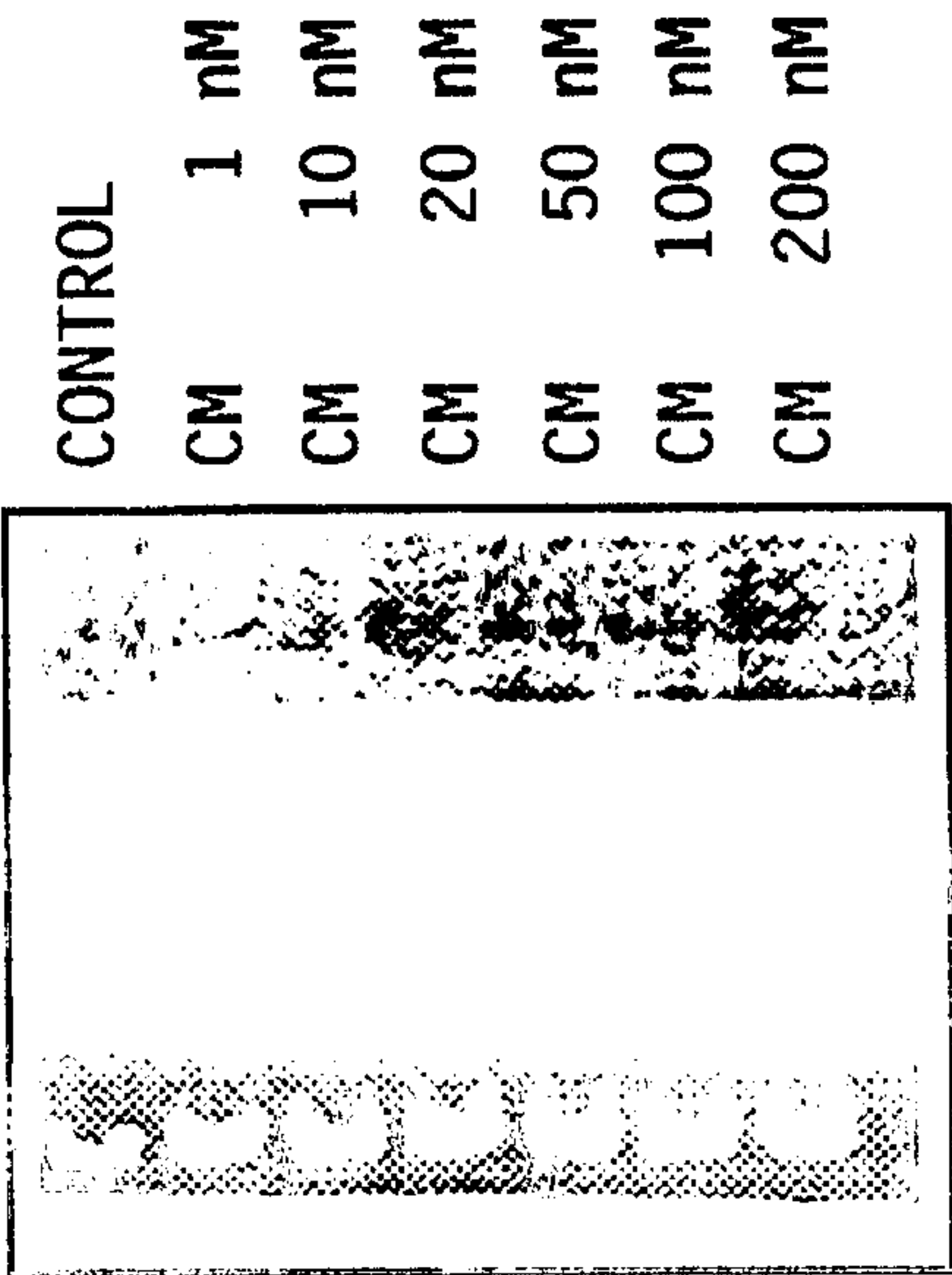


