

US008716232B2

(12) United States Patent

Feghali-Bostwick et al.

(54) USE OF ENDOSTATIN PEPTIDES FOR THE TREATMENT OF FIBROSIS

- (71) Applicants: Carol A. Feghali-Bostwick, Pittsburgh, PA (US); Yukie Yamaguchi, Kanagawa (JP)
- (72) Inventors: Carol A. Feghali-Bostwick, Pittsburgh, PA (US); Yukie Yamaguchi, Kanagawa (JP)
- (73) Assignee: University of Pittsburgh— of the Commonwealth System of Higher Education, Pittsburgh, PA (US)
- (*) Notice: Subject to any disclaimer, the term of this patent is extended or adjusted under 35 U.S.C. 154(b) by 0 days.
- (21) Appl. No.: 13/939,058
- (22) Filed: Jul. 10, 2013

(65) **Prior Publication Data**

US 2013/0316959 A1 Nov. 28, 2013

Related U.S. Application Data

- (62) Division of application No. 13/503,339, filed as application No. PCT/US2010/053831 on Oct. 22, 2010, now Pat. No. 8,507,441.
- (60) Provisional application No. 61/261,280, filed on Nov. 13, 2009, provisional application No. 61/254,143, filed on Oct. 22, 2009.
- (51) Int. Cl.

A61K 38/00	(2006.01)
AUIN 30/00	(2000.01)
A61P 17/00	(2006.01)
A61K 48/00	(2006.01)
C12N 15/00	(2006.01)
C12N 5/00	(2006.01)
C12N 5/02	(2006.01)
C12N 5/07	(2010.01)
C12N 5/10	(2006.01)
C07K 5/00	(2006.01)
C07K 7/00	(2006.01)
C07K 16/00	(2006.01)
C07K 17/00	(2006.01)
C07H 21/02	(2006.01)

- (52) U.S. Cl. USPC 514/18.6; 514/44 R; 435/320.1; 435/325; 435/348; 435/375; 530/324; 536/23.5
- (58) Field of Classification Search None

See application file for complete search history.

(56) **References Cited**

U.S. PATENT DOCUMENTS

6,764,995	B2	7/2004	O'Reilly et al.
2006/0193830	A1	8/2006	Hauswirth et al.

(10) Patent No.: US 8,716,232 B2

(45) **Date of Patent:** May 6, 2014

FOREIGN PATENT DOCUMENTS

EP	1 985 302	10/2008
JP	2007084459 A	4/2007
KR	10-2000-0006175	9/2001
WO	WO 02/068457	9/2002
WO	WO 02068457 A2 *	9/2002
WO	WO 2008/014564	2/2008

OTHER PUBLICATIONS

Chufán et al., "Amidation of Bioactive Peptides: The Structure of the Lyase Domain of the Amidating Enzyme," *Structure* 17:965-973 (Jul. 15, 2009).

Hui et al., "Effect of Recombinant Adenovirus Vector Expressing Human Endostatin on Endothelial Cell Proliferation," *Journal of Clinical Rehabilitative Tissue Engineering Research* 12(50): 9986-9989 (Dec. 9, 2008).

International Search Report from parent PCT Application No. PCT/US2010/53831, 8 pages, (mailed on Jan. 7, 2011).

Isobe et al., "Inhibition of Endostatin/Collagen XVIII Deteriorates Left Ventricular Remodeling and Heart Failure in Rat Myocardial Infarction Model," *Circulation Journal* 74: 109-119 (Jan. 2010).

Kim and Seong, "Peptide Amidation: Production of Peptide Hormones in vivo and in vitro," *Biotechnol Bioprocess Engineering* 6:244-251 (Aug. 2001).

Kim et al., "Endostatin Inhibits Endothelial and Tumor Cellular Invasion by Blocking the Activation and Catalytic Activity of Matrix Metalloproteinase 2," *Cancer Research* 60:5410-5413 (Oct. 1, 2000).

Morbidelli et al., "Angiosuppressive and Angiostimulatory Effects Exerted by Synthetic Partial Sequences of Endostatin," *Clinical Cancer Research* 9: 5358-5369 (Nov. 12, 2003).

Richter et al., "Soluble Endostatin is a Novel Inhibitor of Epithelial Repair in Idiopathic Pulmonary Fibrosis," *Thorax* 64:156-161 (2009).

Rodriguez et al., "Lysyl Oxidase as a Potential Therapeutic Target," Drug News Perspect 21(4): 218-224 (2008).

Szauter et al, "Lysyl Oxidase in Development, Aging and Pathologies of the Skin," *Pathologie Biologie* 53: 448-456 (2005).

Written Opinion from parent PCT Application No. PCT/US2010/53831, 8 pages, (mailed on Jan. 7, 2011).

Yamaguchi et al., "Endostatin Inhibits VEGF-induced Endothelial Cell Migration and Tumor Growth Independently of Zinc Binding," *EMBO* 18:4414-4423 (Aug. 16, 1999).

* cited by examiner

Primary Examiner — Cecilia J Tsang Assistant Examiner — Zachary J Miknis

(74) Attorney, Agent, or Firm - Klarquist Sparkman, LLP

(57) ABSTRACT

C-terminal endostatin polypeptides are disclosed herein. Polynucleotides encoding these polypeptide, host cells transformed with the polynucleotides, and methods of using these polypeptides and polynucleotides are disclosed. Uses of these polypeptide, polynucleotides and expression vectors include the treatment of fibrosis in a subject. Thus, methods are provided for treating fibrosis, including fibrosis of the skin and/or the lung.

23 Claims, 17 Drawing Sheets





FIG. 1D

FIG. 1E







FIG. 2A















FIG. 3B





FIG. 4B



FIG. 5A



FIG. 5B

















FIG. 7B





FIG. 9







FIG. 10B * (bun) 8 6 4 U Bleo Bleo Bleo + E-4 E-4L С

Trichrome



Bleo + E4

Bleo + E-4L



Bleo



H&E

Bleo + E4

Bleo + E-4L



FIG. 11

100x











FIG. 14B V E4 V E4 TGFβ -+ +Pro-MMP2 Active-MMP2

FIG. 15



FIG. 16 1 2 3 4









USE OF ENDOSTATIN PEPTIDES FOR THE TREATMENT OF FIBROSIS

CROSS REFERENCE TO RELATED APPLICATION

This is a divisional of U.S. patent application Ser. No. 13/503,339, filed Apr. 20, 2012, which is the U.S. national stage of PCT Application No. PCT/US2010/053831, filed Oct. 22, 2010, which was published in English under PCT ¹⁰ Article 21(2), which claims the benefit of U.S. Provisional Application No. 61/261,280, filed Nov. 13, 2009 and U.S. Provisional Application No. 61/254,143, filed Oct. 22, 2009. The prior applications are incorporated by reference herein in their entirety. ¹⁵

STATEMENT OF GOVERNMENT SUPPORT

This invention was made with government support under grant AR050840 awarded by the National Institutes of ²⁰ Health; the government has certain rights in the invention.

FIELD

This relates to the field of fibrosis, specifically to the use of ²⁵ C-terminal polypeptides of endostatin for the treatment of fibrosis.

BACKGROUND

Endostatin, a 183 amino acid proteolytic cleavage fragment corresponding to the C-terminus of collagen 18, has anti-tumor activity with no toxic side effects (O'Reilly et al. (1997) Cell, 88: 277-285.; Kisker et al. (2001) Cancer Res, 61:7669-7674; Dhanabal et al. (1999) Cancer Res, 59: 189- 35 197; Yoon et al. (1999) Cancer Res, 59: 6251-6256; Folkman and Kalluri, (2003) Cancer Medicine, 6th edition, pp. 161-194. Hamilton: B. C. Decker Inc.). A number of anti-angiogenic activities have been reported for this protein, such as inhibition of endothelial cell proliferation, migration, and 40 tube formation. This activity has been localized to the N-terminal region of endostatin. Endostatin also suppresses vascular endothelial growth factor (VEGF)-induced vascular permeability (Takahashi et al. (2003) Faseb J, 17: 896-898). Endostatin inhibits endothelial cell migration by inhibiting 45 phosphorylation of focal adhesion kinase via binding to $\alpha 5\beta 1$ integrin (Wickstrom et al. (2002) Cancer Res, 62: 5580-5589). It also has been shown that cell surface glypicans are low-affinity endostatin receptors (Karumanchi et al. (2001) Mol Cell, 7: 811-822). Endostatin has been implicated in 50 several signaling pathways, such as downregulation of c-myc (Shichiri and Hirata (2001) Faseb J, 15: 1044-1053), cyclin-D1 (Hanai et al. (2002) J Biol Chem, 277. 16464-16469) and RhoA activity (Wickstrom et al. (2003) J Biol Chem, 278: 37895-37901), blockage of VEGF signaling (Hajitou et al. 55 (2002) Faseb J, 16: 1802-1804; Kim et al. (2002) J Biol Chem, 277: 27872-27879), and inhibition of the wnt-signaling pathway (Hanai et al. (2002) J Cell Biol, 158: 529-539). Furthermore, endostatin has been shown to bind and inactivate metalloproteinases (Kim et al. (2000) Cancer Res, 60: 60 5410-5413; Nyberg et al. (2003) J Biol Chem, 278: 22404-22411; Lee et al. (2002) FEBS Lett, 519: 147-152) and to regulate a spectrum of genes which suppress angiogenesis (Abdollahi et al. (2004) Mol Cell, 13: 649-663).

The crystal structures of both murine and human endosta- 65 tin have been resolved (Hohenester et al. (1998) Embo J, 17: 1656-1664; Ding et al. (1998) Proc Natl Acad Sci USA, 95:

10443-10448) and show a noncovalently held dimer at high concentration required for crystallization (Ding et al. (1998) Proc Natl Acad Sci USA, 95: 10443-10448). The presence of two disulfide bonds results in a highly folded structure. Endostatin binds one atom of zinc per monomer via the three histidines in the N-terminus of the molecule (histidines 1, 3, and 11) and asparatic 76. The heparin binding property of endostatin is mediated by noncontiguous arginines clustered over the three dimensional globular surface of the molecule (Sasaki et al. (1999) Embo J, 18: 6240-6248).

Excessive deposition of extra cellular matrix (ECM) components such as fibronectin (FN) and type I collagen (Col1 α 1) by organ fibroblasts is defined as fibrosis. Organ fibrosis is the final common pathway for many diseases that result in end-stage organ failure. However, effective therapy for organ fibrosis is still unavailable (see, for example, Bjoraker et al., Am. J. Respir. Crit. Care. Med 2000; 157:199-203). Uncontrollable wound-healing responses, including acute and chronic inflammation, angiogenesis, activation of resident cells, and ECM remodeling, are thought to be involved in the pathogenesis of fibrosis (Wynn, J Clin Invest 2007; 117:524-29; Kalluri et al., Curr Opin Nephrol Hypertens 2000; 9:413-8). TGF- β is the prototype fibrotic cytokine that is increased in fibrotic organs and contributes to the development of fibrosis by stimulating the synthesis of ECM molecules, activating fibroblasts to α -smooth muscle actin $(\alpha$ -SMA)-expressing myofibroblasts, and downregulating matrix metalloproteinases (MMPs) (see, for example, Branton et al., Microbes Infect 1999; 1:1349-65). Despite high expectations, a clinical trial of a monoclonal anti-TGF-β antibody in patients with early SSc failed to show any efficacy (Varga et al., Nature Reviews Rheumatology 2009; 5:200-6). Thus, a need remains for other treatments of fibrosis.

SUMMARY

C-terminal endostatin polypeptides are disclosed herein that have anti-fibrotic activity. In some embodiments, these polypeptides include, consist essentially of or consist of (1) at least 40 consecutive amino acids of the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 13 or SEQ ID NO: 4; (2) at least 40 consecutive amino acids of the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 13 or SEQ ID NO: 4, with at most 5 amino acid substitutions, (3) the amino acid sequence set forth as amino acids 133-180 of SEO ID NO: 2, SEQ ID NO: 13 or SEQ ID NO: 4; or (4) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 13 or SEQ ID NO: 4 with at most 5 amino acid substitutions. These polypeptides have anti-fibrotic activity and do not include amino acids 1-92 of SEQ ID NO: 2, SEQ ID NO: 13, or SEQ ID NO: 4, respectively. Polynucleotides encoding these polypeptides, host cells transformed with the polynucleotides, and methods of using these polypeptides and polynucleotides are disclosed. In one example, the polypeptide includes a modification of the carboxy terminal polypeptide to include an amide.

In some embodiments, methods are disclosed for inhibiting fibrosis in vivo or in vitro. In additional embodiments, methods are disclosed for the treatment of fibrosis in a subject. In some specific non-limiting examples, the subject has scleroderma or pulmonary fibrosis.

The foregoing and other features and advantages will become more apparent from the following detailed description of several embodiments, which proceeds with reference to the accompanying figures.

BRIEF DESCRIPTION OF THE FIGURES

FIGS. 1A-1F. ECM production in recombinant endostatinand endostatin-derived peptide-treated fibroblasts in combination with TGF- β stimulation. A: FN and Col1 α 1 expression 5 in human normal lung fibroblasts (NL) treated with vehicle (V), rE alone, or both with prior TGF- β stimulation. Proteins were detected Western blot. GAPDH was used as a loading control for lysates. B: FN and Col1a1 expression of endostatin polypeptide-treated lung fibroblasts following TGF-B 10 stimulation in primary pulmonary fibroblasts from a healthy control, a patient with SSc, and a patient with IPF. C: Graphical summary of FN and Col1a1 expression in lung fibroblasts obtained using fibroblasts from 4 healthy controls (NL), 3 patients with SSc, and 3 patients with IPF. Intensity of bands 15 was normalized to that of GAPDH and expressed as a ratio to Vehicle (V). Paired-t test was used for statistical analysis. * P<0.04, ** P<0.01. D: Representative result of FN and Col1a1 levels in human skin fibroblasts obtained from a patient with morphea and a patient with SSc. E: Representa- 20 tive result of FN and Col1a1 expression in fibrotic fibroblasts obtained from a patients with IPF treated with V, 5 µg/ml of rE, or endostatin polypeptides alone (left). IPF fibroblasts were treated with different concentrations (5, 10, and 20 µg/ml) of E4. DMSO (V) was added in a volume equivalent to 25 that in the lane corresponding to 20 μ g/ml of E4 (right). F: a-SMA levels in normal lung fibroblasts treated with endostatin polypeptides following TGF-β stimulation.

FIGS. **2A-2**C. Ex vivo human skin fibrosis organ culture model. A: Recombinant TGF-β or 1×PBS (vehicle) was 30 injected intradermally into human skin explants at a concentration of 1, 10, 50 ng/ml. Skin was harvested 1 week postinjection. Representative H&E images are shown in the upper row, and images of Masson trichrome-stained section are shown in the lower row. Magnification, 20×. B: Recombinant 35 endostatin (rE) was injected into human skin at a concentration of 1, 5, 10 µg/ml. 1×PBS was used as a vehicle control (V). Representative H&E images are shown. Magnification, 20×. C: Endostatin polypeptides (E-1, E-2, E-3, and E-4; all at 10 µg/ml) were injected intradermally in human skin. DMSO 40 was used as a vehicle control (V). Representative H&E images are shown. Magnification, 20×.

FIGS. **3**A-**3**B. The effect of recombinant endostatin on TGF-β-induced fibrosis and dermal thickness in human skin. A: Representative H&E images of human skin injected with 45 Vehicle, 10_ng/ml TGF-β alone, or rE (1, 5, and 10 µg/ml) in combination with TGF-β (10 ng/ml). Tissues were harvested one week post-injection. Magnification, 20×. B: Graphical presentation of dermal thickness. Data represent four independent experiments in triplicate using human skin explants 50 from four different donors. Mann-Whitney U test was used for statistical analysis. * P<0.04.

FIGS. 4A-4B. The effect of endostatin polypeptides TGF- β -induced fibrosis and dermal thickness in human skin. A: Representative H&E images of human skin injected with 55 Vehicle, 10 ng/ml TGF- β alone, or E-1, E-2, E-3, or E-4 (10 µg/ml) in combination with TGF- β (10 ng/ml). Magnification, 20×. B: Graphical presentation of dermal thickness data shown in A. Data represent two independent experiments using human skin explants from two donors, and each experiment was done in triplicate. Mann-Whitney U test was used for statistical analysis. * P<0.05, ** P<0.02.

FIGS. 5A-5B. Dose response of E-1 and E-4 in TGF- β induced fibrosis. A: Representative H&E images of human skin injected with E-1 (upper row) or E-4 (lower row) at a 65 concentration of 1, 5, 10, and 20 µg/ml in the presence of TGF- β (10 ng/ml). Magnification, 20x. B: Graphical analysis

of dermal thickness data shown in A. DMSO was used as a vehicle control. Experiments were conducted in duplicate, and dermal thickness was measured in 6 fields from each section. Mann-Whitney U test was used for statistical analysis. * P < 0.02, ** P < 0.01.

FIGS. 6A-6B. The effect of endostatin polypeptides in the development of fibrosis in vivo in mouse skin. A: Mice were injected intradermally with vehicle, 10 ng/ml TGF- β alone, or E-1, E-2, E-3, and E-4 (10 µg/ml) in combination with TGF- β (10 ng/ml). Skin was harvested after 1 week post-injection. Sections were stained with H&E. Magnification, $20 \times$. B: Graphical summary of dermal thickness data shown in A. Data represent four independent experiments, each done in duplicate. Mann-Whitney U test was used for statistical analysis. * P<0.04, ** P<0.01.

FIGS. 7A-7B. Capacity of endostatin polypeptide to inhibit tubular formation in MATRIGEL®. A, Representative images of MATRIGEL® cultures of HUVECs treated with vehicle, rE (50 nM), or E4 (50 nM). An equivalent amount of DMSO was used as vehicle. Magnification 40x. B, Image quantification of the cord formation shown in A. Data shown summarize results of three independent experiments. * P<0.05, one-way ANOVA followed by Bonferroni's test.

FIGS. **8**A-**8**B. The effect of endostatin E-4 on bleomycin induced dermal fibrosis in vivo. A: Mice were injected subcutaneously with 1×PBS as vehicle (V) or Bleomycin (Bleo; 20 µg/mouse) daily. E-4 (10 µg/ml) was mixed with bleomycin on the first day, and daily bleomycin administration was continued without subsequent injections of E4 (Bleo+E-4). Skin was harvested after 10 days. Sections were stained with H&E. Magnification, 100×. B: Graphical summary of dermal thickness data shown in A. Data represent three independent experiments. Mann-Whitney U test was used for statistical analysis. * P<0.001, ** P<0.00001. E4 administration caused a significant attenuation of bleomycin induced dermal fibrosis even with a single administration of E4.

FIG. 9. E4 reverses TGF β -induced dermal fibrosis even if administered 3 days following TGF- β . Mouse skin was treated with TGF β day 1 and E-4L or E-1L (this is E-4 and E-1 administered after a 3 day lag between administration of the fibrotic trigger and the administration of the peptide. E-1 or E-4 was administered intraperitoneally (IP) at day 3 and harvested at day 7. E4 caused a significant decrease of TGFbeta induced dermal fibrosis on day 7. Thus E4 prevents (FIGS. 4-6) and reverses (FIG. 9) dermal fibrosis triggered by TGF β .

FIG. 10A-10C. E4 attenuates bleomycin induced lung fibrosis in vivo. A: Sixty µg of bleomycin was administrated intratracheally in combination with DMSO as a vehicle (Bleo) or E-4 (Bleo+E-4; 10 µg/ml). In some mice, E-4 (10 µg/ml) was administered intratracheally (IT) three days following bleomycin treatment (Bleo+E-4L). PBS was used as a vehicle for bleomycin (V). Lungs were harvested 10 days post-treatment. Representative images stained with H&E (left panel) and Masson trichrome (right panel) are representative of 3 independent experiments. Magnification, 100×. E4 administered concomitantly with bleomycin or three days following bleomycin caused a marked reduction in fibrosis and Masson Trichrome staining. B: Quantification of acid soluble collagen obtained from mouse lungs treated as in panel C with V, Bleo, Bleo+E-4, and Bleo+E-4L. The levels of collagen are presented as µg/mg (lung) from three independent experiments. Unpaired-t test was used for statistical analysis. * P<0.05. E4 polypeptide given 3 days after bleomycin significantly reduced collagen levels in mouse lungs C: Lower magnification (2×) of mouse lung shown in FIG. 9 (BLM+E4/E4L IT day 10). For Bleo+E4L, Bleo was administered first, then there was a lag of three days between Bleo and E4 administration).