FIG. 4D

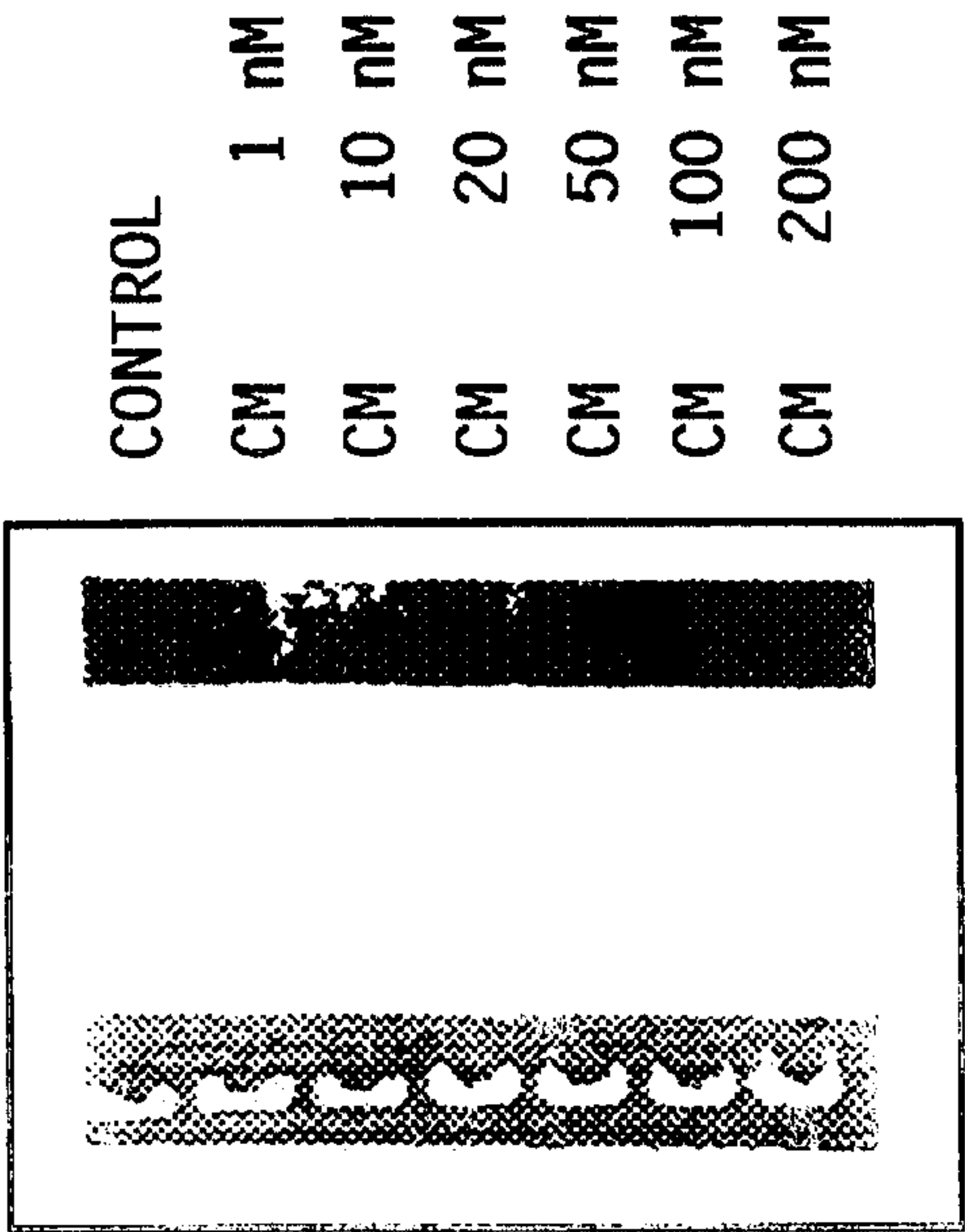