FIG. 11. E4 attenuates bleomycin induced lung fibrosis in vivo whether administered intraperitoneally (IP) or intratracheally (IT). Bleomycin was administered IT at day 1, and E4 was administered either IP or IT at day 3. Lungs were harvested at day 21. E4 caused a significant attenuation of bleomycin induced lung fibrosis on day 21 whether administered IP or IT. Thus E4 is effective at reducing fibrosis irrespective of the route of administration. Results are shown for vehicle alone (V), bleomycin alone (Bleo), bleomycin and E4 administered IP, and bleomycin and E4 administered IT.

FIG. **12**. E4 reduces fibrosis in vivo by reducing levels of lysyl oxidase (LOX), thus reducing crosslinking of collagen and rendering it less stable and more susceptible to pro-¹⁵ teolytic degradation. Lung sections of mice treated with BLM with or without E4 were used in immunohistochemistry to detect LOX. The sections shown are control IgG, phosphate buffered saline, bleomycin and bleomycin followed by treatment with E4. 20

FIG. **13**. E4 reduces fibrosis in vitro by blocking TGFβinduced LOX production in primary human lung fibroblasts. Normal lung fibroblasts in passage 4 were treated with vehicle, E4, TGFbeta, or TGFbeta followed 30 min. later by E4. Media conditioned by the fibroblasts were analyzed using Western blot analysis after 48 hour. Lane 1: Vehicle (DMSO); Lane 2: E-4; Lane 3: TGF β; Lane 4: TGF followed by E4. Similar results were obtained when LOX mRNA levels were examined by real-time PCR.

FIG. **14**A-**14**B. A. E4 reduces fibrosis in vitro by inducing MMP-2 activity in primary human lung fibroblasts, thus resulting in increased degradation of collagen and other matrix proteins. Digital image of a gelatin zymography gel showing increased MMP-2 activity when primary human lung fibroblasts are treated with E-4 following TGFβ (lane 4). Lane 1: Vehicle (DMSO); Lane 2: E-4; Lane 3: TGF β; Lane 4: TGF β followed by E4. B. Digital image showing that both total and active MMP-2 levels are increased in cells treated with TGFbeta and E-4. This suggests E-4 increases levels of MMP-2 pro-enzyme, but also increases levels of active matrix metalloproteinase (MMP-2, also called metalloproteinase (MMP-2, also called metalloproteinase encomposition of the section of the section

FIG. **15**. E4 reduces fibrosis in vitro by inducing expression of ID1, an inhibitor of TGF β , in primary human lung fibroblasts. Real-time PCR analysis was performed to determine the ID1 mRNA levels under the indicated conditions. 45

FIG. **16**. E-4 reduces fibrosis in vitro by reducing levels of the master transcription factor Egr-1 in primary human lung fibroblasts. Reduction of Egr-1 levels parallels that of collagen, SMA, and fibronectin. Lane 1: vehicle (DMSO); Lane 2: E4; Lane 3: TGF β , Lane 4 TGF β followed by E4 after 60 50 minutes. The samples were harvested after 24 hours.

FIG. 17A-17B. The effect of endostatin peptides on established fibrosis triggered by TGF-b in human skin. A: Vehicle (DMSO), E-1, or E-4 (10 mg/ml) was additionally injected to human skin 2 days post-administration of 10 ng/ml TGF-b (V, 55 E-1L and E-4L, respectively). Representative H&E images of human skin were shown. Magnification, 20×. B: Graphical presentation of dermal thickness data shown in A. Data represent two independent experiments using human skin explants from two donors, and each experiment was done in duplicate. Mann-Whitney U test was used for statistical analysis. * P<0.01.

SEQUENCE LISTING

65

The nucleic and amino acid sequences listed in the accompanying sequence listing are shown using standard letter abbreviations for nucleotide bases, and three letter code for amino acids, as defined in 37 C.F.R. 1.822. Only one strand of each nucleic acid sequence is shown, but the complementary strand is understood as included by any reference to the displayed strand.

The Sequence Listing is submitted as an ASCII text file [8123-84102-05_Sequence_Listing.txt, _Jul. 10, 2013, 12.5 KB], which is incorporated by reference herein.

The nucleic and amino acid sequences listed in the accompanying sequence listing are shown using standard letter abbreviations for nucleotide bases, and three letter code for amino acids, as defined in 37 C.F.R. 1.822. Only one strand of each nucleic acid sequence is shown, but the complementary strand is understood as included by any reference to the displayed strand. The Sequence Listing is submitted as an ASCII text file [Sequence_Listing.txt, Oct. 22, 2010, 12.5 kilobytes], which is incorporated by reference herein. In the accompanying sequence listing:

SEQ ID NO: 1 is an exemplary nucleic acid sequence ²⁰ encoding human endostatin.

- SEQ ID NO: 2 is the amino acid sequence of human endostatin.
- SEQ ID NO: 3 is an exemplary nucleic acid sequence encoding mouse endostatin.
- SEQ ID NO:4 is the amino acid sequence of mouse endostatin.

SEQ ID NO: 5 is an exemplary nucleic acid sequence encoding a human immunoglobulin $(Ig)G_1$ protein.

SEQ ID NO: 6 is the amino acid sequence of a human IgG_1 protein.

SEQ ID NO: 7 is an exemplary nucleic acid sequence encoding a linker.

SEQ ID NO: 8 is an amino acid sequence of a linker.

SEQ ID NO: 9 is a portion of the rat endostatin polypeptide.

SEQ ID NO: 10 is a portion of the cow endostatin polypeptide.

SEQ ID NO: 11 is a portion of the human collagen XV polypeptide.

SEQ ID NO: 12 is an exemplary nucleic acid sequence encoding an endostatin.

SEQ ID NO: 13 is an amino acid sequence of an exemplary amino acid sequence of endostatin that differs from SEQ ID NO: 2 by three amino acid substitutions.

DETAILED DESCRIPTION

C-terminal endostatin polypeptides are disclosed herein. In some embodiments, these polypeptides include, consist essentially of, or consist of (1) at least 40 consecutive amino acids of the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13 or SEQ ID NO: 2; (2) at least 40 consecutive amino acids of the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13 or SEQ ID NO: 2, with at most 5 amino acid substitutions, (3) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13 or SEQ ID NO: 2; or (4) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13 or SEQ ID NO: 2 with at most 5 amino acid substitutions; wherein the polypeptide has anti-fibrotic activity and wherein the polypeptide does not comprise amino acids 1-92 of SEQ ID NO: 13 or SEQ ID NO: 2. In some embodiments, the polypeptide is amidated at the C-terminus. Polynucleotides encoding these polypeptides, host cells transformed with the polynucleotides, and methods of using these polypeptides and polynucleotides are disclosed. These methods include the treatment of fibrosis in a subject. For example, methods are

provided for treating fibrotic conditions of the lung and the skin. In some embodiments, the anti-fibrotic C-terminal endostatin polypeptides disclosed herein can selectively inhibiting fribrosis. In some examples, fibrosis is inhibited without inhibiting angiogenesis. Thus, the C-terminal 5 endostatin polypeptides can be used to more specifically and selectively target unwanted fibrosis, without interfering with angiogenesis, to achieve a desired therapeutic outcome. II. Terms

Unless otherwise noted, technical terms are used according to conventional usage. Definitions of common terms in molecular biology may be found in Benjamin Lewin, Genes V, published by Oxford University Press, 1994 (ISBN 0-19-854287-9); Kendrew et al. (eds.), The Encyclopedia of 15 Molecular Biology, published by Blackwell Science Ltd., 1994 (ISBN 0-632-02182-9); and Robert A. Meyers (ed.), Molecular Biology and Biotechnology: a Comprehensive Desk Reference, published by VCH Publishers, Inc., 1995 (ISBN 1-56081-569-8).

In order to facilitate review of the various embodiments of this disclosure, the following explanations of specific terms are provided:

Animal: Living multi-cellular vertebrate organisms, a category that includes, for example, mammals and birds. The 25 term mammal includes both human and non-human mammals. Similarly, the term "subject" includes both human and veterinary subjects.

Amidation or amide derivative: A post-translational modification to form an amide that can enhance the biological 30 activity of the polypeptide. In amidation, the C-terminal amino acid (polypeptide-COOH) is modified to form and amide (polypeptide-CONH₂). The amide may be formed by post-translational C-terminal amidation. The amino acid to be modified can be followed by a glycine, which provides the 35 amide group. The process of post-translational amidation of a polypeptide derived from a precursor proprotein is well characterized and involves three enzymatic steps (Cuttitta, The Anatomical Record, 236:87-93, 1993). Step one involves endoproteolytic cleavage at a pair of basic amino acids near 40 the carboxy terminus of the protein. Step two involves carboxypeptidase-mediated removal of basic residues. Step three is the amidation reaction, which involves oxidation of a terminal glycine to form the amide of the neighboring carboxy terminal amino acid. Glycine is the only known amino 45 acid to function as an amide donor for its neighboring amino acid. Although the free acid and amidated forms of a polypeptide are difficult to distinguish structurally, the amide can be 100-1000 times more biologically active than the free acid form of the polypeptide (Cuttitta, The Anatomical Record, 50 236:87-93, 1993). C-terminal amidation is essential to the biological activity of many polypeptides

cDNA (complementary DNA): A piece of DNA lacking internal, non-coding segments (introns) and regulatory sequences that determine transcription. cDNA is synthesized 55 in the laboratory by reverse transcription from messenger RNA extracted from cells.

Collagen: Proteins that are found in the form of elongated fibrils in mammals that are mostly found in fibrous tissues such as tendon, ligament and skin, and is also abundant in 60 cornea, cartilage, bone, blood vessels, the gut, and intervertebral disc. The tropocollagen or "collagen molecule" is a subunit of larger collagen aggregates such as fibrils. It is approximately 300 nm long and 1.5 nm in diameter, made up of three polypeptide strands (called alpha chains), each pos- 65 sessing the conformation of a left-handed helix. In type I collagen, each triple-helix associates into a right-handed

super-super-coil that is referred to as the collagen microfibril. Endostain is the first 183 amino acids of collagen.

Conservative variants: "Conservative" amino acid substitutions are those substitutions that do not substantially affect or decrease an activity or of a C-terminal endostatin polypeptide, such as the ability of the polypeptide to inhibit fibrosis. Specific, non-limiting examples of a conservative substitution include the following examples:

Original Residue	Conservative Substitutions
Al	Ser
Arg	Lys
Asn	Gln, His
Asp	Glu
Cys	Ser
Gln	Asn
Glu	Asp
His	Asn; Gln
Ile	Leu, Val
Leu	Ile; Val
Lys	Arg; Gln; Glu
Met	Leu; Ile
Phe	Met; Leu; Tyr
Ser	Thr
Thr	Ser
Trp	Tyr
Tyr	Trp; Phe
Val	Ile; Leu

The term conservative variation also includes the use of a substituted amino acid in place of an unsubstituted parent amino acid, provided that antibodies raised to the substituted polypeptide also immunoreact with the unsubstituted polypeptide. Non-conservative substitutions are those that reduce an activity, such as the ability of a protein to inhibit fibrosis.

Consists Essentially Of/Consists Of: With regard to a polypeptide, a polypeptide that consists essentially of a specified amino acid sequence if it does not include any additional amino acid residues. However, the polypeptide can include additional non-peptide components, such as labels (for example, fluorescent, radioactive, or solid particle labels), sugars or lipids. With regard to a polypeptide, a polypeptide that consists of a specified amino acid sequence does not include any additional amino acid residues, nor does it include additional biological components, such as nucleic acids lipids, sugars, nor does it include labels. A polypeptide that consists or consists essential of a specified amino acid sequence can be glycosylated or have an amide modification. With regard to a polynucleotide, a polynucleotide that consists essentially of a specified nucleic acid sequence if it does not include any additional nucleic acid residues. However, the polypeptide can include additional non-nucleic acid components, such as labels (for example, fluorescent, radioactive, or solid particle labels) or polypeptides. With regard to a polynucleotide, a polynucleotide that consists of a specified nucleic acid sequence does not include any additional nucleic acid residues, nor does it include additional biological components, such as proteins, nor does it include labels.

Degenerate variant: A polynucleotide encoding a C-terminal endostatin polypeptide that includes a sequence that is degenerate as a result of the genetic code. There are 20 natural amino acids, most of which are specified by more than one codon. Therefore, all degenerate nucleotide sequences are included in this disclosure as long as the amino acid sequence of the C-terminal endostatin polypeptide encoded by the nucleotide sequence is unchanged.

Expression Control Sequences: Nucleic acid sequences that regulate the expression of a heterologous nucleic acid sequence to which it is operatively linked. Expression control sequences are operatively linked to a nucleic acid sequence when the expression control sequences control and regulate the transcription and, as appropriate, translation of the nucleic acid sequence. Thus, expression control sequences can include appropriate promoters, enhancers, transcription terminators, a start codon (i.e., ATG) in front of a proteinencoding gene, splicing signal for introns, maintenance of the correct reading frame of that gene to permit proper translation of mRNA, and stop codons. The term "control sequences" is intended to include, at a minimum, components whose presence can influence expression, and can also include additional components whose presence is advantageous, for example, leader sequences and fusion partner sequences. Expression control sequences can include a promoter.

A promoter is a minimal sequence sufficient to direct transcription. Also included are those promoter elements which 20 are sufficient to render promoter-dependent gene expression controllable for cell-type specific, tissue-specific, or inducible by external signals or agents; such elements may be located in the 5' or 3' regions of the gene. Both constitutive and inducible promoters are included (see e.g., Bitter et al., 25 Methods in Enzymology 153:516-544, 1987). For example, when cloning in bacterial systems, inducible promoters such as pL of bacteriophage lambda, plac, ptrp, ptac (ptrp-lac hybrid promoter) and the like can be used. In one embodiment, when cloning in mammalian cell systems, promoters 30 derived from the genome of mammalian cells (such as the metallothionein promoter) or from mammalian viruses (such as the retrovirus long terminal repeat; the adenovirus late promoter; the vaccinia virus 7.5K promoter) can be used. Promoters produced by recombinant DNA or synthetic tech- 35 niques can also be used to provide for transcription of the nucleic acid sequences.

Endostatin: A 183 amino acid proteolytic cleavage fragment corresponding to the C-terminus of collagen XVIII. C-terminal polypeptides of endostatin include consecutive amino acids from the C-terminal region, which is from amino acid 93 to amino acid 183. Exemplary human endostatin polypeptides are set forth in SEQ ID NO: 2 and SEQ ID NO: 13.

Fibrosis: The formation or development of excess fibrous 45 connective tissue in an organ or tissue as a reparative or reactive process, as opposed to a formation of fibrous tissue as a normal constituent of an organ or tissue. Skin and lungs are susceptible to fibrosis. Exemplary fibrotic conditions are scle-roderma idiopathic pulmonary fibrosis, morphea, fibrosis as a 50 result of Graft-Versus-Host Disease (GVHD), keloid and hypertrophic scar, and subepithelial fibrosis, endomyocardial fibrosis, uterine fibrosis, myelofibrosis, retroperitoneal fibrosis, nephrogenic systemic fibrosis, scarring after surgery, asthma, cirrhosis/liver fibrosis, aberrant wound healing, 55 glomerulonephritis, and multifocal fibroses.

Heterologous: Originating from separate genetic sources or species. A polypeptide that is heterologous to endostatin originates from a nucleic acid that does not encode endostatin. In specific, non-limiting examples, a polypeptide comoprising an C-terminal endostatin polypeptide and a heterologous amino acid sequence includes an Ig (such as IgG₁), β -galactosidase, a maltose binding protein, and albumin, hepatitis B surface antigen, or an immunoglobulin amino acid sequence. Generally, an antibody that specifically binds to a 65 protein of interest such as endostatin will not specifically bind to a heterologous protein.

Host cells: Cells in which a vector can be propagated and its DNA expressed. The cell may be prokaryotic or eukaryotic. The cell can be mammalian, such as a human cell. The term also includes any progeny of the subject host cell. It is understood that all progeny may not be identical to the parental cell since there may be mutations that occur during replication. However, such progeny are included when the term "host cell" is used.

Idiopathic Pulmonary Fibrosis: A condition also known as cryptogenic fibrosing alveolitis (CFA) that is a chronic, progressive form of lung disease characterized by fibrosis of the supporting framework (interstitium) of the lungs. By definition, the term is used only when the cause of the pulmonary fibrosis is unknown ("idiopathic"). When lung tissue from patients with IPF is examined under a microscope by a pathologist, it shows a characteristic set of histologic/pathologic features known as usual interstitial pneumonia (UIP). UIP is characterized by progressive scarring of both lung that involves the supporting framework (interstitium) of the lung.

Inhibiting or treating a disease: Inhibiting a disease, such as fibrosis, refers to inhibiting the full development of a disease. In several examples, inhibiting a disease refers to lessening symptoms of a fibrosis, such as the formation of scar tissue or an increase in range of motion or a decrease in pain. "Treatment" refers to a therapeutic intervention that ameliorates a sign or symptom of a disease or pathological condition related to the disease, such as the fibrosis.

Isolated: An "isolated" biological component (such as a nucleic acid or protein or organelle) has been substantially separated or purified away from other biological components in the cell of the organism in which the component naturally occurs, i.e., other chromosomal and extra-chromosomal DNA and RNA, proteins and organelles. Nucleic acids and proteins that have been "isolated" include nucleic acids and proteins purified by standard purification methods. The term also embraces nucleic acids and proteins prepared by recombinant expression in a host cell as well as chemically synthesized nucleic acids.

Keloid or keloidal scar: A type of scar, which depending on its maturity, is composed of mainly either type III (early) or type I (late) collagen. It is a result of an overgrowth of granulation tissue (collagen type 3) at the site of a healed skin injury which is then slowly replaced by collagen type 1. Keloids are firm, rubbery lesions or shiny, fibrous nodules, and can vary from pink to flesh-colored or red to dark brown in color. A keloid scar is benign, non-contagious, and usually accompanied by severe itchiness, sharp pains, and changes in texture. In severe cases, it can affect movement of skin. Keloids are different than hypertrophic scars, which are raised scars that do not grow beyond the boundaries of the original wound.

Label: A detectable compound or composition that is conjugated directly or indirectly to another molecule to facilitate detection of that molecule. Specific, non-limiting examples of labels include fluorescent tags, enzymatic linkages, and radioactive isotopes.

Linker sequence: A linker sequence is an amino acid sequence that covalently links two polypeptide domains. Linker sequences can be included in the between the C-terminal endostatin polypeptides disclosed herein to provide rotational freedom to the linked polypeptide domains and thereby to promote proper domain folding and presentation to the MHC. By way of example, in a recombinant polypeptide comprising two C-terminal endostatin polypeptides, linker sequences can be provided between them, such as a polypeptide comprising C-terminal endostatin polypeptide-linker-Cterminal endostatin polypeptide. Linker sequences, which are generally between 2 and 25 amino acids in length, are well known in the art and include, but are not limited to, the glycine(4)-serine spacer (×3) described by Chaudhary et al., *Nature* 339:394-397, 1989.

Lysyl oxidase (LOX): Lysyl oxidase is an extracellular copper enzyme that catalyzes formation of aldehydes from 5 lysine residues in collagen and elastin precursors. These aldehydes are highly reactive, and undergo spontaneous chemical reactions with other lysyl oxidase-derived aldehyde residues, or with unmodified lysine residues. This results in crosslinking collagen and elastin, which is essential for stabilization of collagen fibrils and for the integrity and elasticity of mature elastin.