FIG. 4A

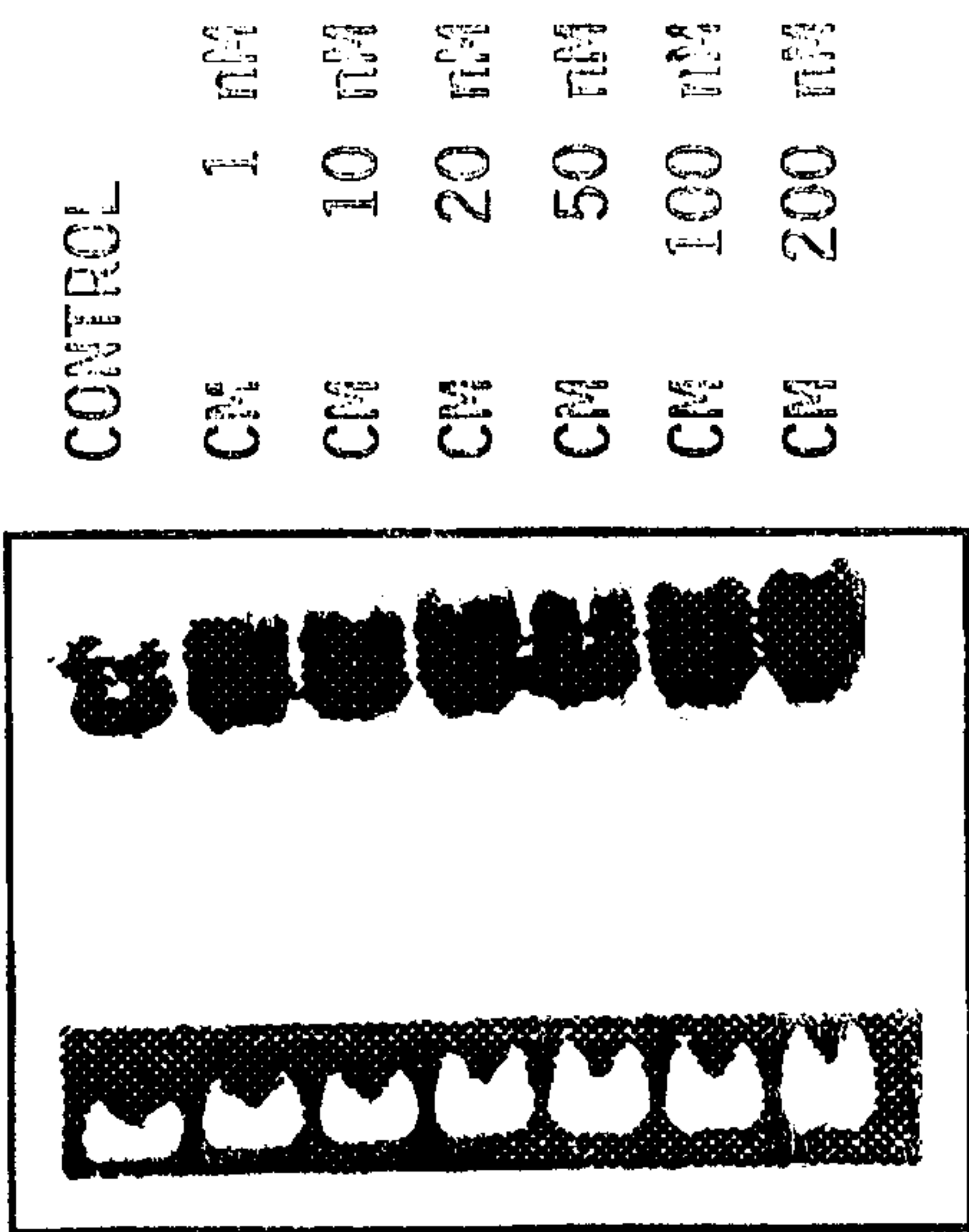