Mammal: This term includes both human and non-human mammals. Similarly, the term "subject" includes both human and veterinary subjects.

Matrix metalloproteinase-2: A 72 kDa type IV collagenase also known as gelatinase A. Proteins of the matrix metalloproteinase (MMP) family are involved in the breakdown of extracellular matrix in normal physiological processes, such as embryonic development, reproduction, and tissue remod- 20 eling, as well as in disease processes, such as arthritis and metastasis. Most MMP's are secreted as inactive proproteins which are activated when cleaved by extracellular proteinases. MMP-2 degrades type IV collagen, the major structural component of basement membranes. MMP-2 also degrades 25 additional substrates such as native and denatured collagen I and fibronectin (see the clip.ubc.ca/archive/mmp_timp_ folder/mmp_substrates.shtm website).

Oligonucleotide: A linear polynucleotide sequence of up to about 100 nucleotide bases in length.

Open reading frame (ORF): A series of nucleotide triplets (codons) coding for amino acids without any internal termination codons. These sequences are usually translatable into a polypeptide.

Operably linked: A first nucleic acid sequence is operably 35 linked with a second nucleic acid sequence when the first nucleic acid sequence is placed in a functional relationship with the second nucleic acid sequence. For instance, a promoter is operably linked to a coding sequence if the promoter affects the transcription or expression of the coding sequence, 40 such as a sequence that encodes a C-terminal endostatin polypeptide. Generally, operably linked DNA sequences are contiguous and, where necessary to join two protein-coding regions, in the same reading frame.

Peptide Modifications: C-terminal endostatin polypeptides 45 include synthetic embodiments of polypeptides described herein. In addition, analogs (non-peptide organic molecules), derivatives (chemically functionalized polypeptide molecules obtained starting with the disclosed polypeptide sequences) and variants (homologs) of these proteins can be 50 utilized in the methods described herein. Each polypeptide of this disclosure is comprised of a sequence of amino acids, which may be either L- and/or D-amino acids, naturally occurring and otherwise.

Peptides can be modified by a variety of chemical tech- 55 niques to produce derivatives having essentially the same activity as the unmodified polypeptides, and optionally having other desirable properties. For example, carboxylic acid groups of the protein, whether carboxyl-terminal or side chain, can be provided in the form of a salt of a pharmaceutically-acceptable cation or esterified to form a C_1 - C_{16} ester, or converted to an amide of formula NR₁R₂ wherein R₁ and R₂ are each independently H or C_1 - C_{16} alkyl, or combined to form a heterocyclic ring, such as a 5- or 6-membered ring. Amino groups of the polypeptide, whether amino-terminal or side chain, can be in the form of a pharmaceutically-acceptable acid addition salt, such as the HCl, HBr, acetic, benzoic,

toluene sulfonic, maleic, tartaric and other organic salts, or can be modified to C_1 - C_{16} alkyl or dialkyl amino or further converted to an amide.

Hydroxyl groups of the polypeptide side chains may be converted to C_1 - C_{16} alkoxy or to a C_1 - C_{16} ester using wellrecognized techniques. Phenyl and phenolic rings of the polypeptide side chains may be substituted with one or more halogen atoms, such as fluorine, chlorine, bromine or iodine, or with C_1 - C_{16} alkyl, C_1 - C_{16} alkoxy, carboxylic acids and esters thereof, or amides of such carboxylic acids. Methylene groups of the polypeptide side chains can be extended to homologous C_2 - C_4 alkylenes. Thiols can be protected with any one of a number of well-recognized protecting groups, such as acetamide groups. Those skilled in the art will also recognize methods for introducing cyclic structures into the polypeptides of this invention to select and provide conformational constraints to the structure that result in enhanced stability.

Peptidomimetic and organomimetic embodiments are envisioned, whereby the three-dimensional arrangement of the chemical constituents of such peptido- and organomimetics mimic the three-dimensional arrangement of the polypeptide backbone and component amino acid side chains, resulting in such peptido- and organomimetics of C-terminal endostatin polypeptide having measurable or enhanced ability to treat fibrosis. For computer modeling applications, a pharmacophore is an idealized three-dimensional definition of the structural requirements for biological activity. Peptidoand organomimetics can be designed to fit each pharmacophore with current computer modeling software (using computer assisted drug design or CADD). See Walters, "Computer-Assisted Modeling of Drugs," in Klegerman & Groves, eds., 1993, Pharmaceutical Biotechnology, Interpharm Press: Buffalo Grove, Ill., pp. 165-174 and Principles of Pharmacology, Munson (ed.) 1995, Ch. 102, for descriptions of techniques used in CADD. Also included are mimetics prepared using such techniques.

Pharmaceutically acceptable carriers: The pharmaceutically acceptable carriers of use are conventional. *Remington's Pharmaceutical Sciences*, by E. W. Martin, Mack Publishing Co., Easton, Pa., 15th Edition (1975), describes compositions and formulations suitable for pharmaceutical delivery of the fusion proteins herein disclosed.

In general, the nature of the carrier will depend on the particular mode of administration being employed. For instance, parenteral formulations usually comprise injectable fluids that include pharmaceutically and physiologically acceptable fluids such as water, physiological saline, balanced salt solutions, aqueous dextrose, glycerol or the like as a vehicle. For solid compositions (such as powder, pill, tablet, or capsule forms), conventional non-toxic solid carriers can include, for example, pharmaceutical grades of mannitol, lactose, starch, or magnesium stearate. In addition to biologically neutral carriers, pharmaceutical compositions to be administered can contain minor amounts of non-toxic auxiliary substances, such as wetting or emulsifying agents, preservatives, and pH buffering agents and the like, for example sodium acetate or sorbitan monolaurate.

A "therapeutically effective amount" is a quantity of a composition to achieve a desired effect in a subject being treated. For instance, this can be the amount necessary to induce an immune response, inhibit fibrosis, reduce scar volume or to measurably alter outward symptoms of the fibrotic condition. When administered to a subject, a dosage will generally be used that will achieve target tissue concentrations (for example, in skin cells or lung tissue) that has been shown to achieve an in vitro effect.

55

60

Polynucleotide: The term polynucleotide or nucleic acid sequence refers to a polymeric form of nucleotide at least 10 bases in length. A recombinant polynucleotide includes a polynucleotide that is not immediately contiguous with both of the coding sequences with which it is immediately con-5 tiguous (one on the 5' end and one on the 3' end) in the naturally occurring genome of the organism from which it is derived. The term therefore includes, for example, a recombinant DNA which is incorporated into a vector; into an autonomously replicating plasmid or virus; or into the 10 genomic DNA of a prokaryote or eukaryote, or which exists as a separate molecule (e.g., a cDNA) independent of other sequences. The nucleotides can be ribonucleotides, deoxyribonucleotides, or modified forms of either nucleotide. The term includes single- and double-stranded forms of DNA.

Peptide or Polypeptide: Any chain of amino acids, regardless of length or post-translational modification (e.g., glycosylation or phosphorylation). In one embodiment, the polypeptide is a C-terminal endostatin polypeptide. A polypeptide can be between 5 and 60 amino acids in length. In one embodiment, a polypeptide is from about 10 to about 55 amino acids in length. In yet another embodiment, a polypeptide is from about 20 to about 50 amino acids in length. In yet another embodiment, polypeptide is about 50 amino acids in length. With regard to polypeptides, the word "about" indicates integer amounts. Thus, in one example, a polypeptide ²⁵ "about" 50 amino acids in length is from 49 to 51 amino acids in length.

Post-translational modification: The modification of a newly formed protein; may involve deletion of amino acids, chemical modification of certain amino acids (for example, 30 amidation, acetylation, phosphorylation, glycosylation, formation of pyroglutamate, oxidation/reduction of sulfa group on a methionine, or addition of similar small molecules) to certain amino acids

Probes and primers: A probe comprises an isolated nucleic 35 acid attached to a detectable label or reporter molecule. Primers are short nucleic acids, preferably DNA oligonucleotides, of about 15 nucleotides or more in length. Primers may be annealed to a complementary target DNA strand by nucleic acid hybridization to form a hybrid between the primer and the target DNA strand, and then extended along the target DNA strand by a DNA polymerase enzyme. Primer pairs can be used for amplification of a nucleic acid sequence, for example by polymerase chain reaction (PCR) or other nucleic-acid amplification methods known in the art. One of skill in the art will appreciate that the specificity of a particu- 45 lar probe or primer increases with its length. Thus, for example, a primer comprising 20 consecutive nucleotides will anneal to a target with a higher specificity than a corresponding primer of only 15 nucleotides. Thus, in order to obtain greater specificity, probes and primers can be selected 50that comprise about 20, 25, 30, 35, 40, 50 or more consecutive nucleotides

Purified: The C-terminal endostatin polypeptides disclosed herein can be purified (and/or synthesized) by any of the means known in the art (see, e.g., Guide to Protein Purification, ed. Deutscher, Meth. Enzymol. 185, Academic Press, San Diego, 1990; and Scopes, Protein Purification: Principles and Practice, Springer Verlag, New York, 1982). Substantial purification denotes purification from other proteins or cellular components. A substantially purified protein is at least about 60%, 70%, 80%, 90%, 95%, 98% or 99% pure. Thus, in one specific, non-limiting example, a substantially purified protein is 90% free of other proteins or cellular components.

Thus, the term purified does not require absolute purity; rather, it is intended as a relative term. For example, a purified 65 nucleic acid is one in which the nucleic acid is more enriched than the nucleic acid in its natural environment within a cell.

In additional embodiments, a nucleic acid or cell preparation is purified such that the nucleic acid or cell represents at least about 60% (such as, but not limited to, 70%, 80%, 90%, 95%, 98% or 99%) of the total nucleic acid or cell content of the preparation, respectively.

Recombinant: A recombinant nucleic acid is one that has a sequence that is not naturally occurring or has a sequence that is made by an artificial combination of two otherwise separated segments of sequence. This artificial combination is often accomplished by chemical synthesis or, more commonly, by the artificial manipulation of isolated segments of nucleic acids, e.g., by genetic engineering techniques.

Scleroderma: A chronic autoimmune disease characterized by fibrosis (or hardening), vascular alterations, and autoantibodies. There are two major forms, one is a systemic form that includes limited cutaneous scleroderma mainly affects the hands, arms and face, although pulmonary hypertension is frequent. Diffuse cutaneous scleroderma (or systemic sclerosis) is rapidly progressing and affects a large area of the skin and one or more internal organs, frequently the kidneys, esophagus, heart and lungs. Systemic scleroderma in both of its forms can be fatal. The other form of scleroderma is a localized form that has two subtypes: morphea and linear scleroderma. The disclosed endostatin peptides can be used to treat any form of scleroderma.

Selectively hybridize: Hybridization under moderately or highly stringent conditions that excludes non-related nucleotide sequences.

In nucleic acid hybridization reactions, the conditions used to achieve a particular level of stringency will vary, depending on the nature of the nucleic acids being hybridized. For example, the length, degree of complementarity, nucleotide sequence composition (for example, GC v. AT content), and nucleic acid type (for example, RNA versus DNA) of the hybridizing regions of the nucleic acids can be considered in selecting hybridization conditions. An additional consideration is whether one of the nucleic acids is immobilized, for example, on a filter.

A specific example of progressively higher stringency conditions is as follows: 2×SSC/0.1% SDS at about room temperature (hybridization conditions); 0.2×SSC/0.1% SDS at about room temperature (low stringency conditions); 0.2× SSC/0.1% SDS at about 42° C. (moderate stringency conditions); and 0.1×SSC at about 68° C. (high stringency conditions). One of skill in the art can readily determine variations on these conditions (e.g., Molecular Cloning: A Laboratory Manual, 2nd ed., vol. 1-3, ed. Sambrook et al., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1989). Washing can be carried out using only one of these conditions, e.g., high stringency conditions, or each of the conditions can be used, e.g., for 10-15 minutes each, in the order listed above, repeating any or all of the steps listed. However, as mentioned above, optimal conditions will vary, depending on the particular hybridization reaction involved, and can be determined empirically.

Sequence identity: The similarity between amino acid sequences is expressed in terms of the similarity between the sequences, otherwise referred to as sequence identity. Sequence identity is frequently measured in terms of percentage identity (or similarity or homology); the higher the percentage, the more similar the two sequences are. Homologs or variants of a C-terminal endostatin polypeptide will possess a relatively high degree of sequence identity when aligned using standard methods.

Methods of alignment of sequences for comparison are well known in the art. Various programs and alignment algorithms are described in: Smith and Waterman, Adv. Appl. Math. 2:482, 1981; Needleman and Wunsch, J. Mol. Biol. 48:443, 1970; Higgins and Sharp, Gene 73:237, 1988; Higgins and Sharp, CABIOS 5:151, 1989; Corpet et al., Nucleic Acids Research 16:10881, 1988; and Pearson and Lipman, Proc. Natl. Acad. Sci. USA 85:2444, 1988. Altschul et al., Nature Genet. 6:119, 1994, presents a detailed consideration of sequence alignment methods and homology calculations.

The NCBI Basic Local Alignment Search Tool (BLAST) 5 (Altschul et al., *J. Mol. Biol.* 215:403, 1990) is available from several sources, including the National Center for Biotechnology Information (NCBI, Bethesda, Md.) and on the internet, for use in connection with the sequence analysis programs blastp, blastn, blastx, tblastn and tblastx. A description 10 of how to determine sequence identity using this program is available on the NCBI website on the internet.

Homologs and variants of a C-terminal endostatin polypeptide are typically characterized by possession of at least 75%, for example at least 80%, sequence identity 15 counted over the full length alignment with the amino acid sequence of endostatin using the NCBI Blast 2.0, gapped blastp set to default parameters. For comparisons of amino acid sequences of greater than about 30 amino acids, the Blast 2 sequences function is employed using the default BLO- 20 SUM62 matrix set to default parameters, (gap existence cost of 11, and a per residue gap cost of 1). When aligning short peptides (fewer than around 30 amino acids), the alignment should be performed using the Blast 2 sequences function, employing the PAM30 matrix set to default parameters (open 25 gap 9, extension gap 1 penalties). Proteins with even greater similarity to the reference sequences will show increasing percentage identities when assessed by this method, such as at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, or at least 99% sequence identity. When less than the 30 entire sequence is being compared for sequence identity, homologs and variants will typically possess at least 80% sequence identity over short windows of 10-20 amino acids, and can possess sequence identities of at least 85% or at least 90% or 95% depending on their similarity to the reference 35 sequence. Methods for determining sequence identity over such short windows are available at the NCBI website on the internet. One of skill in the art will appreciate that these sequence identity ranges are provided for guidance only; it is entirely possible that strongly significant homologs could be 40 obtained that fall outside of the ranges provided.

Therapeutically effective amount: A quantity of compound, such as the C-terminal endostatin polypeptide sufficient to achieve a desired effect in a subject being treated. For instance, this can be the amount necessary to treat or amelio-45 rate fibrosis, such as skin or lung fibrosis, in a subject. In some embodiments, it is the amount necessary to treat a subject by a measurable amount over a period of time, or to measurably inhibit progression of disease, in a subject. In other embodiments, a therapeutically effective amount is the amount necson system to prophylactically inhibit a disease.

An effective amount of a C-terminal endostatin polypeptide may be administered in a single dose, or in several doses, for example daily, during a course of treatment. However, the effective amount will be dependent on the compound applied, 55 the subject being treated, the severity and type of the affliction, and the manner of administration of the compound. 16

Transduced: A transduced cell is a cell into which has been introduced a nucleic acid molecule by molecular biology techniques. As used herein, the term transduction encompasses all techniques by which a nucleic acid molecule might be introduced into such a cell, including transfection with viral vectors, transformation with plasmid vectors, and introduction of naked DNA by electroporation, lipofection, and particle gun acceleration.

Vector: A nucleic acid molecule as introduced into a host cell, thereby producing a transformed host cell. A vector may include nucleic acid sequences that permit it to replicate in a host cell, such as an origin of replication. A vector may also include one or more selectable marker gene and other genetic elements known in the art. Vectors include plasmid vectors, including plasmids for expression in gram negative and gram positive bacterial cell. Exemplary vectors include those for expression in *E. coli* and *Salmonella*. Vectors also include viral vectors, such as, but are not limited to, retrovirus, orthopox, avipox, fowlpox, capripox, suipox, adenoviral, herpes virus, alpha virus, baculovirus, Sindbis virus, vaccinia virus and poliovirus vectors. Vectors also include vectors for expression in yeast cells and insect cells.

Unless otherwise explained, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this disclosure belongs. The singular terms "a," "an," and "the" include plural referents unless context clearly indicates otherwise. Similarly, the word "or" is intended to include "and" unless the context clearly indicates otherwise. It is further to be understood that all base sizes or amino acid sizes, and all molecular weight or molecular mass values, given for nucleic acids or peptides are approximate, and are provided for description. Although methods and materials similar or equivalent to those described herein can be used in the practice or testing of this disclosure, suitable methods and materials are described below. The term "comprises" means "includes." All publications, patent applications, patents, and other references mentioned herein are incorporated by reference in their entirety. In case of conflict, the present specification, including explanations of terms, will control. In addition, the materials, methods, and examples are illustrative only and not intended to be limiting.

C-terminal Endostatin Polypeptides

C-terminal endostatin polypeptides and variants thereof are disclosed herein that inhibit fibrosis, such as found in fibrotic conditions, such as but not limited to scleroderma. The polypeptides comprise a C-terminal amino acid sequence of an endostatin protein, but do not include full length endostatin. The endostatin protein can be a mammalian protein, such as from a human, a non-human primate, a canine, a feline, an equine, a bovine, an ovine, a sheep, or a rodent (e.g., mouse or rat). An exemplary nucleotide sequence encoding human endostatin (the amino acid sequence set forth as SEQ ID NO: 2) is:

(SEQ ID NO: 1) ATGCACAGCC ACCGCGACTT CCAGCCGGTG CTCCACCTGG TTGCGCTCAA CAGCCCCCTG TCAGGCGGCA TGCGGGGCAT CCGCGGGGCC GACTTCCAGT GCTTCCAGCA GGCGCGGGCC GTGGGGGCTGG CGGGCACCTT CCGCGCCTTC CTGTCCTCGC GCCTGCAGGA CCTGTACAGC ATCGTGCGCC GTGCCGACCG CGCAGCCGTG CCCATCGTCA ACCTCAAGGA CGAGCTGCTG

-continued TTTCCCAGCT GGGAGGCTCT GTTCTCAGGC TCTGAGGGTC CGCTGAAGCC CGGGGCACGC ATCTTCTCCT TTAACGGCAA GGACGTCTG ACCCACCCA CCTGGCCCCA GAAGAGCGTG TGGCATGGCT CGGACCCCAA CGGGCGCAGG CTGACCGAGA GCTACTGTGA GACGTGGCGG ACGGAGGCTC CCTCGGCCAC GGGCCAGGCC TACTCGCTGC TGGGGGGGCAG GCTCCTGGGG CAGAGTGCCG CGAGCTGCCA TCACGCCTAC ATCGTGCTAT GCATTGAGAA CAGCTTCATG ACTGCCTCCA AGTAG

See also GENBANK® Accession Nos. NM030582.3; NM130444.2; NM130445.2, all of which are incorporated herein by reference.