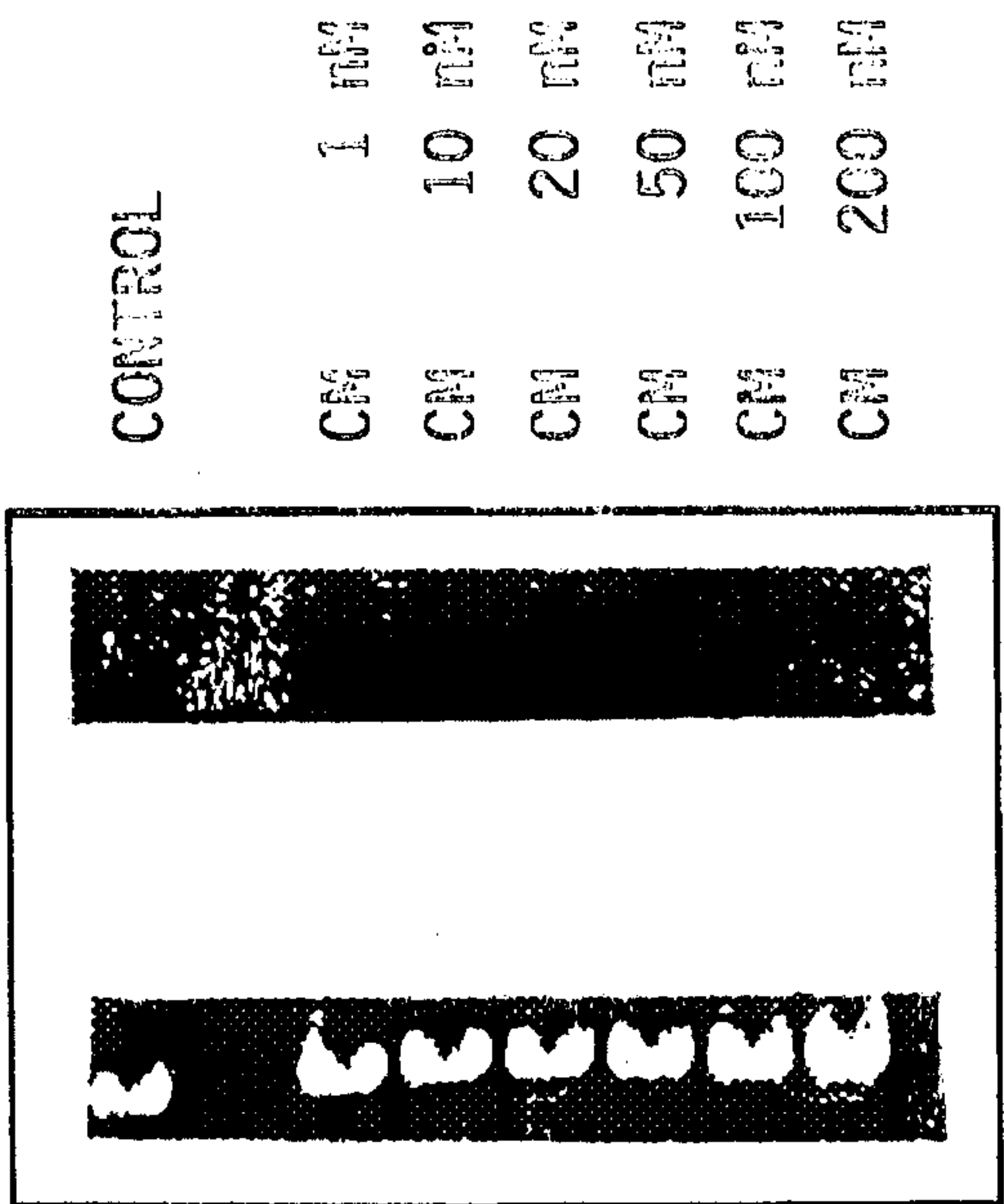