Another exemplary nucleotide sequence encoding a human endostatin (the amino acid sequence set forth as SEQ ID NO: 13) is:

(SEQ ID NO: 12) CACAGCCACCGC GACTTCCAGC CGGTGCTCCACCTGGTTGCG CTCAACAGCC CCCTGTCAGG

CGGCATGCGGGGCATCCGCGGGGCCGACTTCCAGTGCTTCCAGCAGGCGGGGCCGTGGGGCTGGCGGGGCACCTTCCGGCCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATCGTGCGCCGTGCCGACCGCCGAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTTCCCAGCTTGGGAGCTCTGTTCTCAGGCTCGAGGGCACCCCACCTGGAAGCCCGGGGCACGCATCGTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGGCATGGCTCCGGACCCAACGGGCGCAGGCTGCCTGGCGAGGGCACCCCACCTGGGGCAGGCTCCTGGGGCAGAGAGTGCCGCGAGACCCACGGGCCAGGCCTCCTCGCTGCTGCGGGGGCAGGCTCCTGGGGCAGAGAGTGCCGCGAGAGTGCCATCACGCCTACATCGTGCTCTGCATGAGAACAGCTTCATGACTGCCTCCAAGTAGTCCCAAGTAGCCACGGCCCCACGCGCCCCACGCCCACCTGCTGCACTCTCCAAGTAGTGCATCACCCCTACATCGTGCTCTGCATGAGAACAGCTTCATGACTGCCTCCAAGTAGTCCCAAGTAGCCACGCGCACACCCGCCACGGCCTCATGACTGCCTCCAAGTAGTGCCGCACGAGCCCACGCCACGACCCCACGCCCACGCCTCATGACTGCCTCCAAGTAGTGCCACGCACACGCCCACGCCACGACCTCATGACTGCTCATGACTGCCTCCAAGTAGTGCCACCACGCCCCACGCCACGACCCCACGCCACGCCTCATGACTGCCTCCAAGTAGTGCCACCACGCCCCACGCCACGCCTGCTGCACCACGCTCATGACTGCCTCCAAGTAGTGCCACCACGCCTGCCACGCCACGCCTGCCACGCCACGCCTGCTGCACCACGCCCTCCAAGTGCTGCCACCACGCCTGCCACGCCACCACCCCACCTGCCCGCCACCACCCCCACCCCTGCCACGCCTGCTGCCACGCCCTCCAAGCCCTGCCACCACGCCTGCCACGCCACGCCCCACCCCCCCCC

An exemplary human endostatin protein is:

 $(SEQ ID NO: 2) \\ HSHRDFQPVL HLVALNSPLS GGMRGIRGAD FQCFQQARAV GLAGTFRAFL SSRLQDLYSI \\ VRRADRAAVP IVNLKDELLF PSWEALFSGS EGPLKPGARI FSF$ **M**GKDVL**T** $HPTWPQKSVW \\ HGSDPNGRRL TESYCETWRT EAPSATGQA$ **Y** $SLLGGRLLGQ SAASCHHAYI VLCIENSFMTASK \\ (SEQ ID NO: 2) \\ SRLQDLYSI \\ SRLQDYSI \\ SRLDYSI \\$

This protein is 183 amino acids in length, and is identical to GENBANK® Accession number AAF01310 except that it is lacking the initiator methionine of AAF01310).

Another exemplary endostatin protein is:

(SEQ ID NO: 13) HSHRDFQPVL HLVALNSPLS GGMRGIRGAD FQCFQQARAV GLAGTFRAFL SSRLQDLYSI VRRADRAAVP IVNLKDELLF PSWEALFSGS EGPLKPGARI FSF<u>D</u>GKDVL<u>R</u> HPTWPQKSVW HGSDPNGRRL TESYCETWRT EAPSATGQA**S** SLLGGRLLGQ SAASCHHAYI VLCIENSFMTASK

65

See also GENBANK® Accession No. CAB90482, which is ₆₀ incorporated herein by reference. SEQ ID NO: 2 is identical to SEQ ID NO: 13, with the exception of three amino acid substitutions, indicated by underlining.

An exemplary nucleotide sequence encoding mouse endostatin is:

(SEQ ID NO: 3)

CATACTCATCAGGACTTTCAGCCAGTGCTCCACCTGGTGGCACTGAACACCCCCCTGTCTGGAGGCATGCGTGGTATCCGTGGAGCAGATTTCCAGTGCTTCCAGCAAGCCCGAGCCGTGGGGCTGTCGGGCACCTTCCGGGCTTTCCTGTCCTCTAGGCTGCAGGACGAGGTGCTATCTGTGCGCCGTGCTGACCGGGGGTCTGTGCCCATCGTCAACCTGCAAGCAGAGGTGCTATCTCCCAGCTGGGACTCCCTGTTTTCTGGCTCCCAGGGTCAACTGCAACCCGGGGCCCGCATCTTTTCTTTTGACGGCAGAGATGTCCTGAGACACCCAGCCTGGCCGCAGAAAGGCGAACTCACGGCTCGGACCCCAGTGGGCGGAGGCTGATGGAGAGTACTGTGAGACATGGCGAACAGGAAACTACTGGGGCTGCCACAACAGCTACATCGTCCTGTGCATTGAGAATAGCTCATGACCTCTTTCTCCAAA.AAAAAGCTGCGAAAACAAACAAAC

An exemplary mouse endostatin protein is:

SFSK

(SEQ ID NO: 4) HTHQDFQPVL HLVALNTPLS GGMRGIRGAD FQCFQQARAV GLSGTFRAFL SSRLQDLYSI VRRADRGSVP IVNLKDEVLS PSWDSLFSGS QGQLQPGARI FSFDGRDVLR HPAWPQKSVW HGSDPSGRRL MESYCETWRT ETTGATGQAS SLLSGRLLEQ KAASCHNSYI VLCIENSFMT

This protein is identical to GENBANK® Accession number AAF69009. The nucleotide and amino acid sequences of other species are also publicly available. 30 125 to 183, 130 to 183, 131 to 183, 132 to 183, 134 to 183, 135 to 183, 120 to 180, 125 to 180, 130 to 180, 131 to 180, 132 to 180, 133 to 180, 134 to 180 or 135 to 180 of SEQ ID NO: 2,

In one embodiment, the C-terminal endostatin polypeptide comprises about 10 to about 60 consecutive amino acids of the C-terminal region of an endostatin protein, but does not 35 include a full length endostatin protein, or the N-terminal region of an endostatin protein. The peptide can include from about 10 to about 55 consecutive amino acids or from about 20 to about 54 consecutive amino acids of the C-terminal region of an endostatin protein, such as about 53 consecutive amino acids of the C-terminal region of an endostatin protein (such as SEQ ID NO: 2). For example, the peptide may include about 40, about 45, about, 46, about 47, about 48, about 49, about 50, about 51, about 52 or about 53 consecu- 45 tive amino acids of the C-terminal region on an endostatin protein, such as amino acids 93 to 183 of endostatin, for example SEQ ID NO: 2, SEQ ID NO: 13 or SEQ ID NO: 4. In the context of an amino acid or nucleic acid sequence, "about" means within one residue (one more or one less than the 50 specified number).

The endostatin peptide can include 40, 45, 46, 47, 48, 49 50, 51, 52, 53 consecutive amino acids of the C-terminal region of an endostatin protein. In some examples the peptide consists of 40, 45, 46, 47, 48, 49 50, 51, 52, 53 consecutive stamino acids of the C-terminal region of an endostatin protein, such as but not limited to SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. In some embodiments, the peptide includes or consists of at least 30 amino acids of amino acids 133 to 180 of endostatin, or a variant thereof that has anti-fibrotic 60 activity.

The endostatin peptide may include, consist of or consist essentially of about amino acid 120, 125, 130, 131, 132, 133, 134 or 135 to about amino acids 175, 180, 181, 182 or 183 of an endostatin protein, such as SEQ ID NO: 2, SEQ ID NO: 4 65 or SEQ ID NO: 13. In several examples, the peptide includes, consists of or consists essentially of amino acid 120 to 183,

125 to 183, 130 to 183, 131 to 183, 132 to 183, 134 to 183, 135 to 183, 120 to 180, 125 to 180, 130 to 180, 131 to 180, 132 to 180, 133 to 180, 134 to 180 or 135 to 180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. In specific examples, the peptide includes, consists of, or consists essentially of amino acids 133-180 of SEQ ID NO: 2, amino acids 133-180 of SEQ ID NO: 4 or amino acids 133-180 of SEQ ID NO: 13. In this context, "consists essentially of" means that a peptide does not include additional amino acid residues but can include additional components, such as a label.

Other endostatin peptide variants disclosed herein may comprise, consist of or consist essentially of an amino acid sequence that has at least about 70%, 80%, 90%, 95%, 98% or 99% identity or homology with a C-terminal endostatin polypeptide. C-terminal endostatin polypeptides do not include a full length endostatin protein or the N-terminal region of an endostatin protein (such as amino acids 1-92 of SEQ ID NO: 2).

In some non-limiting examples, C-terminal endostatin polypeptides can include substitutions, such as conservative amino acid substitutions, in a naturally occurring C-terminal endostatin polypeptide (see SEQ ID NO: 2, 4 or 13) in at most about 1, 2, 3, 4, 5 substitutions would be expected to retain anti-fibrotic activity. The C-terminal endostatin polypeptide can include at most 1, at most 2, at most 3 or at most 4 amino acid substitutions, such as conservative amino acid substitutions.

Peptides that are similar to the sequences described above may contain substitutions, deletions or additions. The differences are preferably in regions that are not significantly conserved among different species. Such regions can be identified by aligning the amino acid sequences of endostatin proteins from various animal species. Thus, the endostatin peptide can include, consist essentially of, or consist of at least 40, at least 45, at least 46, at least 47, at least 48, at least 50, at least 51, at least 52 or all of the amino acids set forth as amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. Alternatively, the endostatin peptide can include at most 1, 2, 3, 4 or 5 amino acid substitutions in one of these sequences, provided the peptide has anti-fibrotic activity. The peptide can be 40, 45, 46, 47, 48, 49, 50, 51, 52 or 53 amino acids in length. The peptide does not include the entire 5 sequence of endostatin, or the N-terminal region, such as of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. In additional embodiments, the peptide is at most 40, 45, 46, 47, 48, 49, 50, 51, 52 or 53 amino acids in length, such as peptide that is 40, 45, 46, 47, 48, 49, 50, 51, 52 or 53 amino acids in 10 length.

In some embodiments, the peptide is modified, such as to include a C-terminal amide. Any of the C-terminal endostatin polypeptides disclosed herein can include a C-terminal amide.

The following is an alignment of the human (amino acid 133-180 of endostatin), mouse, rat, cow collagen XVIII amino acid sequences, and human collagen XV:

Human	SYCETWRTEAPS SYCETWRTE	A TGQASSLL <i>G</i> GRLI ATGQASSLL GRL	l g Q S A A SCH HA L Q AASCH H	YIVLCIENSFMT +YIVLCIENSFMT
Mouse	SYCETWRTETTG	atgqassll s grli	EQKAASCHNS	YIVLCIENSFMT
Rat	SYCETWRTEATG	V TGQASSLL S GRLI	EQKAESCHNS	YIVLCIENSFMT
Cow	SYCETWRT DSRA	ATGQASSLL A GRLI	EQKAAGCHNA	FIVLCIENSFMT
HumXV	NYCEAWRTADTA	VTGLASPLSTGKII	DQKAYSCANR	LIVLCIENSFMT

(amino acids 133-141, amino acids 145-153, amino acids 155-158, 162-166, amino acids 169-180 for the human ³⁰ sequence above, see SEQ ID NO: 2, for the mouse sequence above, see SEQ ID NO: 4, for the rat sequence above, see SEQ ID NO: 9, for the cow sequence above, see SEQ ID NO: 10; human collagen XV is SEQ ID NO: 11) 35

In some embodiments, the highlighted amino acids above show those regions of SEQ ID NO: 2, 4 or 13 should be conserved to preserve anti-fibrotic activity of the polypeptide. In further embodiments, the underlined amino acids should be retained in order to preserve anti-fibrotic activity of the ⁴⁰ polypeptide. Thus, in some embodiments, the C-terminal endostatin polypeptide comprises amino acids 133-141, 145-153, 155-158, 162-166 and 169-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. In one example, the A at position ⁴⁵ 145 is conserved.

E4 is amino acids 133-180 of human endostatin (see amino acids 133-180 of SEQ ID NO: 2) and includes a C-terminal amide. In additional embodiments, a region that should be retained in the peptide to retain anti-fibrotic activity is one or both potential phosphorylation sites in the first seven amino acids of E4 that are conserved: SYCE and TWR (amino acids 1-4 and 5-7 of E4, respectively, see also amino acids 133-136 and 137-139 of SEQ ID NO: 2 or SEQ ID NO: 13). In several embodiments, regions that can be retained in the peptide to retain anti-fibrotic activity are one or both potential myristilation sites: GQaySL and GQsaAS (amino acids 15-20 and 27-32 of E4, respectively, see amino acids 147-152 and amino acids 159-164 of SEQ ID NO: 2 or SEQ ID NO: 13). Thus, in $_{60}$ some embodiments, the C-terminal endostatin polypeptide includes zero, or at most 1, at most 2, at most 3, at most 4, or at most 5 substitutions in amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13, wherein the substitutions are not in amino acids 133-141, 145-153, 155- 65 158, 162-166 and 169-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13, and includes a C-terminal amide.

In other embodiments, amino acids can be substituted that differ between the human and mouse sequences without affecting anti-fibrotic activity. In other embodiments, the bolded and italicized amino acids shown above in the human sequence are those amino acids that can be substituted while preserving anti-fibrotic activity. For example, amino acids 142-144, 154, 159-161, and 181-183 of the amino acid sequence can be altered in the C-terminal endostatin polypeptide. These amino acids can be substituted, for example, with those found in another species, as shown above (SEQ ID NOs: 9-10). For example, the C-terminal endostatin polypeptide can include amino acids 133-180 of SEQ ID NO: 2 or SEQ ID NO: 13, wherein amino acids 142-144, 154, and amino acids 159-161 are substituted. This polypeptide can include a C-terminal amide.

Other amino acids that can be substituted, inserted or deleted at these or other locations can be identified by

mutagenesis studies coupled with biological assays. The above alignment is provided only as a guideline.

Also encompassed herein are C-terminal endostatin polypeptides that are fused to a heterologous peptide, such as a peptide that can be used for detecting; purifying; stabilizing; or solubilizing the endostatin polypeptide. These polypeptides do not include a full length endostatin protein or an N-terminal region of an endostatin protein. In one example, a C-terminal polypeptide can be linked to an immunoglobulin (Ig) constant heavy or light chain domain or portion thereof at its N-terminus. For example, a polypeptide, such as but not limited to amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13(e.g., E4) may be linked to a CH1, CH2 and/or CH3 domain of a heavy chain. If the constant region is from a light chain, it can be from a kappa or lambda light chain. If the constant region is from a heavy chain, it can be from an antibody of any one of the following classes of antibodies: IgG, IgA, IgE, IgD, and IgM. IgG can be an IgG₁, IgG₂, IgG₃ or IgG₄. The constant domain may be an Fc fragment. The constant domain can be from a mammalian antibody, such as a human antibody. Soluble receptor-IgG fusion proteins are common immunological reagents and methods for their construction are known in the art (see, for example, U.S. Pat. Nos. 5,225,538, 5,726,044; 5,707,632; 750,375, 5,925,351, 6,406,697 and Bergers et al. Science 1999 284: 808-12). In one example, the immunoglobulin is the constant part of the heavy chain of human IgG, particularly IgG₁, where dimerization between two heavy chains takes place at the hinge region. It is recognized that inclusion of the CH2 and CH3 domains of the Fc region as part of the fusion polypeptide increases the in vivo circulation half-life of the polypeptide comprising the Fc region, and that of the oligomer or dimer comprising the polypeptide.

An Fc portion of human IgG_1 which includes the hinge region, and domains CH2 and CH3 has the nucleotide sequence:

(SEQ ID NO: 5) GAG CCC AAA TCT TGT GAC AAA ACT CAC ACA TGC CCA CCG TGC CCA GCA CCT GAA CTC CTG GGG GGA CCG TCA GTC TTC CTC TTC CCC CCA AAA CCC AAG GAC ACC CTC ATG ATC TCC CGG ACC CCT GAG GTC ACA TGC GTG GTG GTG GAC GTG AGC CAC GAA GAC CCT GAG GTC AAG TTC AAC TGG TAC GTG GAC GGC GTG GAG GTG CAT AAT GCC AAG ACA AAG CCG CGG GAG GAG CAG TAC AAC AGC ACG TAC CGT GTG GTC AGC GTC CTC ACC GTC CTG CAC CAG GAC TGG CTG AAT GGC AAG GAG TAC AAG TGC AAG GTC TCC AAC AAA GCC CTC CCA GCC CCC ATC GAG AAA ACC ATC TCC AAA GCC AAA GGG CAG CCC CGA GAA CCA CAG GTG TAC ACC CTG CCC CCA TCC CGG GAT GAG CTG ACC AAG AAC CAG GTC AGC CTG ACC TGC CTG GTC AAA GGC TTC TAT CCC AGC GAC ATC GCC GTG GAG TGG GAG AGC AAT GGG CAG CCG GAG AAC AAC TAC AAG ACC ACG CCT CCC GTG CTG GAC TCC GAC GGC TCC TTC TTC CTC TAC AGC AAG CTC ACC GTG GAC AAG AGC AGG TGG CAG CAG GGG AAC GTC TTC TCA TGC TCC GTG ATG CAT GAG GCT CTG CAC AAC CAC TAC ACG CAG AAG AGC CTC TCC CTG TCT CCG GGT AAA TGA.

which encodes a polypeptide having the amino acid sequence:

amino acid sequence: Ser Arg Gly Gly Leu Val Pro Arg Gly Ser Gly Ser Pro Gly Leu Gln (SEQ ID NO: 8). A C-terminal

(SEQ ID NO: 6) Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gin Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gin Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gin Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arq Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gin Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gin Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tvr Thr Gin Lvs Ser Leu Ser Leu Ser Pro Glv Lys.

60

Constant Ig domains can also contain one or more mutations ⁵⁰ that reduce or eliminate one or more effector function, e.g., binding to Fc receptors and complement activation (see, for example, Morrison, Annu. Rev. Immunol., 10, pp. 239-65 (1992); Duncan and Winter (1988) Nature 332: 738-740; and Xu et al. (1994) J Biol. Chem. 269: 3469-3474). For example, mutations of amino acids corresponding to Leu 235 and Pro 331 of human IgG₁ to Glu and Ser respectively, are provided. Such constructs are further described in U.S. Pat. No. 6,656, 728.

The C-terminal endostatin polypeptide can also be linked to a linker sequence with a thrombin cleavage site, such as between the C-terminal endostatin polypeptide and a heterologous polypeptide. An exemplary nucleotide sequence encoding such a site has the following nucleotide sequence: 5' 65 TCT AGA GGT GGT CTA GTG CCG CGC GGC AGC GGT TCC CCC GGG TTG CAG 3' (SEQ ID NO: 7), which encodes a polypeptide having the

endostatin polypeptide can also be fused to a signal sequence. For example, when prepared recombinantly, a nucleic acid encoding the peptide can be linked at its 5' end to a signal sequence, such that the peptide is secreted from the cell.

Peptides can be used as a substantially pure preparation, such as wherein at least about 90% of the peptides in the preparation are the desired peptide. Compositions comprising at least about 50%, 60%, 70%, or 80% of the desired peptide may also be used. Peptides can be denatured or nondenatured and may be aggregated or non-aggregated as a result thereof.

Other C-terminal endostatin polypeptides that are encompassed herein are those that include modified amino acids. Exemplary peptides are derivative peptides that may be one modified by glycosylation, pegylation, phosphorylation or any similar process that retains at least one biological function of the peptide from which it was derived. Peptides may

24

also comprise one or more non-naturally occurring amino acids. For example, nonclassical amino acids or chemical amino acid analogs can be introduced as a substitution or addition into peptides. Non-classical amino acids include, but are not limited to, the D-isomers of the common amino acids, 5 2,4-diaminobutyric acid, alpha-amino isobutyric acid, 4-aminobutyric acid, Abu, 2-amino butyric acid, gamma-Abu, epsilon-Ahx, 6-amino hexanoic acid, Aib, 2-amino isobutyric acid, 3-amino propionic acid, ornithine, norleucine, norvaline, hydroxyproline, sarcosine, citrulline, homocitrulline, 10 cysteic acid, t-butylglycine, t-butylalanine, phenylglycine, cyclohexylalanine, beta-alanine, fluoro-amino acids, designer amino acids such as beta-methyl amino acids, Calpha-methyl amino acids, Nalpha-methyl amino acids, and amino acid analogs in general. Furthermore, the amino acid 15 can be D (dextrorotary) or L (levorotary). In other specific embodiments, branched versions of the peptides listed herein are provided, such as by substituting one or more amino acids within the sequence with an amino acid or amino acid analog with a free side chain capable of forming a peptide bond with 20 one or more amino acids (and thus capable of forming a "branch"). Cyclical peptides are also contemplated.