FIG. 4C



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FIG. 5-1

1	H1	ACE	1	1.952	1.249	.316
2	CH3	ACE	1	1.927	2.333	.203
3	H2	ACE	1	1.576	2.782	1.133
4	H3	ACE	1	1.243	2.593	-.605
5	C	ACE	1	3.323	2.853	-.119
6	O	ACE	1	4.272	2.077	-.210
7	N	ALA	2	3.439	4.166	-.310
8	HN	ALA	2	2.632	4.767	-.211
9	CA	ALA	2	4.695	4.857	-.543
10	HA	ALA	2	5.425	4.521	.195
11	CB	ALA	2	5.211	4.572	-1.957
12	HB1	ALA	2	4.475	4.894	-2.694
13	HB2	ALA	2	6.142	5.115	-2.122
14	HB3	ALA	2	5.399	3.506	-2.083
15	C	ALA	2	4.443	6.350	-.361
16	O	ALA	2	3.292	6.788	-.383
17	N	ASN	3	5.504	7.130	-.187
18	HN	ASN	3	6.426	6.699	-.147
19	CA	ASN	3	5.484	8.570	.015
20	HA	ASN	3	4.836	9.029	-.733
21	CB	ASN	3	4.964	8.871	1.432
22	HB2	ASN	3	5.670	8.464	2.158
23	HB3	ASN	3	4.000	8.383	1.578
24	CG	ASN	3	4.760	10.357	1.704
25	OD1	ASN	3	4.723	11.177	.792
26	ND2	ASN	3	4.654	10.739	2.969
27	HND1	ASN	3	4.656	10.052	3.709
28	HND2	ASN	3	4.487	11.714	3.168
29	C	ASN	3	6.925	9.054	-.180
30	O	ASN	3	7.816	8.242	-.407
31	N	VAL	4	7.196	10.354	-.082
32	HN	VAL	4	6.433	11.007	.038
33	CA	VAL	4	8.563	10.863	-.110
34	HA	VAL	4	9.001	10.579	-1.068
35	CB	VAL	4	8.546	12.404	-.038
36	HB	VAL	4	7.922	12.763	-.857
37	CG1	VAL	4	7.958	12.951	1.272
38	HG11	VAL	4	8.590	12.684	2.119
39	HG12	VAL	4	7.900	14.038	1.213
40	HG13	VAL	4	6.955	12.563	1.438
41	CG2	VAL	4	9.949	12.988	-.245
42	HG21	VAL	4	10.376	12.608	-1.174
43	HG22	VAL	4	9.889	14.075	-.306
44	HG23	VAL	4	10.605	12.718	.583