Also included are peptide derivatives which are differentially modified during or after synthesis, such as by benzylation, glycosylation, acetylation, phosphorylation, amidation, 25 pegylation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to an antibody molecule or other cellular ligand, etc. In specific embodiments, the peptides are acetylated at the N-terminus and/or amidated at the C-terminus. 30

In one example, the peptide includes a carboxy terminal amide. One specific non-limiting example of this type of C-terminal endostatin polypeptide is E4 (see, for example, amino acids 133-180 of SEQ ID NO: 13), which is described in detail in the examples section below. This peptide, or any of 35 the C-terminal endostatin polypeptides disclosed herein can be amidated at the C-terminus.

Also provided are derivatives of C-terminal endostatin polypeptides, such as chemically modified peptides and peptidomimetics. Peptidomimetics are compounds based on, or 40 derived from, peptides and proteins. Peptidomimetics can be obtained by structural modification of known peptide sequences using unnatural amino acids, conformational restraints, isosteric replacement, and the like. The subject peptidomimetics constitute the continum of structural space 45 between peptides and non-peptide synthetic structures; peptidomimetics may be useful, therefore, in delineating pharmacophores and in helping to translate peptides into nonpeptide compounds with the activity of the parent peptides.

Mimetopes of the C-terminal endostatin polypeptides are 50 included in the present disclosure. Such peptidomimetics can have such attributes as being non-hydrolyzable (e.g., increased stability against proteases or other physiological conditions which degrade the corresponding peptide), increased specificity and/or potency for stimulating cell dif- 55 ferentiation. For illustrative purposes, peptide analogs can be generated using, for example, benzodiazepines (e.g., see Freidinger et al. in Peptides: Chemistry and Biology, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988), substituted gama lactam rings (Garvey et al. in Peptides: 60 Chemistry and Biology, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p123), C-7 mimics (Huffman et al. in Peptides: Chemistry and Biology, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p. 105), keto-methylene pseudopeptides (Ewenson et al. (1986) J Med 65 Chem 29:295; and Ewenson et al. in Peptides: Structure and Function (Proceedings of the 9th American Peptide Sympo-

sium) Pierce Chemical Co. Rockland, Ill., 1985), β -turn dipeptide cores (Nagai et al. (1985) Tetrahedron Lett 26:647; and Sato et al. (1986) J Chem Soc Perkin Trans 1: 1231), β -aminoalcohols (Gordon et al. (1985) Biochem Biophys Res Commun 126:419; and Dann et al. (1986) Biochem Biophys Res Commun 134:71), diaminoketones (Natarajan et al. (1984) Biochem Biophys Res Commun 124:141), and methyleneamino-modified (Roark et al. in Peptides: Chemistry and Biology, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988, p134). Also, see generally, Session III: Analytic and synthetic methods, in Peptides: Chemistry and Biology, G. R. Marshall ed., ESCOM Publisher: Leiden, Netherlands, 1988).

In addition to a variety of side-chain replacements which can be carried out to generate peptidomimetics, the present disclosure specifically contemplates the use of conformationally restrained mimics of peptide secondary structure. Numerous surrogates have been developed for the amide bond of peptides. Frequently exploited surrogates for the amide bond include the following groups (i) trans-olefins, (ii) fluoroalkene, (iii) methyleneamino, (iv) phosphonamides, and (v) sulfonamides. Additionally, peptidomimietics based on more substantial modifications of the backbone of a peptide can be used. Peptidomimetics which fall in this category include (i) retro-inverso analogs, and (ii) N-alkyl glycine analogs (so-called peptoids). Furthermore, the methods of combinatorial chemistry can be used to produce peptidomimetics. For example, one embodiment of a so-called "peptide morphing" strategy focuses on the random generation of a library of peptide analogs that comprise a wide range of peptide bond substitutes. In an exemplary embodiment, the peptidomimetic can be derived as a retro-inverso analog of the peptide. Such retro-inverso analogs can be made according to the methods known in the art, such as that described by the Sisto et al. U.S. Pat. No. 4,522,752. A retro-inverso analog can be generated as described, for example in PCT Publication No. WO 00/01720. A mixed peptide, such as one including some normal peptide linkages, can be generated. As a general guide, sites which are most susceptible to proteolysis are typically altered, with less susceptible amide linkages being optional for mimetic switching. The final product, or intermediates thereof, can be purified by HPLC.

In some embodiments, peptides can include at least one amino acid or every amino acid that is a D stereoisomer. Other peptides can include at least one amino acid that is reversed. The amino acid that is reversed may be a D stereoisomer. Every amino acid of a peptide may be reversed and/or every amino acid can be a D stereoisomer. In another illustrative embodiment, a peptidomimetic can be derived as a retroenantio analog of a peptide. Retro-enantio analogs such as this can be synthesized with commercially available D-amino acids (or analogs thereof) and standard solid- or solutionphase peptide-synthesis techniques, as described, for example in PCT Publication No. WO 00/01720. The final product can be purified by HPLC to yield the pure retroenantio analog. In still another illustrative embodiment, transolefin derivatives can be made for the subject peptide. Transolefin analogs can be synthesized according to the method of Y. K. Shue et al. (1987) Tetrahedron Letters 28:3225 and as described in PCT Publication WO 00/01720. It is further possible to couple pseudodipeptides synthesized by the above method to other pseudodipeptides, to make peptide analogs with several olefinic functionalities in place of amide functionalities. Still another class of peptidomimetic derivatives include the phosphonate derivatives. The synthesis of such phosphonate derivatives can be adapted from known synthesis schemes (see, for example, Loots et al. in Peptides: Chemistry and Biology, (Escom Science Publishers, Leiden, 1988, p. 118)); Petrillo et al. in Peptides: Structure and Function (Proceedings of the 9th American Peptide Symposium, Pierce Chemical Co. Rockland, Ill., 1985).

Many other peptidomimetic structures are known in the art 5 and can be readily adapted for use in the subject peptidomimetics. For example, a peptidomimetic may incorporate the 1-azabicyclo[4.3.0]nonane surrogate (see Kim et al. (1997) J. Org. Chem. 62:2847), or an N-acyl piperazic acid (see Xi et al. (1998) J. Am. Chem. Soc. 120:80), or a 2-substituted 10 piperazine moiety as a constrained amino acid analogue (see Williams et al. (1996) J. Med. Chem. 39:1345-1348). In still other embodiments, certain amino acid residues can be replaced with aryl and bi-aryl moieties, such as monocyclic or bicyclic aromatic or heteroaromatic nucleus, or a biaromatic, 15 aromatic-heteroaromatic, or biheteroaromatic nucleus. The subject peptidomimetics can be optimized such as by combinatorial synthesis techniques combined with high throughput screening. Moreover, other examples of mimetopes include, but are not limited to, protein-based compounds, carbohy- 20 drate-based compounds, lipid-based compounds, nucleic acid-based compounds, natural organic compounds, synthetically derived organic compounds, anti-idiotypic antibodies and/or catalytic antibodies, or fragments thereof. A mimetope can be obtained by, for example, screening libraries of natural 25 and synthetic compounds for compounds capable of inhibiting fibrosis. A mimetope can also be obtained, for example, from libraries of natural and synthetic compounds, in particular, chemical or combinatorial libraries (i.e., libraries of compounds that differ in sequence or size but that have the same 30 building blocks). A mimetope can also be obtained by, for example, rational drug design. In a rational drug design procedure, the three-dimensional structure of a compound of the present invention can be analyzed by, for example, nuclear magnetic resonance (NMR) or x-ray crystallography. The 35 three-dimensional structure can then be used to predict structures of potential mimetopes by, for example, computer modelling. The predicted mimetope structures can then be produced by, for example, chemical synthesis, recombinant DNA technology, or by isolating a mimetope from a natural 40 source (for example, plants, animals, bacteria and fungi).

All of the C-terminal endostatin polypeptides of use in the disclosed methods have anti-fibrotic activity. For example, they can reduce or inhibit fibrosis by a factor of at least about 50%, 60%, 70% 80%, 90%, or 2 fold, 5 fold, 10 fold, 30 fold 45 or 100 fold, as compared to a control, such as in an assay described herein.

The C-terminal endostatin polypeptides (including amidated forms of the peptides) can be readily synthesized by automated solid phase procedures well known in the art. 50 Techniques and procedures for solid phase synthesis are described in Solid Phase Peptide Synthesis: A Practical Approach, by E. Atherton and R. C. Sheppard, published by IRL, Oxford University Press, 1989. Alternatively, these peptides may be prepared by way of segment condensation, as 55 described, for example, in Liu et al., Tetrahedron Lett. 37:933-936, 1996; Baca et al., J. Am. Chem. Soc. 117:1881-1887, 1995; Tam et al., Int. J. Peptide Protein Res. 45:209-216, 1995; Schnolzer and Kent, Science 256:221-225, 1992; Liu and Tam, J. Am. Chem. Soc. 116:4149-4153, 1994; Liu 60 and Tam, Proc. Natl. Acad. Sci. USA 91:6584-6588, 1994; and Yamashiro and Li, Int. J. Peptide Protein Res. 31:322-334, 1988). Other methods useful for synthesizing peptides of the present disclosure are described in Nakagawa et al., J. Am. Chem. Soc. 107:7087-7092, 1985. Peptides of the disclosure 65 can also be readily purchased from commercial suppliers of synthetic peptides. Such suppliers include, for example,

Advanced ChemTech (Louisville, Ky.), Applied Biosystems (Foster City, Calif.), Anaspec (San Jose, Calif.), and Cell Essentials (Boston, Mass.).

Polynucleotides Encoding the C-terminal Endostatin Polypeptides and Host Cells

Polynucleotides encoding the C-terminal endostatin polypeptides disclosed herein are also provided. These polynucleotides include DNA, cDNA and RNA sequences which encode the peptide of interest. Silent mutations in the coding sequence result from the degeneracy (i.e., redundancy) of the genetic code, whereby more than one codon can encode the same amino acid residue. Thus, for example, leucine can be encoded by CTT, CTC, CTA, CTG, TTA, or TTG; serine can be encoded by TCT, TCC, TCA, TCG, AGT, or AGC; asparagine can be encoded by AAT or AAC; aspartic acid can be encoded by GAT or GAC; cysteine can be encoded by TGT or TGC; alanine can be encoded by GCT, GCC, GCA, or GCG; glutamine can be encoded by CAA or CAG; tyrosine can be encoded by TAT or TAC; and isoleucine can be encoded by ATT, ATC, or ATA. Tables showing the standard genetic code can be found in various sources (see, for example, Stryer, 1988, Biochemistry, 3.sup.rd Edition, W.H. 5 Freeman and Co., NY).

A nucleic acid encoding a C-terminal endostatin polypeptide can be cloned or amplified by in vitro methods, such as the polymerase chain reaction (PCR), the ligase chain reaction (LCR), the transcription-based amplification system (TAS), the self-sustained sequence replication system (3SR) and the $Q\beta$ replicase amplification system (QB). For example, a polynucleotide encoding the protein can be isolated by polymerase chain reaction of cDNA using primers based on the DNA sequence of the molecule. A wide variety of cloning and in vitro amplification methodologies are well known to persons skilled in the art. PCR methods are described in, for example, U.S. Pat. No. 4,683,195; Mullis et al., Cold Spring Harbor Symp. Quant. Biol. 51:263, 1987; and Erlich, ed., PCR Technology, (Stockton Press, NY, 1989). Polynucleotides also can be isolated by screening genomic or cDNA libraries with probes selected from the sequences of the desired polynucleotide under stringent hybridization conditions.

The polynucleotides encoding a C-terminal endostatin polypeptide include a recombinant DNA which is incorporated into a vector in an autonomously replicating plasmid or virus or into the genomic DNA of a prokaryote or eukaryote, or which exists as a separate molecule (such as a cDNA) independent of other sequences. The nucleotides can be ribonucleotides, deoxyribonucleotides, or modified forms of either nucleotide. The term includes single and double forms of DNA.

In one embodiment, vectors are used for expression in yeast such as S. cerevisiae or Kluvveromyces lactis. Several promoters are known to be of use in yeast expression systems such as the constitutive promoters plasma membrane H+-AT-Pase (PMA1), glyceraldehyde-3-phosphate dehydrogenase (GPD), phosphoglycerate kinase-1 (PGK1), alcohol dehydrogenase-1 (ADH1), and pleiotropic drug-resistant pump (PDR5). In addition, many inducible promoters are of use, such as GAL1-10 (induced by galactose), PHO5 (induced by low extracellular inorganic phosphate), and tandem heat shock HSE elements (induced by temperature elevation to 37° C.). Promoters that direct variable expression in response to a titratable inducer include the methionine-responsive MET3 and MET25 promoters and copper-dependent CUP1 promoters. Any of these promoters may be cloned into multicopy (2µ) or single copy (CEN) plasmids to give an additional level of control in expression level. The plasmids can include nutritional markers (such as URA3, ADE3, HIS1, and others) for selection in yeast and antibiotic resistance (AMP) for propagation in bacteria. Plasmids for expression on K. lactis are known, such as pKLAC1. Thus, in one example, after amplification in bacteria, plasmids can be introduced into the corresponding yeast auxotrophs by methods similar to bacterial transformation. The polynucleotides can also be designed to express in insect cells.

The C-terminal endostatin polypeptides can be expressed in a variety of yeast strains. For example, seven pleiotropic drug-resistant transporters, YOR1, SNQ2, PDR5, YCF1, PDR10, PDR11, and PDR15, together with their activating transcription factors, PDR1 and PDR3, have been simultaneously deleted in yeast host cells, rendering the resultant 15 strain sensitive to drugs. Yeast strains with altered lipid composition of the plasma membrane, such as the erg6 mutant defective in ergosterol biosynthesis, can also be utilized. Proteins that are highly sensitive to proteolysis can be expressed in a yeast lacking the master vacuolar endopeptidase Pep4, 20 which controls the activation of other vacuolar hydrolases. Heterologous expression in strains carrying temperature-sensitive (ts) alleles of genes can be employed if the corresponding null mutant is inviable.

Viral vectors can also be prepared encoding the C-terminal 25 endostatin polypeptides disclosed herein. A number of viral vectors have been constructed, including polyoma, SV40 (Madzak et al., 1992, J. Gen. Virol., 73:15331536), adenovirus (Berkner, 1992, Cur. Top. Microbiol. Immunol., 158:39-6; Berliner et al., 1988, Bio Techniques, 6:616-629; Gorziglia 30 et al., 1992, J. Virol., 66:4407-4412; Quantin et al., 1992, Proc. Nad. Acad. Sci. USA, 89:2581-2584; Rosenfeld et al., 1992, Cell, 68:143-155; Wilkinson et al., 1992, Nucl. Acids Res., 20:2233-2239; Stratford-Perricaudet et al., 1990, Hum. Gene Ther., 1:241-256), vaccinia virus (Mackett et al., 1992, 35 Biotechnology, 24:495-499), adeno-associated virus (Muzyczka, 1992, Curr. Top. Microbiol. Immunol., 158:91-123; On et al., 1990, Gene, 89:279-282), herpes viruses including HSV and EBV (Margolskee, 1992, Curr. Top. Microbiol. Immunol., 158:67-90; Johnson et al., 1992, J. Virol., 40 66:29522965; Fink et al., 1992, Hum. Gene Ther. 3:11-19; Breakfield et al., 1987, Mol. Neurobiol., 1:337-371; Fresse et al., 1990, Biochem. Pharmacol., 40:2189-2199), Sindbis viruses (H. Herweijer et al., 1995, Human Gene Therapy 6:1161-1167; U.S. Pat. Nos. 5,091,309 and 5,2217,879), 45 alphaviruses (S. Schlesinger, 1993, Trends Biotechnol. 11:18-22; I. Frolov et al., 1996, Proc. Natl. Acad. Sci. USA 93:11371-11377) and retroviruses of avian (Brandyopadhyay et al., 1984, Mol. Cell Biol., 4:749-754; Petropouplos et al., 1992, J. Virol., 66:3391-3397), murine (Miller, 1992, Curr. 50 Top. Microbiol. Immunol., 158:1-24; Miller et al., 1985, Mol. Cell Biol., 5:431-437; Sorge et al., 1984, Mol. Cell Biol., 4:1730-1737; Mann et al., 1985, J. Virol., 54:401-407), and human origin (Page et al., 1990, J. Virol., 64:5370-5276; Buchschalcher et al., 1992, J. Virol., 66:2731-2739). Bacu- 55 lovirus (Autographa californica multinuclear polyhedrosis virus; AcMNPV) vectors are also known in the art, and may be obtained from commercial sources (such as PharMingen, San Diego, Calif.; Protein Sciences Corp., Meriden, Conn.; Stratagene, La Jolla, Calif.).

Thus, in one embodiment, the polynucleotide encoding a C-terminal endostatin polypeptide is included in a viral vector. Suitable vectors include retrovirus vectors, orthopox vectors, avipox vectors, fowlpox vectors, capripox vectors, suipox vectors, adenoviral vectors, herpes virus vectors, alpha 65 virus vectors, baculovirus vectors, Sindbis virus vectors, vaccinia virus vectors and poliovirus vectors. Specific exemplary

60

vectors are poxvirus vectors such as vaccinia virus, fowlpox virus and a highly attenuated vaccinia virus (MVA), adenovirus, baculovirus and the like.

Pox viruses of use include orthopox, suipox, avipox, and capripox virus. Orthopox include vaccinia, ectromelia, and raccoon pox. One example of an orthopox of use is vaccinia. Avipox includes fowlpox, canary pox and pigeon pox. Capripox include goatpox and sheeppox. In one example, the suipox is swinepox. Examples of pox viral vectors for expression as described for example, in U.S. Pat. No. 6,165,460, which is incorporated herein by reference. Other viral vectors that can be used include other DNA viruses such as herpes virus and adenoviruses, and RNA viruses such as retroviruses and polio.

Suitable vectors are disclosed, for example, in U.S. Pat. No. 6,998,252, which is incorporated herein by reference. In one example, a recombinant poxvirus, such as a recombinant vaccinia virus is synthetically modified by insertion of a chimeric gene containing vaccinia regulatory sequences or DNA sequences functionally equivalent thereto flanking DNA sequences which in nature are not contiguous with the flanking vaccinia regulatory DNA sequences that encode a C-terminal endostatin polypeptide. The recombinant virus containing such a chimeric gene is effective at expressing the C-terminal endostatin polypeptide. In one example, the vaccine viral vector comprises (A) a segment comprised of (i) a first DNA sequence encoding a C-terminal endostatin polypeptide and (ii) a poxvirus promoter, wherein the poxvirus promoter is adjacent to and exerts transcriptional control over the DNA sequence encoding a C-terminal endostatin polypeptide; and, flanking said segment, (B) DNA from a nonessential region of a poxvirus genome. The viral vector can encode a selectable marker. In one example, the poxvirus includes, for example, a thymidine kinase gene (see U.S. Pat. No. 6,998,252, which is incorporated herein by reference).

Poxviral vectors that encode a C-terminal endostatin polypeptide include at least one expression control element operationally linked to the nucleic acid sequence encoding the C-terminal endostatin polypeptide. The expression control elements are inserted in the poxviral vector to control and regulate the expression of the nucleic acid sequence. Examples of expression control elements of use in these vectors includes, but is not limited to, lac system, operator and promoter regions of phage lambda, yeast promoters and promoters derived from polyoma, adenovirus, retrovirus or SV40. Additional operational elements include, but are not limited to, leader sequence, termination codons, polyadenylation signals and any other sequences necessary for the appropriate transcription and subsequent translation of the nucleic acid sequence encoding the C-terminal endostatin polypeptide in the host system. The expression vector can contain additional elements necessary for the transfer and subsequent replication of the expression vector containing the nucleic acid sequence in the host system. Examples of such elements include, but are not limited to, origins of replication and selectable markers. It will further be understood by one skilled in the art that such vectors are easily constructed using conventional methods (Ausubel et al., (1987) in "Current Protocols in Molecular Biology," John Wiley and Sons, New York, N.Y.) and are commercially available.