FIG. 5-2

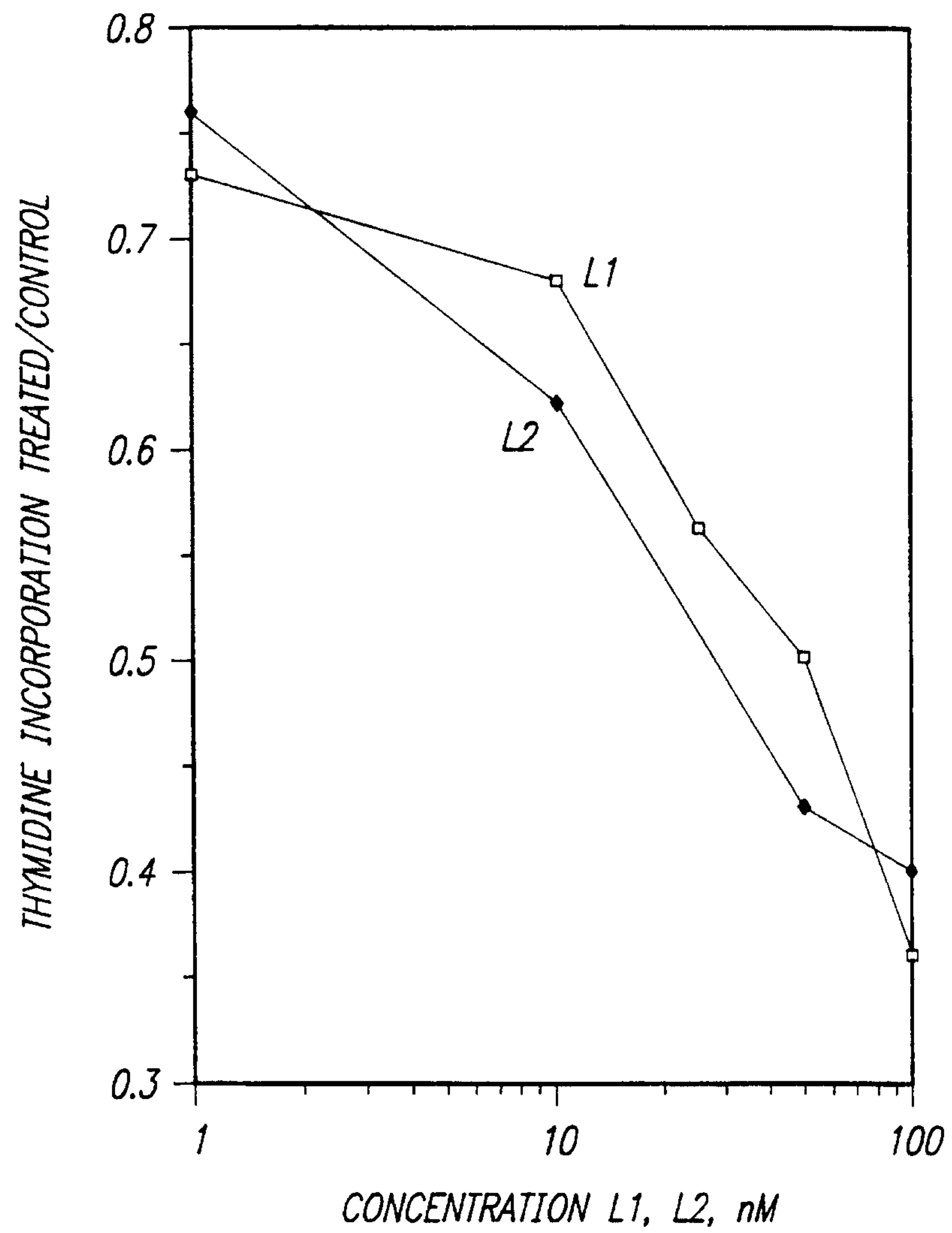
45	C	VAL	4	9.414	10.217	.994
46	O	VAL	4	10.604	9.986	.807
47	N	ALA	5	8.808	9.966	2.157
48	HN	ALA	5	7.829	10.176	2.244
49	CA	ALA	5	9.520	9.482	3.331
50	HA	ALA	5	10.411	10.096	3.479
51	CB	ALA	5	8.626	9.645	4.563
52	HB1	ALA	5	7.723	9.042	4.457
53	HB2	ALA	5	9.169	9.314	5.450
54	HB3	ALA	5	8.350	10.693	4.688
55	C	ALA	5	9.968	8.025	3.185
56	O	ALA	5	11.076	7.684	3.587
57	N	GLU	6	9.079	7.168	2.682
58	HN	GLU	6	8.245	7.512	2.233
59	CA	GLU	6	9.204	5.720	2.725
60	HA	GLU	6	10.254	5.429	2.779
61	CB	GLU	6	8.456	5.210	3.964
62	HB2	GLU	6	7.393	5.439	3.867
63	HB3	GLU	6	8.837	5.712	4.854
64	CG	GLU	6	8.602	3.703	4.203
65	HG2	GLU	6	9.655	3.443	4.313
66	HG3	GLU	6	8.176	3.138	3.375
67	CD	GLU	6	7.861	3.290	5.467
68	OE1	GLU	6	6.706	3.745	5.612
69	OE2	GLU	6	8.468	2.548	6.266
70	C	GLU	6	8.583	5.188	1.438
71	O	GLU	6	7.633	5.781	.930
72	N	ASN	7	9.123	4.107	.882
73	HN	ASN	7	9.827	3.588	1.392
74	CA	ASN	7	8.673	3.542	-.382
75	HA	ASN	7	7.595	3.676	-.478
76	CB	ASN	7	9.390	4.236	-1.545
77	HB2	ASN	7	10.466	4.108	-1.419
78	HB3	ASN	7	9.155	5.301	-1.530
79	CG	ASN	7	8.962	3.664	-2.892
80	OD1	ASN	7	7.842	3.197	-3.055
81	ND2	ASN	7	9.840	3.699	-3.887
82	HND1	ASN	7	10.757	4.091	-3.746
83	HND2	ASN	7	9.556	3.315	-4.774
84	C	ASN	7	8.974	2.049	-.371
85	O	ASN	7	9.950	1.634	.253
86	N	ALA	8	8.148	1.245	-1.035
87	HN	ALA	8	7.449	1.658	-1.644

FIG. 5-3

88	CA	ALA	8	8.215	-.205	-1.003
89	HA	ALA	8	9.244	-.529	-.837
90	CB	ALA	8	7.341	-.717	.150
91	HB1	ALA	8	6.300	-.440	-.024
92	HB2	ALA	8	7.415	-1.800	.241
93	HB3	ALA	8	7.675	-.269	1.087
94	C	ALA	8	7.762	-.755	-2.355
95	O	ALA	8	7.562	-.002	-3.304
96	N	NME	9	7.602	-2.077	-2.453
97	HN	NME	9	7.783	-2.641	-1.638
98	CT	NME	9	7.166	-2.736	-3.673
99	HT1	NME	9	6.431	-2.126	-4.201
100	HT2	NME	9	8.026	-2.904	-4.322
101	HT3	NME	9	6.712	-3.695	-3.424

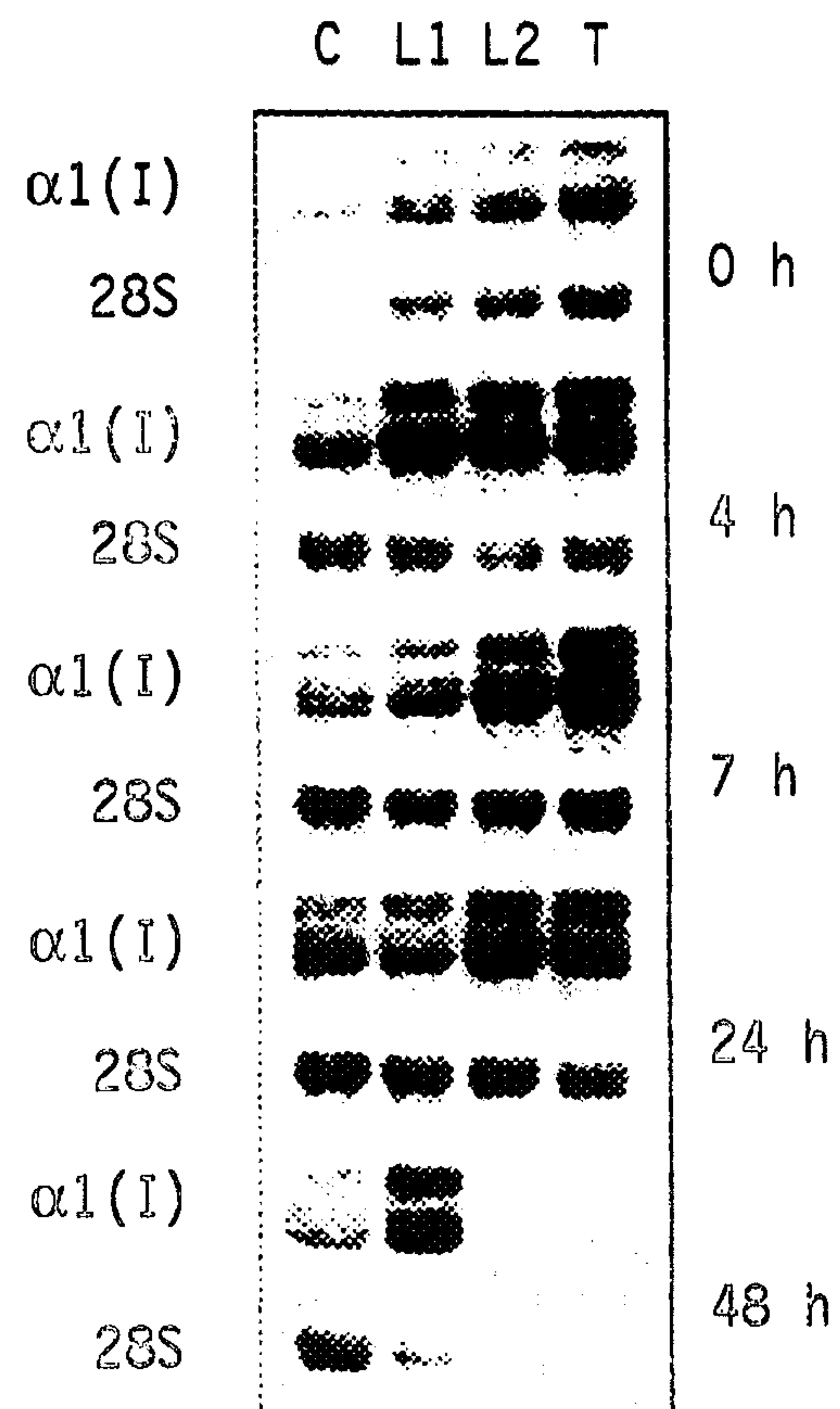
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FIG. 6