Basic techniques for preparing recombinant DNA viruses containing a heterologous DNA sequence encoding the C-terminal endostatin polypeptide, are known in the art. Such techniques involve, for example, homologous recombination between the viral DNA sequences flanking the DNA sequence in a donor plasmid and homologous sequences present in the parental virus (Mackett et al., 1982, Proc. Natl.

Acad. Sci. USA 79:7415-7419). In particular, recombinant viral vectors such as a poxviral vector can be used in delivering the gene. The vector can be constructed for example by steps known in the art, such as steps analogous to the methods for creating synthetic recombinants of the fowlpox virus 5 described in U.S. Pat. No. 5,093,258, incorporated herein by reference. Other techniques include using a unique restriction endonuclease site that is naturally present or artificially inserted in the parental viral vector to insert the heterologous DNA.

DNA sequences encoding a C-terminal endostatin polypeptide can be expressed in vitro by DNA transfer into a suitable host cell. The cell may be prokaryotic or eukaryotic. The term also includes any progeny of the subject host cell. It is understood that all progeny may not be identical to the 15 parental cell since there may be mutations that occur during replication. Methods of stable transfer, meaning that the foreign DNA is continuously maintained in the host, are known in the art

As noted above, a polynucleotide sequence encoding a 20 C-terminal endostatin polypeptide can be operatively linked to expression control sequences. An expression control sequence operatively linked to a coding sequence is ligated such that expression of the coding sequence is achieved under conditions compatible with the expression control sequences. 25 The expression control sequences include, but are not limited to, appropriate promoters, enhancers, transcription terminators, a start codon (i.e., ATG) in front of a protein-encoding gene, splicing signal for introns, maintenance of the correct reading frame of that gene to permit proper translation of 30 mRNA, and stop codons.

Hosts cells can include microbial, yeast, insect and mammalian host cells. Methods of expressing DNA sequences having eukaryotic or viral sequences in prokaryotes are well known in the art. Non-limiting examples of suitable host cells 35 include bacteria, archea, insect, fungi (for example, yeast), plant, and animal cells (for example, mammalian cells, such as human). Exemplary cells of use include Escherichia coli, Bacillus subtilis, Saccharomyces cerevisiae, Salmonella typhimurium, SF9 cells, C129 cells, 293 cells, Neurospora, 40 and immortalized mammalian myeloid and lymphoid cell lines. Techniques for the propagation of mammalian cells in culture are well-known (see, Jakoby and Pastan (eds), 1979, Cell Culture. Methods in Enzymology, volume 58, Academic Press, Inc., Harcourt Brace Jovanovich, N.Y.). Examples of 45 commonly used mammalian host cell lines are VERO and HeLa cells, CHO cells, and WI38, BHK, and COS cell lines, although cell lines may be used, such as cells designed to provide higher expression desirable glycosylation patterns, or other features. As discussed above, techniques for the trans- 50 formation of yeast cells, such as polyethylene glycol transformation, protoplast transformation and gene guns are also known in the art (see Gietz and Woods Methods in Enzymology 350: 87-96, 2002).

Transformation of a host cell with recombinant DNA can 55 be carried out by conventional techniques as are well known to those skilled in the art. Where the host is prokaryotic, such as, but not limited to, E. coli, competent cells which are capable of DNA uptake can be prepared from cells harvested after exponential growth phase and subsequently treated by 60 the CaCl₂ method using procedures well known in the art. Alternatively, MgCl₂ or RbCl can be used. Transformation can also be performed after forming a protoplast of the host cell if desired, or by electroporation.

When the host is a eukaryote, such methods of transfection 65 of DNA as calcium phosphate coprecipitates, conventional mechanical procedures such as microinjection, electropora-

tion, insertion of a plasmid encased in liposomes, or virus vectors can be used. Eukaryotic cells can also be co-transformed with polynucleotide sequences encoding a C-terminal endostatin polypeptide, and a second foreign DNA molecule encoding a selectable phenotype, such as the herpes simplex thymidine kinase gene. Another method is to use a eukaryotic viral vector, such as simian virus 40 (SV40) or bovine papilloma virus, to transiently infect or transform eukaryotic cells and express the protein (see for example, Eukaryotic Viral Vectors, Cold Spring Harbor Laboratory, Gluzman ed., 1982).

Therapeutic Methods and Pharmaceutical Compositions

The C-terminal endostatin polypeptides disclosed herein, or nucleic acids encoding the C-terminal endostatin polypeptides, can be used to treat fibrosis. In several examples, the C-terminal endostatin polypeptides, or nucleic acid encoding these polypeptides are of use to decrease fibrosis, such as in a subject. Thus, in several embodiments, the methods include administering to a subject a therapeutically effective amount of one or more of the C-terminal endostatin polypeptides disclosed herein, or polynucleotides encoding these polypeptides, in order to decrease fibrosis. In some examples, the C-terminal endostatin polypeptide comprises or consists of amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13. However, any of the C-terminal endostatin polypeptides disclosed herein can be used to decrease fibrosis. In some embodiments, the peptides can be administered as a unit dose.

Suitable subjects include those with a fibrosis of the skin or lungs, but fibrosis of any tissue can be treated using the methods disclosed herein. In one example, the subject has scleroderma. In other examples, the subject has idiopathic pulmonary fibrosis, morphea, fibrosis as a result of Graft-Versus-Host Disease (GVHD), a keloid or hypertrophic scar, subepithelial fibrosis, endomyocardial fibrosis, uterine fibrosis, myelofibrosis, retroperitoneal fibrosis, nephrogenic systemic fibrosis, scarring after surgery, asthma, cirrhosis/liver fibrosis, aberrant wound healing, glomerulonephritis, and multifocal fibrosclerosis.

In further examples, the methods are used to treat the systemic form of scleroderma, such as limited cutaneous scleroderma or diffuse cutaneous scleroderma (or systemic sclerosis). The methods can be used to treat the localized form of scleroderma, including morphea and linear scleroderma.

The methods can include selecting a subject in need of treatment, such as a subject with a fibrotic disease, such as scleroderma, idiopathic pulmonary fibrosis, morphea, a keloid scar, a hypertrophic scar, or subepithelial fibrosis. In exemplary applications, compositions are administered to a subject having a fibrotic disease, such as scleroderma, idiopathic pulmonary fibrosis, morphea, a keloid scar, a hypertrophic scar, or subepithelial fibrosis, or any of the disorders listed above, in an amount sufficient to reduce the fibrosis. Amounts effective for this use will depend upon the severity of the disease, the general state of the patient's health, and the robustness of the patient's immune system. In one example, a therapeutically effective amount of the compound is that which provides either subjective relief of a symptom(s) or an objectively identifiable improvement as noted by the clinician or other qualified observer.

A method is provided herein for decreasing skin thickness. The method includes administering a therapeutically effective amount of a C-terminal endostatin polypeptide, thereby decreasing skin thickness. In another embodiment, a method is provided for decreasing lung fibrosis. The method includes administering a therapeutically effective amount of a C-terminal endostatin polypeptide, thereby decreasing skin thickness. Any of the C-terminal endostatin polypeptides disclosed herein can be used in these methods. In some embodiments, the C-terminal endostatin polypeptide comprises, or consists 5 of, amino acids 133-180 of SEQ ID NO: 2, SEQ ID NO: 4 or SEQ ID NO: 13.

Methods are provided herein for decreasing lysyl oxidase (LOX), such as transforming growth factor (TGF)- β induced LOX. The method includes contacting a cell with an effective 10 amount of a C-terminal endostatin polypeptide, thereby decreasing LOX. The methods can be practiced in vivo or in vitro. In some embodiments, the methods include comparing the amount of LOX produced by a cell contacted with a C-terminal endostatin polypeptide to a control. The control 15 can be a standard value, or the amount of LOX produced by a cell not contacted with the C-terminal endostatin polypeptide, such as a cell contacted with a carrier.

Methods are provided herein for increasing matrix metalloproteinase-2 (MMP-2). The method includes contacting a 20 cell with an effective amount of a C-terminal endostatin polypeptide, thereby increasing MMP-2 production. The methods can be practiced in vivo or in vitro. In some embodiments, the methods include comparing the amount of MMP-2 produced by a cell contacted with a C-terminal endostatin 25 polypeptide to a control The control can be a standard value, or the amount of MMP-2 produced by a cell not contacted with the C-terminal endostatin polypeptide, such as a cell contacted with a carrier.

A C-terminal endostatin polypeptide can be administered 30 by any means known to one of skill in the art (see Banga, A., "Parenteral Controlled Delivery of Therapeutic Peptides and Proteins," in Therapeutic Peptides and Proteins, Technomic Publishing Co., Inc., Lancaster, Pa., 1995) either locally or systemically, such as by intradermal, intrathecal, intramuscu- 35 lar, subcutaneous, intraperitoneal or intravenous injection, but even oral, nasal, transdermal or anal administration is contemplated. In one embodiment, administration is by subcutaneous, intradermal, or intramuscular injection. In another embodiment, administration is by intraperitoneal or intrath- 40 ecal administration. To extend the time during which the peptide or protein is available to stimulate a response, the peptide or protein can be provided as an implant, an oily injection, or as a particulate system. The particulate system can be a microparticle, a microcapsule, a microsphere, a 45 nanocapsule, or similar particle. (see, e.g., Banga, supra).

For treatment of the skin, a therapeutically effective amount of at least one C-terminal endostatin polypeptide, or a nucleic acid encoding the peptide, can be locally administered to the affected area of the skin, such as in the form of an 50 ointment. In one embodiment, the ointment is an entirely homogenous semi-solid external agent with a firmness appropriate for easy application to the skin. Such an ointment can include fats, fatty oils, lanoline, VASELINE®, paraffin, wax, hard ointments, resins, plastics, glycols, higher alcohols, 55 glycerol, water or emulsifier and a suspending agent. Using these ingredients as a base, a decoy compound can be evenly mixed. Depending on the base, the mixture can be in the form of an oleaginous ointment, an emulsified ointment, or a watersoluble ointment oleaginous ointments use bases such as 60 plant and animal oils and fats, wax, VASELINE® and liquid paraffin. Emulsified ointments are comprised of an oleaginous substance and water, emulsified with an emulsifier. They can take either an oil-in-water form (O/W) or a water-in-oilform (W/O). The oil-in-water form (O/W) can be a hydro- 65 philic ointment. The water-in-oil form (W/O) initially lacks an aqueous phase and can include hydrophilic VASELINE

and purified lanoline, or it can contain a water-absorption ointment (including an aqueous phase) and hydrated lanoline. A water-soluble ointment can contain a completely watersoluble Macrogol base as its main ingredient.

Pharmaceutically acceptable carriers include a petroleum jelly, such as VASELINE®, wherein the petroleum jelly contains 5% stearyl alcohol, or petroleum jelly alone, or petroleum jelly containing liquid paraffin. Such carriers enable pharmaceutical compositions to be prescribed in forms appropriate for consumption, such as tablets, pills, sugarcoated agents, capsules, liquid preparations, gels, ointments, syrups, slurries, and suspensions. When locally administered into cells in an affected area or a tissue of interest, the at least one C-terminal endostatin polypeptide, or polynucleotide encoding the peptide can be administered in a composition that contains a synthetic or natural hydrophilic polymer as the carrier. Examples of such polymers include hydroxypropyl cellulose and polyethylene glycol. One or more C-terminal endostatin polypeptides, or polynucleotide encoding the polypeptides, can be mixed with a hydrophilic polymer in an appropriate solvent. The solvent is then removed by methods such as air-drying, and the remainder is then shaped into a desired form (for example, a sheet) and applied to the target site. Formulations containing such hydrophilic polymers keep well as they have a low water-content. At the time of use, they absorb water, becoming gels that also store well. In the case of sheets, the firmness can be adjusted by mixing a polyhydric alcohol with a hydrophilic polymer similar to those above, such as cellulose, starch and its derivatives, or synthetic polymeric compounds. Hydrophilic sheets thus formed can be used. A therapeutically effective amount of one or more C-terminal endostatin polypeptide, or polynucleotide encoding the peptide can also be incorporated into bandages and dressings.

For administration by inhalation, the C-terminal endostatin polypeptide, or polynucleotide encoding the peptide can be conveniently delivered in the form of an aerosol spray presentation from pressurized packs or a nebulizer, with the use of a suitable propellant, such as dichlorodifluoromethane, trichlorofluoromethane, dichlorotetrafluoroethane, carbon dioxide or other suitable gas. In the case of a pressurized aerosol, the dosage unit can be determined by providing a valve to deliver a metered amount. Capsules and cartridges for use in an inhaler or insufflator can be formulated containing a powder mix of the compound and a suitable powder base such as lactose or starch.

In some embodiments, the C-terminal endostatin polypeptide, such as, but not limited to E4, can be administered by inhalation. For example, the C-terminal endostatin polypeptide can be administered in an aerosolized form, such as using a nebulizer or a metered dose inhaler. Technologies of use include micropump nebulizers (such as the AEROGEN GO® system), jet nebulizers designed to produce large fine particle fractions (such as the PARILC STAR®), jet nebulizers developing less shear during atomization (such as the HUDSON MICROMIST®), and ultrasonic nebulizers (such as the DeVilbiss ULTRA-NEB®).

The endostatin polypeptide can be dissolved in a carrier, such as saline, and atomized using the devices above. The associated aerosols can be collected using a NEXT GEN-ERATION IMPACTOR® (NGI) (MSP Corp., Shoreview, Minn.), which uses a series of aerodynamic stages to separate and collect the aerosol into separate fractions based on droplet size. Since droplet size is the primary determinant of deposition location in the lungs, this device allows us to specifically isolate the portion of the liquid aerosol that will deposit in the small airways and alveoli. Aerosol particle size is often expressed in terms of mass median aerodynamic diameter (MMAD), a parameter that is based on particle size, shape, and density. For a spherical particle, MMAD is equal to MMD ($p^{1/2}$), in which MMD is mass median diameter and r is the bulk density. For a nonspherical particle, MMAD is equal to MMD (p/x)^{1/2}, in which X is the shape factor. Thus, particles with larger than unit density will have actual diameters smaller than their MMAD.

The site of particle deposition within the respiratory tract is demarcated based on particle size. In one example, particles 10 of about 1 to about 500 microns are utilized, such as particles of about 25 to about 250 microns, or about 10 to about 25 microns are utilized. In other embodiments, particles of about 1 to 50 microns are utilized. For use in a metered dose inhaler, for administration to lungs particles of less than about 10 15 microns, such as particles of about 2 to about 8 microns, such as about 1 to about 5 microns, such as particles of 2 to 3 microns, can be utilized.

A therapeutically effect amount of a C-terminal endostatin polypeptide, or polynucleotide encoding the peptide can be 20 administered in the pharmaceutically acceptable carrier. Pharmacologically acceptable carriers (e.g., physiologically or pharmaceutically acceptable carriers) are well known in the art, and include, but are not limited to buffered solutions as a physiological pH (e.g. from a pH of about 7.0 to about 8.0, 25 or at a pH of about 7.4). One specific, non-limiting example of a physiologically compatible buffered solution is phosphate buffered saline. Other pharmacologically acceptable carriers include penetrants, which are particularly suitable for pharmaceutical formulations that are intended to be topically 30 applied (for example in the application of surgical wounds to promote healing).

The pharmacological compositions disclosed herein facilitate the use of at least one C-terminal endostatin polypeptide, or polynucleotide encoding the peptide, either in vivo or ex 35 vivo, to decrease fibrosis. Such a composition can be suitable for delivery of the active ingredient to any suitable subject, and can be manufactured in a manner that is itself known, e.g., by means of conventional mixing, dissolving, granulating, emulsifying, encapsulating, entrapping or lyophilizing pro- 40 cesses. Pharmacological compositions can be formulated in a conventional manner using one or more pharmacologically (e.g., physiologically or pharmaceutically) acceptable carriers, as well as optional auxiliaries that facilitate processing of the active compounds into preparations which can be used 45 pharmaceutically. Proper formulation is dependent upon the route of administration chosen. Thus, for injection, the active ingredient can be formulated in aqueous solutions. For transmucosal administration, penetrants appropriate to the barrier to be permeated are used in the formulation. Such penetrants 50 are generally known in the art.

For oral administration, the active ingredient can be combined with carriers suitable for incorporation into tablets, pills, capsules, liquids, gels, syrups, slurries, suspensions and the like. The active ingredient can be formulated for 55 parenteral administration by injection, such as by bolus injection or continuous infusion. Such compositions can take such forms as suspensions, solutions or emulsions in oily or aqueous vehicles, and can contain formulatory agents such as suspending, stabilizing and/or dispersing agents. Other phar-60 macological excipients are known in the art.

Optionally, the at least one C-terminal endostatin polypeptide, or polynucleotide encoding the peptide can be contained within or conjugated with a heterologous protein, hydrocarbon or lipid, whether for in vitro or in vivo administration. 65 Co-administration can be such that the at least one C-terminal endostatin polypeptide, or polynucleotide encoding the pep36

tide is administered before, at substantially the same time as, or after the protein, hydrocarbon, or lipid. In one embodiment, the at least one C-terminal endostatin polypeptide, or polynucleotide encoding the peptide is administered at substantially the same time, as the protein, hydrocarbon, or lipid.

Other delivery systems can include time-release, delayed release or sustained release delivery systems. Such systems can avoid repeated administrations of the compositions of the invention described above, increasing convenience to the subject and the physician. Many types of release delivery systems are available and known to those of ordinary skill in the art. They include polymer based systems such as poly(lactideglycolide), copolyoxalates, polycaprolactones, polyesteramides, polyorthoesters, polyhydroxybutyric acid, and polyanhydrides. Microcapsules of the foregoing polymers containing drugs are described in, for example, U.S. Pat. No. 5,075,109. Delivery systems also include non-polymer systems, such as lipids including sterols such as cholesterol, cholesterol esters and fatty acids or neutral fats such as monodi- and tri-glycerides; hydrogel release systems; silastic systems; peptide based systems; wax coatings; compressed tablets using conventional binders and excipients; partially fused implants; and the like. Specific examples include, but are not limited to: (a) erosional systems in which the at least one C-terminal endostatin polypeptide, or polynucleotide encoding the peptide is contained in a form within a matrix such as those described in U.S. Pat. Nos. 4,452,775; 4,667,014; 4,748,034; 5,239,660; and 6,218,371 and (b) diffusional systems in which an active component permeates at a controlled rate from a polymer such as described in U.S. Pat. Nos. 3,832,253 and 3,854,480. In addition, pump-based hardware delivery systems can be used, some of which are adapted for implantation.

Use of a long-term sustained release implant may be particularly suitable for treatment of chronic conditions, such as scleroderma. Long-term release, as used herein, means that the implant is constructed and arranged to deliver therapeutic levels of the active ingredient for at least 30 days, and preferably 60 days. Long-term sustained release implants are well known to those of ordinary skill in the art and include some of the release systems described above. These systems have been described for use with oligodeoxynucleotides (see U.S. Pat. No. 6,218,371). For use in vivo, nucleic acids and peptides are preferably relatively resistant to degradation (such as via endo- and exo-nucleases). Thus, modifications, such as the inclusion of a C-terminal amide, can be used.

The therapeutically effective amount of C-terminal endostatin polypeptide, or polynucleotide encoding the peptide will be dependent on the C-terminal endostatin polypeptide, or polynucleotide encoding the peptide, that is utilized, the subject being treated, the severity and type of the affliction, and the manner of administration. For example, a therapeutically effective amount of a polynucleotide encoding the peptide can vary from about 0.01 µg per kilogram (kg) body weight to about 1 g per kg body weight, such as about 1 µg to about 5 mg per kg body weight, or about 5 µg to about 1 mg per kg body weight. The exact dose is readily determined by one of skill in the art based on the potency of the specific compound the age, weight, sex and physiological condition of the subject.

With regard to the administration of nucleic acids, one approach to administration of nucleic acids is direct treatment with plasmid DNA, such as with a mammalian expression plasmid. As described above, the nucleotide sequence encoding a N-terminal endostatin peptide can be placed under the control of a promoter to increase expression of the molecule.

When a viral vector is utilized for administration in vivo, it is desirable to provide the recipient with a dosage of each recombinant virus in the composition in the range of from about 10^5 to about 10^{10} plaque forming units/mg mammal, although a lower or higher dose can be administered. The 5 composition of recombinant viral vectors can be introduced into a mammal either prior to any evidence of a cancer, or to mediate regression of the disease in a mammal afflicted with the cancer. Examples of methods for administering the composition into mammals include, but are not limited to, expo-10sure of cells to the recombinant virus ex vivo, or injection of the composition into the affected tissue or intravenous, subcutaneous, intradermal or intramuscular administration of the virus. Alternatively the recombinant viral vector or combination of recombinant viral vectors may be administered locally 15 by direct injection into the cancerous lesion in a pharmaceutically acceptable carrier. Generally, the quantity of recombinant viral vector, carrying the nucleic acid sequence of one or more C-terminal endostatin polypeptides to be administered is based on the titer of virus particles. An exemplary range of 20 the immunogen to be administered is 10^5 to 10^{10} virus particles per mammal, such as a human.

In one specific, non-limiting example, a pharmaceutical composition for intravenous administration would include about 0.1 μ g to 10 mg of C-terminal endostatin polypeptide 25 per patient per day. Dosages from 0.1 up to about 100 mg per patient per day can be used, particularly if the agent is administered to a secluded site and not into the circulatory or lymph system, such as into a body cavity or into a lumen of an organ. Actual methods for preparing administrable compositions 30 will be known or apparent to those skilled in the art and are described in more detail in such publications as *Remingtons Pharmaceutical Sciences*, 19th Ed., Mack Publishing Company, Easton, Pa., 1995.

Single or multiple administrations of the compositions are 35 administered depending on the dosage and frequency as required and tolerated by the subject. In one embodiment, the dosage is administered once as a bolus, but in another embodiment can be applied periodically until a therapeutic result is achieved. Generally, the dose is sufficient to treat or 40 ameliorate symptoms or signs of disease without producing unacceptable toxicity to the subject. Systemic or local administration can be utilized.

In a further method, an additional agent is administered. In one example, this administration is sequential. In other 45 examples, the additional agent is administered simultaneously with the C-terminal endostatin polypeptide.

For the treatment of scleroderma, examples of additional agents that can be used with a C-terminal endostatin polypeptides include nifedipine, amlodipine, diltiazem, felodipine, or nicardipine. An investigational drug GLEEVEC®, is also used for the treatment of scleroderma. GLEEVEC® or other tyrosine kinase inhibitors can be used with the C-terminal endostatin polypeptides disclosed herein. Patients with lung involvement of scleroderma benefit from oxygen therapy; the C-terminal endostatin polypeptides disclosed herein can be administered with this therapy.

For the treatment of fibrosis of the skin and scleroderma, additional agents of use are d-penicillamine, colchicine, Relaxin, steroids, and cyclosporine. C-terminal endostatin ⁶⁰ polypeptides also can be used in combination with immunosuppressive agents. Additionally, the C-terminal endostatin polypeptides can be used with methotrexate, cyclophosphamide, azathioprine, mycophenolate, glitazones, endothelin receptor antagonists, or Fulvestrant (ICI-182,780). ⁶⁵

The disclosure is illustrated by the following non-limiting Examples.

38

EXAMPLES

Excessive deposition of extra cellular matrix (ECM) components such as fibronectin (FN) and type I collagen (Col1 α 1) by organ fibroblasts is defined as fibrosis. Organ fibrosis is the final common pathway for many diseases that result in end-stage organ failure. However, effective therapy for organ fibrosis is still unavailable (see, for example, Bjoraker et al., Am. J. Respir. Crit. Care. Med 2000; 157:199-20; Varga and Abraham, J Clin Invest 2007; 117:557-67; Wynn, J Clin Invest 2007; 117:524-29). Uncontrollable wound-healing responses, including acute and chronic inflammation, angiogenesis, activation of resident cells, and ECM remodeling, are thought to be involved in the pathogenesis of fibrosis (Wynn, J Clin Invest 2007; 117:524-29; Kalluri and Sukhatme, Curr Opin Nephrol Hypertens 2000; 9:413-8). TGF- β is the prototype fibrotic cytokine that is increased in fibrotic organs and contributes to the development of fibrosis by stimulating the synthesis of ECM molecules, activating fibroblasts to α -smooth muscle actin (α -SMA)-expressing myofibroblasts, and downregulating matrix metalloproteinases (MMPs) (Branton, Microbes Infect 1999; 1:1349-65; Varga and Pasche Nature Reviews Rheumatology 2009; 5:200-6). Despite high expectations, a clinical trial of a monoclonal anti-TGF- β antibody in patients with early SSc failed to show any efficacy (Varga and Pasche, Nature Reviews Rheumatology 2009; 5:200-6).

Endostatin is a 20-kDa internal fragment of the carboxyterminus of collagen XVIII. It was originally identified in the supernatant of a cultured murine hemangioendothelioma cell line with potent antiangiogenic activity (O'Reilly et al., Cell 1997; 88:277-85). Endostatin inhibits endothelial proliferation and tube formation in vitro, and tumor growth in vivo (Dhanabal et al., Biochem Biophys Res Commun 1999; 258: 345-52). Studies have been conducted to assess endostatin's anti-tumor properties, including clinical trials (Folkman, Exp Cell Res 2006; 312:594-607). The NH₂-terminal domain of endostatin has been reported as the functional domain responsible for inhibiting angiogenesis (Tjin Than Sjin et al., Cancer Res 2005; 65:3656-63). Although the exact molecular mechanism of its effect remains unclear, integrins, glypicans, flk-1, and nucleolin have been reported as endostatin receptors (Sudhakar et al., Proc Natl Acad Sci USA 2003; 100:4766-71; Karumanchi et al., Mol Cell 2001; 7:811-22). Recent studies have shown that endostatin is increased in serum and/or BALF obtained from IPF and SSc patients with pulmonary fibrosis (for example, Sumi, J Clin Lab Anal 2005; 19:146-9).

In the studies presented herein, the effects of endostatin on fibrosis were evaluated. The effect of endostatin and endostatin-derived peptides on fibrosis in vitro was assessed using primary human fibroblasts, ex vivo using human skin, and in vivo in mice skin treated with TGF- β . Surprisingly, the findings demonstrate that a carboxy-terminal peptide of endostatin has anti-fibrotic activity and provide a novel therapy for fibrotic disorders.

Example 1

Materials and Methods

Reagents and antibodies. The full-length recombinant human endostatin (rE) was purchased from Sigma-Aldrich (St. Louis, Mo.). Recombinant human TGF- β was from R&D Systems Inc. (Minneapolis, Minn.). Mouse monoclonal antihuman fibronectin (FN) antibody, goat polyclonal anti-human type I Collagen α I chain (Col1 α 1) antibody, and mouse monoclonal anti-human GAPDH antibody were from Santa Cruz Biotechnology (Santa Cruz, Calif.). Mouse monoclonal anti-human α -smooth muscle actin (α -SMA) antibody was from Sigma-Aldrich.

Synthesis of human endostatin peptides. Peptides were synthesized by the solid-phase on Liberty Microwave Syn- 5 thesizer (CEM Corporation, 3100 Smith Farm Road, Mathews, N.C. 28106) using FMOC synthesis protocol. Briefly, synthesis was performed by stepwise addition of activated amino acids to the solid support (Wang resin and PEG-PS) starting from the carboxy terminus to the amino 10 terminus. Activation of amino acids was performed by DIPEA/HOBT/TBTU chemistry. At the end of the synthesis, peptides were cleaved off the resin with reagent R (90% TFA, 5% Thioanisole, 3% Ethanedithiol, and 2% Anisole) and subjected to multiple ether extractions. The crude peptides 15 were analyzed, characterized, and purified by Gel filtration (G-25 column), Reversed-Phase High Performance Liquid Chromatography (RP-HPLC, 486 and 600E by Waters Corporation). The correct mass was confirmed by MALDI-TOF Mass Spectroscopy (The Voyager-DE STR Biospectrometry 20 Workstation). Sequences of the peptides are shown in Table 1 and correspond to amino acids 1-45 (E1); 71-115 (E2); 133-180 (E3), 133-180A (E4) which differs from E3 by the presence of a carboxy-terminal amide. The purity of all peptides was >98%. All peptides were dissolved in DMSO at a con- 25 centration 5 mg/ml, and diluted in $1 \times PBS$ to $1-20 \mu g/ml$.

Primary fibroblast culture. Human primary lung and skin fibroblasts were cultured. The explanted lungs of normal organ donors, patients with SSc or IPF, and clinically involved skin of SSc patients, a morphea patient and healthy 30 donors were used for primary fibroblast culture. Approximately 2-cm pieces of peripheral lung and skin were minced and fibroblasts were cultured in Dulbecco's modified Eagle's medium (DMEM; Mediatech, Herndon, Va.) supplemented with 10% FBS, penicillin, streptomycin, and anti-mycotic 35 agent, as previously described (Feghali et al., Arthritis Rheum 1999; 42:1451-7). All the cells were used between passages 3-6.

Western blot analysis. Cellular lysates were obtained from cultured fibroblasts as previously described (Pilewski et al., 40 Am J Pathol 2005; 166:399-407). Briefly, 2.0×10⁵ primary fibroblasts were cultured in 35-mm wells in 0.5% FBS-containing medium supplemented with 10 ng/ml of human recombinant TGF-β or PBS as vehicle control for 24 h, following which 5 µg/ml of human rE, endostatin peptides (E1- 45 E4), or DMSO (vehicle) was added for 48 h. In some experiments, endostatin peptides were used without $TGF-\beta$ stimulation. Cellular lysates were analyzed by western blot. Signals were detected following incubation with horseradish peroxidase-conjugated secondary antibody and chemilumi- 50 nescence (Perkin Elmer Life Sciences, Inc., Boston, Mass.). The intensity of individual bands with expected molecular sizes was semi-quantitatively analyzed using the image/J® software available at on the internet (/rsb.info.nih.gov/ij/index.html), and normalized to individual GAPDH intensity. 55

Ex vivo human skin assays. Human abdominal skin was obtained from corrective plastic surgery. As previously described (Yasuoka et al., The Open Rheumatol J 2008; 2:17-22), subcutaneous fat tissue was removed uniformly and skin tissue was cut into 1.5 cm×1.5 cm sections. The following 60 were injected intradermally in a total volume of 100 µl 1×PBS: rE alone (1-10 µg/ml), endostatin peptides alone (10 µg/ml), rE or endostatin peptides (1-20 µg/ml) in combination with TGF- β (10 ng/ml), and TGF- β alone (10 ng/ml). In some experiments, human skin was first injected with TGF- β for 48 65 h followed by recombinant endostatin (rE) administration in the same injection site as TGF- β . Independent experiments

were conducted in duplicate or triplicate as indicated in the figure legends. Explants containing complete epidermal and dermal layers were cultured in an air liquid interface with the epidermal and keratin layers side up and exposed to air. The culture medium was replaced every other day. After 1 or 2 weeks, skin tissue corresponding to an area with 8-mm diameter centered around the injection site was harvested using disposable 8-mm ACUPUNCH® (Acuderm Inc., Lauderdale, Fla.). Skin tissue was fixed in 10% formalin prior to embedding in paraffin.

In vivo mouse experiments. CB57BL6/J male mice were purchased from The Jackson Laboratory (Bar Harbor, Me.). Human rE (10 μ g/ml) or Endostatin peptides (10 μ g/ml) in combination with TGF- β (10 ng/ml), or TGF- β alone were injected intradermally on the back of mice in a total volume of 100 μ l 1×PBS. Mice were injected in two different skin sites and sacrificed one week post-injection. Skin surrounding the injection site was harvested and fixed in 10% formalin prior to embedding in paraffin.

Measurement of skin dermal thickness: Six μ m sections of paraffin-embedded human and mouse skin tissues were stained with hematoxylin and eosin (H&E). In some experiments, sections were stained with Masson trichrome which identifies collagens. Images were taken on a Nikon Eclipse 800 microscope. The thickness of the dermis was measured in 6 random fields of each section using the image/J® software. Data are shown in arbitrary units.

Tubular Formation Assay. The ability of endostatin peptide to inhibit angiogenesis was examined in tubular formation assay using MATRIGEL® culture. Human umbilical vein endothelial cells (HUVECs) were maintained in endothelial cell basal medium-2 (EBM-2; Clonetics, San Diego, Calif.) supplemented with EBM-2 MV SINGLEQUOTS®. HUVECs (5×10⁴) were cultured in duplicate on 24-well MATRIGEL® plates (BD Biosciences, San Diego, Calif.) alone, or in the presence of rE or E4 peptide (50 nM) in EBM-2 at 37° C. DMSO was used as vehicle control. After 24 hours, images were captured using a converted microscope. The degree of cord formation was quantified by measuring the area occupied by tubes in 6 random fields per well. Three independent experiments were performed.

Statistical analysis. All continuous variables were expressed as the mean±standard deviation. Comparisons between 2 groups were tested for statistical significance using the paired t-test or Mann-Whitney U test as appropriate. Comparison among 3 groups was performed using ANOVA followed by Bonferroni's test.

Example 2

Human Endostatin Inhibits FN and Col1α1 Production in TGF-β-treated Human Primary Lung and Skin Fibroblasts in vitro

To evaluate whether endostatin modulates production of ECM components in fibroblasts, FN and Col1 α 1 expression was examined in normal human lung fibroblasts by western blot analysis. Cells were treated with 5 µg/ml rE for 48 h with or without pre-stimulation with human TGF- β for 24 h. As shown FIG. 1A, rE dramatically reduced FN and Col1 α 1 levels in TGF- β pre-treated fibroblasts. To define the functional domain of endostatin that mediates its inhibitory effect, four different peptides were synthesized corresponding to different regions of endostatin (Table 1).

4	1			

CONH2

TABLE I
Amino acid sequence of human endostatin fragments.
E1(1-45 of SEQ ID NO: 2)
${\tt H-}^1{\tt HSHRDFQPVLHLVALNSPLSGGMRGIRGADFQCFQQARAVGLAGT^{45}-OH}$
E2 (71-115 of SEQ ID NO: 2)
$\texttt{H-}^{71}\texttt{IVNLKDELLFPSWEALFSGSEGPLKPGARIFSFDGKDVLRHPTWP^{115}-\texttt{OH}$
E3 (133-180 of SEQ ID NO: 2)
$\texttt{H-}^{\texttt{133}}\texttt{SYCETWRTEAPSATGQASSLLGGRLLGQSAASCH\texttt{HAYIVLCIENSFMT}^{\texttt{180}}-\texttt{OH}$
E4 (133-180A of SEQ ID NO: 2)
H- ¹³³ SYCETWRTEAPSATGOASSLLGGRLLGOSAASCHHAYIVLCIENSFMT ¹⁸⁰ -

As shown in FIGS. 1B and 1C, a fragment from the carboxy terminus of endostatin (E4) significantly suppressed FN and Col1 α 1 production in TGF- β treated cells compared with normal lung fibroblasts treated with TGF- β alone (P=0.03, in both comparisons). On the other hand, E1 peptide, located in the amino terminal region of endostatin, had no effect. In 25 addition to healthy fibroblasts, lung fibroblasts obtained from SSc and IPF patients, who had clinical lung fibrosis, were used in parallel assays with similar results (FIGS. 1B and 1C). Having demonstrated anti-fibrotic effects of rE and E4 in lung fibroblasts, the effects of these peptides was examined on skin 30 model to assess the effects of fibrogenic factors and for evalufibroblasts since skin is a major organ affected by fibrosis in SSc. Primary fibroblasts obtained from the skin of healthy controls, patients with systemic sclerosis (SSc) or localized scleroderma (morphea) were treated with rE or E4. Similarly to lung fibroblasts, rE and E4 reduced TGF- β -induced ECM production in dermal fibroblasts. Representative results are shown in FIG. 1D.

Example 3

Endostatin Peptides Reverse the Fibrotic Phenotype of Primary Lung Fibroblasts from Patients with SSc and IPF

Since it has been shown that TGF- β is upregulated in fibrotic tissue, it was examined if matrix production in fibrotic lung fibroblasts was altered by treatment with endostatin peptide in the absence of TGF- β stimulation. As shown in FIG. 1E left panel, both FN and Col1 α 1 levels decreased in 50 E4-treated fibroblasts. In addition, the same fibroblasts were treated with different concentrations of E4 to identify the optimal anti-fibrotic dose. E4 dose-dependently reduced Col1 α 1 levels when compared to vehicle control (FIG. 1E, right panel), but had a modest effect on FN levels. The reduc- 55 tion in ECM was more modest than that observed following TGF- β stimulation. Taken together, the results indicate that E4 can reduce baseline production of ECM components in fibroblasts from a fibrotic milieu and thus reverse the fibrotic phenotype. 60

Myofibroblasts, activated fibroblasts which express α -SMA, are induced by TGF- β stimulation and play a central role in fibrosis. Therefore, the effects of endostatin peptides on α-SMA expression in normal lung fibroblasts was examined. As shown in FIG. 1F, TGF-ß stimulation greatly 65 increased α-SMA expression. Interestingly, E4, and to a lesser extent E3, decreased TGF- β -induced α -SMA levels

suggesting that the carboxy-terminal region of endostatin can prevent the activation of fibroblasts and their transition to a myofibroblastic phenotype.

Example 4

Endostatin Reduces Dermal Thickness and Prevents TGF-β-induced Fibrosis in Human Skin

Cultured human skin explants can be used as an organ ating the efficacy of inhibitors/therapies to halt the progression of fibrosis and potentially reverse it (Yasuoka, The Open Rheumatol J 2008; 2:17-22). To evaluate the efficacy of endostatin as a potential therapeutic agent for fibrosis, this ex vivo human skin model was used. Since TGF- β is a wellknown pro-fibrotic factor that plays a central role in fibrosis, human recombinant TGF-ß was first injected intradermally to assess the level of fibrosis. As shown in FIG. 2A, TGF- β injection dramatically increased dermal thickness in a dosedependent manner one week post-injection. The fibrotic effect of TGF- β (10 ng/ml) resolved by two weeks. The baseline effects of rE (1, 5, and 10 μ g/ml) or endostatin peptides (10 µg/ml) were also examined individually. Although rE and E1-4 did not significantly alter dermal thickness, rE, E3, and E4 showed a tendency towards reduction in human dermal thickness (FIGS. 2B and 2C). It was determined if rE could inhibit fibrosis in TGF-β-treated human skin. TGF- β and rE were injected simultaneously. One week post-administration, rE in combination with TGF-β significantly reduced dermal thickness in a dose-dependent manner (FIG. 3). To assess the effects of rE on reversing fibrosis, the peptide was injected 2 days after TGF-ß administration. Similarly to co-treatment, delayed rE also significantly ameliorated TGF-\beta-induced dermal fibrosis. The findings indicate that human endostatin can prevent the development and progression of fibrosis and also reverse TGF-β-induced fibrosis in human skin.

Example 5

Endostatin Peptides Reduce TGF-\beta-induced Fibrosis in Human Skin ex vivo and Reverse Existing Fibrosis

To determine which part of endostatin is responsible for inhibiting TGF-\beta-induced fibrosis in human skin explants, endostatin peptides (10 µg/ml) were administrated in the

35

65

presence of 10 ng/ml of TGF- β . Representative images are shown in FIG. **4**A. E3 and E4 significantly abolished the development of fibrosis as measured by dermal thickness when compared to TGF- β alone (P=0.04, 0.01, respectively; FIG. **4**). The dermal thickness of skin explants injected with different concentrations of E1 or E4 in combination with TGF- β was examined. As shown in FIG. **5**, unlike E1, E4 at concentrations of 5-20 µg/ml clearly ameliorated TGF- β induced skin fibrosis, indicating that the C-terminus of endostatin can suppress fibrosis (see FIG. **17**).