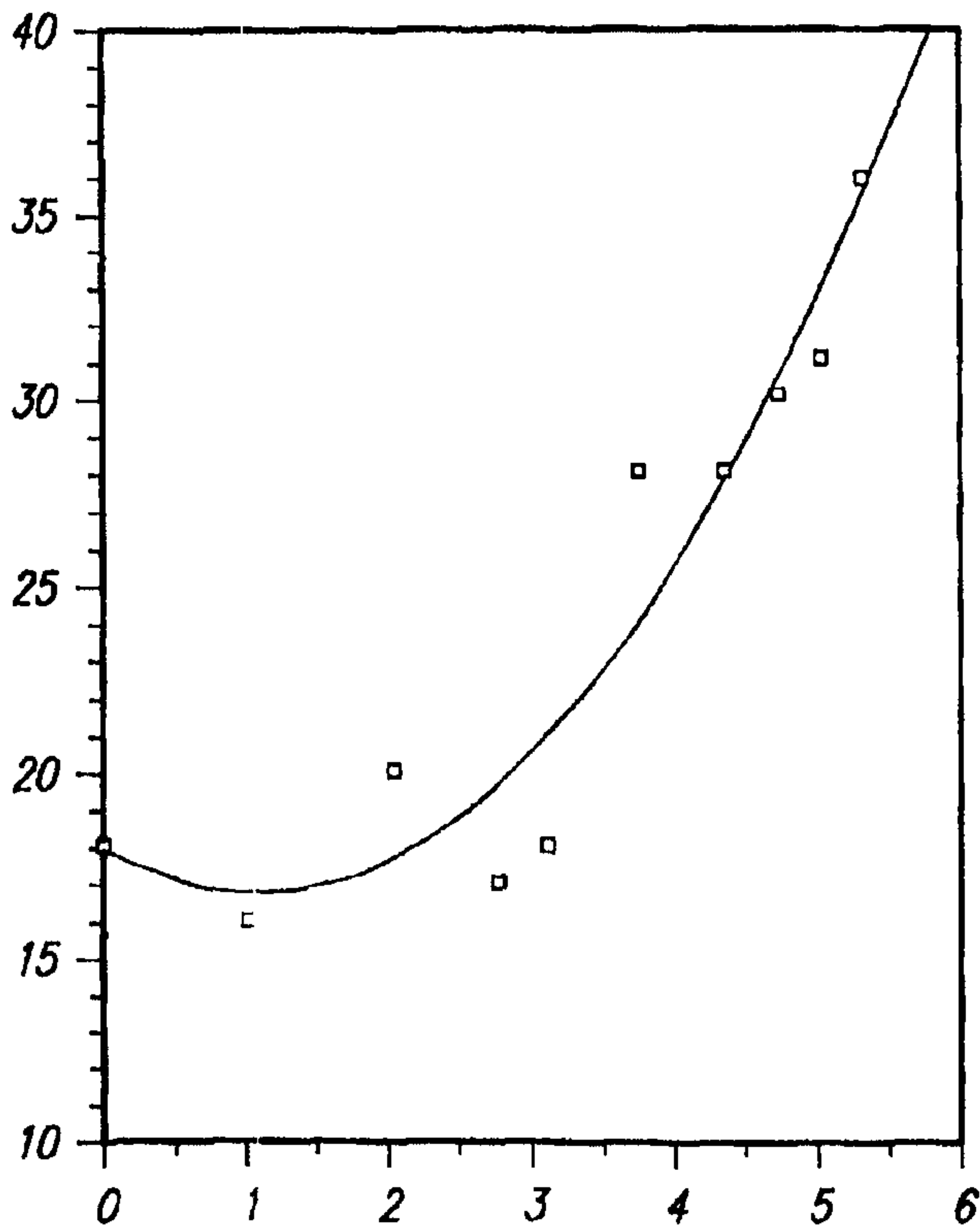


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FIG. 7



PERCENT INHIBITION OF CELL PROLIFERATION



LOG picoMolar CYTOMODULIN