Example 6

Endostatin Peptides Reduce TGF-β-induced Fibrosis in vivo in Mouse Skin

The anti-fibrotic effect of endostatin peptides was further assessed in vivo. rE and endostatin peptides in combination with TGF- β were injected in the skin of mice. One week post-injection, mice appeared healthy and showed no signs of ²⁰ distress. As shown in FIG. **6**, human TGF- β strongly increased dermal thickness in mouse skin (P=0.004). Peptides E3 and E4 from the carboxy terminus of human endostatin peptide prevented dermal fibrosis induced by TGF- β (P=0.01, 0.007, respectively). In addition, E2 significantly ²⁵ reduced dermal thickness (P=0.03). E1, a peptide corresponding to the amino terminus of endostatin did not alter TGF- β -induced dermal fibrosis. These results confirmed those obtained in our human skin model and emphasize the importance of the C-terminal domain of endostatin in pre- ³⁰ venting TGF- β -induced fibrosis in vivo and ex vivo.

Example 7

The C-terminal Peptide of Endostatin has Modest Anti-angiogenic Activity

The anti-angiogenic effect of endostatin has been attributed to its amino terminal domain (Tjin Tham Sjin et al., Cancer Res 2005; 65:3656-63). To evaluate the anti-angio-40 genic capacity of the carboxy terminal regions of endostatin, the effect of E4 on in vitro tubular formation was examined using MATRIGEL. As shown in FIG. 7, the capacity of rE to inhibit tubular structure formation by HUVECs was significant, confirming previous reports. On the other hand, the 45 ability of E4 to suppress angiogenesis was modest, suggesting that the region of endostatin corresponding to E4 does not significantly contribute to its anti-angiogenic activity.

Thus, E4, a peptide corresponding to the carboxy terminal region of endostatin, ameliorates TGF- β -induced fibrosis and 50 even reverses it. E4 suppressed TGF- β -induced ECM production and downregulated α -SMA levels in primary lung and skin fibroblasts. In vivo and ex vivo analyses revealed that E4 impedes the increase of skin dermal thickness triggered by TGF- β . Furthermore, the anti-angiogeneic capacity of E4 55 was low compared to that of rE. Taken together, the findings suggest that the domains of endostatin responsible for its anti-fibrotic and anti-angiogenic capacity are distinct. Other endostatin peptides (for example, E2 and E3) are shown to have anti-fibrotic activity. 60

The anti-angiogenic activity of endostatin has been the focus of numerous investigations directed at the development of anti-tumor therapy. Recently, elevated serum and BALF levels of endostatin in fibrotic disorders such as idiopathic pulmonary fibrosis (IPF) and systemic sclerosis (SSc) were reported. Endostatin levels were relatively increased in IPF patients with severe respiratory dysfunction and in SSc

patients with pulmonary fibrosis, severe skin fibrosis, and with cutaneous scars, compared to patients without those clinical manifestations (Sumi J Clin Lab Anal 2005; 19:146-9; Richter et al., Thorax 2009; 64:156-61). In addition, collagen XVIII expression was increased in cultured dermal fibroblasts of SSc patients (Tan et al., Arthritis Rheum 2005; 52:865-76) and in whole lung extracts of patients with IPF (Yang et al., Am J Respir Crit Care Med 2007; 175:45-54). In this regard, since endostatin is a proteolytic product of collagen XVIII cleaved by several proteases including MMPs and cathepsin L (Wen e al., Cancer Res 1999; 59:6052-6; Felbor, EMBO J 2000; 19:1187-94), and since MMPs are also upregulated in SSc and IPF (Richter et. al., Thorax 2009; 64:156-61, Toubi et al., Clin Exp Rheumatol 2002; 20:221-4), the observations that cleaved endostatin levels are elevated in those patients is plausible. However, it is unclear how endostatin may be involved in the pathogenesis of fibrosis.

Without being bound by theory, increased endostatin in fibrotic tissues may constitute a negative feedback regulatory loop which, although unsuccessful, is directed at halting the progression of fibrosis. Since endostatin was originally identified in aberrant "angiogenic" endothelial cancer cells as a product that likely controls/inhibits its "angiogenic" capacity (O'Reilly et al., Cell 1997; 88:277-85), it is plausible that endostatin in fibrosis serves a similar regulatory function.

Bloch W et al reported reduced connective tissue but normal vessel density in recombinant endostatin-treated mouse skin using a wound healing model (Bloch et al., FASEB J 2000; 14:2373-6). Furthermore, a peptide from the N-terminal region of endostatin prevented the progression of peritoneal sclerosis in a mouse model (Tanabe et al., Kidney Int 2007; 71:227-38); the peptide under investigation corresponded to the N-terminus of endostatin encompassing amino acids 1-27.

In contrast, the C-terminal region of endostatin, but not the N-terminus, is shown herein to be responsible for its antifibrotic effects. In fact, the peptide corresponding to the N-terminal domain of endostatin contributed to the fibrotic phenotype in some of the assays. Studies directed at defining the specific amino acid sequence responsible for endostatin's anti-angiogenic capacity (Richter et al., Thorax 2009; 64:156-61; Cattaneo et al., Exp Cell Res 2003; 283:230-6; Xu et al., Curr Protein Pept Sci 2008; 9:275-83) have shown that the entire angio-suppressive activity of endostatin was located in a 27-amino-acid peptide in the N-terminal domain (Richter et al., Thorax 2009; 64:156-61). Thus, the functional domain of endostatin that mediates its anti-fibrotic activity is different from that responsible for its anti-angiogenic capacity, implying different mechanisms for inhibiting angiogenesis and fibrosis. The anti-fibrotic C-terminal endostatin polypeptides disclosed herein are therefore capable of selectively inhibiting fribrosis without inhibiting angiogenesis. The C-terminal endostatin polypeptides can be used to more specifically and selectively target unwanted fibrosis without interfering with angiogenesis that may impact a desired therapeutic outcome.

The C-terminal endostatin polypeptide also reduces α -SMA expression in TGF- β -treated fibroblasts. In addition, the matrix reducing effects of E4 on normal fibroblasts was 60 modest compared to that in fibrotic fibroblasts. This suggests that the therapeutic effect of endostatin C-terminal peptide in fibrosis could be due, in part, to hindrance of fibroblast activation by TGF- β and other fibrosis promoting growth factors.

In 2005, ENDSTAR®, a recombinant human endostatin purified from *E. coli* containing an additional nine-amino acid sequence produced as a his-tagged protein was approved for the treatment of non-small-cell lung cancer in China (Sun et al., J Clin Oncol 2005 (ASCO Annual meeting proceedings); 23:7138). Despite its effectiveness, the treatment had several disadvantages including a requirement for high doses, the protein's short half-life, poor stability and easy inactivation (see, for example, Crystal, Nat Biotechnol 1999; 17:336-7; Hu et al., Acta Pharmacol Sin 2008; 29:1357-69). The small synthetic peptides disclosed herein could overcome these obstacles. E4 significantly inhibited fibrosis compared to rE and even E3 in vitro, in vivo, and ex vivo. In addition, E4 had minimal anti-angiogenic activity compared to rE, confirming 10 that the anti-angiogenic activity of endostatin resides in its N-terminal domain. The only difference between E3 and E4 was the presence of an amide-bond in the C-terminus of E4. Without being bound by theory, this amide renders the peptide more resistant to carboxy degradation by carboxypepti- 15 dases or other degrading molecules, thus stabilizing the peptide and likely maintaining its biological activity (Yang et. al. Am J Respir Crit Care Med 2007; 175:45-54).

Unfortunately, there are no effective therapies for organ fibrosis. The C-terminal domain of endostatin, corresponding ²⁰ to amino acid sequence 133-180 with amide-bond formation, suppressed ECM production by primary skin and lung fibroblasts and ameliorated dermal fibrosis induced by TGF- β in vivo and ex vivo in human skin. The findings presented herein demonstrate that E4 could be used for the treatment of fibrotic ²⁵ disorders, including IPF, SSc, morphea, as well as Graftversus-host disease, keloid and hypertrophic scar, and other organ fibrosis such as subepithelial fibrosis in asthma.

Example 8

Confirmation of the Efficacy of E4

E4, a peptide representing the carboxy terminus of human endostatin, can attenuate fibrosis triggered by multiple fibro- 35 genic factors. The anti-fibrotic effects of E4 can be detected whether administered concomitantly with or following the fibrogenic trigger. The efficacy of E4 was confirmed in four pre-clinical models of fibrosis: a) bleomycin-induced dermal fibrosis in vivo in mouse skin, b) TGF- β induced dermal 40 fibrosis in mouse skin, and c) bleomycin-induced pulmonary fibrosis. E4 peptide or a control peptide (E1; representing the amino terminal region of endostatin) were administered at the same time as TGF- β or bleomycin or 3-4 days following TGF- β or bleomycin. Mice were sacrificed one and two 45 weeks after TGF- β -initiation of dermal fibrosis, and two and three weeks after bleomycin-induced pulmonary fibrosis. Two different modes of administration of the E4 peptide were also tested. It was confirmed that intraperitoneal and intratracheal administration was effective. The amount of E4 that was 50 administered ws 10 μ g/ml in a total volume of 100 μ l for skin and IP injections and 50 µl for IT administration.

For these studies, fibrosis was assessed by measurement of dermal thickness on H&E skin sections (skin), assessment of collagen levels by Masson Trichrome staining (skin and 55 lung), and measurement of collagen levels by Sircol assay (lung). Furthermore, to confirm the mechanism by which E4 exerts its anti-fibrotic effects, the production of extra-cellular matrix (ECM) components, the levels of enzymes that promote matrix stabilization and thus accumulation and levels of 60 those that degrade ECM components, and levels of transcription factors downstream of the pro-fibrotic triggers were evaluated. Results were assessed using the unpaired t test and the 3-way ANOVA (for the ID1 data).

E4 caused a significant attenuation of bleomycin induced dermal fibrosis even with a single administration of E4 (FIG.

8). E4 caused a significant decrease of TGFbeta induced dermal fibrosis on day 7. Thus E4 prevents (FIG. **8**) and reverses (FIG. **9**) dermal fibrosis triggered by TGFbeta.

E4 administered concomitantly with bleomycin or three days following bleomycin caused a marked reduction in fibrosis and Masson Trichrome staining (see FIG. 9 and FIG. 10). E4 peptide given three days after bleomycin significantly reduced collagen levels in mouse lungs (FIG. 10, panel B).

E4 caused a statistically significant reduction in both TGF β and bleomycin induced skin (FIG. 8) and lung fibrosis (FIG. 10) regardless of the mode of administration. Intraperitoneal and intratracheal administration of E4 were both effective in blocking dermal and pulmonary fibrosis. For example, E4 caused a significant attenuation of bleomycin induced lung fibrosis on day 21 whether administered intraperitoneally or intratracheally (FIG. 11). Thus E4 is effective at reducing fibrosis irrespective of the administration mode.

The results also evidenced that E4 exerts its anti-fibrotic effects via multiple pathways. E4 reduces levels of lysyl oxidase (LOX), and enzyme responsible for the cross-linking of collagen, elastin, and other extracellular matrix (ECM) molecules and thus the stabilization of the ECM. E4 can make collagen less stable and more susceptible to proteolytic degradation. FIG. **12** shows lung sections of mice treated with bleomycin with or without E4.

E4-mediated reduction of LOX was detected also was detected in vitro. Normal lung fibroblasts in passage 4 were treated with vehicle, E4, TGFβ, or TGF-β followed 30 min-³⁰ utes later by E4 (FIG. **13**). Media conditioned by the fibroblasts were analyzed using western blot analysis after 48 hrs. Treatment with E4 significantly reduced the level of LOX. Similar results were obtained when LOX mRNA levels were examined by real-time PCR

E4 also promotes the degradation of ECM components via induction and activation of matrix metalloprotease (MMP-2), an enzyme that degrades several ECM molecules including fibronectin and native and denatured collagens (FIG. 14). In addition, E4 increases levels of inhibitor of differentiation (ID)-1, a transcription factor that inhibits TGF- β effects (see FIG. 15). It was determined in a Western bloat analysis that E4 reduces the levels of the master switch transcription factor, Egr-1 (see FIG. 16) in primary human lung fibroblasts, treated and harvested after 24 hours. The reduction of Egr-1 levels parallels a reduction in collagen, SMA and fibronectin. Egr-1 is known to mediate the effects of several fibrotic agents (including TGF- β and bleomycin).

Thus, E4 exerted significant anti-fibrotic effects. This peptide significantly attenuates the fibrogenic effects of TGFbeta and bleomycin whether administered simultaneously with these fibrotic triggers or a few days following the initiation of fibrosis, suggesting that E4, and other C-terminal endostatin polypeptides is also effective at reversing established fibrosis. The anti-fibrotic effects of E4 were noted whether it was administered intratracheally or intraperitoneally to mice in which pulmonary fibrosis was induced by bleomycin and dermal fibrosis was induced by TGFbeta. Furthermore, E4 exerted its anti-fibrotic effects via multiple pathways that include destabilization of ECM through reduction of LOX and thus decrease of ECM crosslinking, induction of ECM degradation via activation of MMP-2, suppression of Egr-1 levels, and induction of the TGF^β, thereby inhibiting transcription factor ID-1.

Thus, several in vitro assays and four in vivo and ex vivo pre-clinical models of fibrosis suggest that C-terminal endostatin polypeptides, as exemplified by E4, are an effective anti-fibrotic peptide that can block and reverse fibrosis in two organs, lung and skin. These anti-fibrotic effects as well as the lack of anti-angiogenic effects characteristic of endostatin render E4 an attractive therapeutic peptide for organ fibrosis. **48**

It will be apparent that the precise details of the methods or compositions described may be varied or modified without departing from the spirit of the described invention. We claim all such modifications and variations that fall within the scope and spirit of the claims below.

SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 13 <210> SEQ ID NO 1 <211> LENGTH: 555 <212> TYPE: DNA <213> ORGANISM: Homo sapiens <400> SEQUENCE: 1 atgcacagec accgegactt ccageeggtg ctecacetgg ttgegeteaa cageeeectg 60 tcaggeggca tgeggggcat cegeggggee gaetteeagt getteeagea ggegegggee 120 gtgggggetgg egggeacett eegegeette etgteetege geetgeagga eetgtacage 180 240 atcgtgcgcc gtgccgaccg cgcagccgtg cccatcgtca acctcaagga cgagctgctg tttcccaget gggaggetet gtteteagge tetgagggte egetgaagee eggggeaege 300 atetteteet ttaacggcaa ggacgteetg acceaceeca eetggeeeca gaagagegtg 360 tggcatggct cggaccccaa cgggcgcagg ctgaccgaga gctactgtga gacgtggcgg 420 acqqaqqctc cctcqqccac qqqccaqqcc tactcqctqc tqqqqqqcaq qctcctqqqq 480 cagagtgccg cgagctgcca tcacgcctac atcgtgctat gcattgagaa cagcttcatg 540 actgcctcca agtag 555 <210> SEO ID NO 2 <211> LENGTH: 183 <212> TYPE: PRT <213> ORGANISM: Homo sapiens <400> SEOUENCE: 2 His Ser His Arg Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn 5 10 1 15 Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln 25 20 30 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala 40 35 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala 55 Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe 65 70 80 Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro 85 90 95 Gly Ala Arg Ile Phe Ser Phe Asn Gly Lys Asp Val Leu Thr His Pro 100 105 110 Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg 115 120 125 Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser 130 135 140 Ala Thr Gly Gln Ala Tyr Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln 145 150 155 160 Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn 165 170 175

Ser Phe Met Thr Ala Ser Lys 180

<210> SEQ ID NO 3 <211> LENGTH: 552 <212> TYPE: DNA <213> ORGANISM: Mus musculus <400> SEQUENCE: 3 catactcatc aggactttca gccagtgctc cacctggtgg cactgaacac ccccctgtct 60 ggaggcatgc gtggtatccg tggagcagat ttccagtgct tccagcaagc ccgagccgtg 120 180 gggetgtegg geacetteeg ggettteetg teetetagge tgeaggatet etatageate gtgcgccgtg ctgaccgggg gtctgtgccc atcgtcaacc tgaaggacga ggtgctatct 240 cccagctggg actccctgtt ttctggctcc cagggtcaac tgcaacccgg ggcccgcatc 300 ttttcttttg acggcagaga tgtcctgaga cacccagcct ggccgcagaa gagcgtatgg 360 cacggetegg acceeagtgg geggaggetg atggagagtt actgtgagae atggegaact 420 gaaactactg gggctacagg tcaggcctcc tccctgctgt caggcaggct cctggaacag 480 aaagetgega getgecacaa cagetacate gteetgtgea ttgagaatag etteatgace 540 tctttctcca aa 552 <210> SEQ ID NO 4 <211> LENGTH: 184 <212> TYPE: PRT <213> ORGANISM: Mus musculus <400> SEOUENCE: 4 His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn 10 1 5 15 Thr Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln 20 25 30 Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ser Gly Thr Phe Arg Ala 35 40 45 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala 50 55 60 Asp Arg Gly Ser Val Pro Ile Val Asn Leu Lys Asp Glu Val Leu Ser 65 70 75 80 Pro Ser Trp Asp Ser Leu Phe Ser Gly Ser Gln Gly Gln Leu Gln Pro 85 90 95 Gly Ala Arg Ile Phe Ser Phe Asp Gly Arg Asp Val Leu Arg His Pro 105 100 110 Ala Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Ser Gly Arg 115 125 120 Arg Leu Met Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Thr Thr Gly 130 135 140 Ala Thr Gly Gln Ala Ser Ser Leu Leu Ser Gly Arg Leu Leu Glu Gln 155 145 150 160 Lys Ala Ala Ser Cys His Asn Ser Tyr Ile Val Leu Cys Ile Glu Asn 165 170 175 Ser Phe Met Thr Ser Phe Ser Lys 180

<210> SEQ ID NO 5 <211> LENGTH: 699 <212> TYPE: DNA <213> ORGANISM: Homo sapiens

<400)> SH	EQUEI	NCE :	5												
gago	ccaa	aat d	cttgi	gaca	aa aa	actca	acaca	a tgo	ccca	cgt	gcco	cagea	acc 1	cgaa	ctcctg	60
aaaa	gaco	cgt d	cagto	ette	ct ct	tcc	cccca	a aaa	accca	aagg	acad	ccct	cat 🤉	gatci	cccgg	120
acco	ctga	agg t	tcaca	atgco	gt g	gtggt	ggad	c gto	gageo	cacg	aaga	accct	cga 🤉	ggtca	aagttc	180
aact	ggta	acg t	tgga	cggc	gt g	gaggt	cgcat	c aat	geea	aaga	caaa	ageeq	geg (ggago	gagcag	240
taca	acaç	gca (cgta	ccgt	gt ge	gtcaç	gcgto	c cto	cacco	gtcc	tgca	accaç	gga (ctgg	ctgaat	300
ggca	agga	agt a	acaa	gtgca	aa go	gtcto	ccaa	c aaa	ageed	ctcc	cago	cccc	cat	cgaga	aaaacc	360
atct	ccaa	aag d	ccaaa	agggo	ca go	cccc	gagaa	a cca	acago	gtgt	acad	ccct	gee (cccat	cccgg	420
gatg	jaget	cga d	ccaa	gaaco	ca go	gtcaç	geete	g aco	ctgco	tgg	tcaa	aaggo	ett (ctato	cccagc	480
gaca	tcgo	ccg t	tggaq	gtgg	ga ga	agcaa	atggg	g caq	geegé	gaga	acaa	actad	caa q	gacca	acgcct	540
cccg	gtgct	gg a	actco	cgaco	gg ct	cctt	cctto	c cto	ctaca	agca	agci	caco	gt 🤅	ggaca	aagagc	600
aggt	ggca	agc a	aggg	gaaco	gt ci	tcto	catgo	c tco	gtga	atgc	atga	agget	cct q	gcaca	aaccac	660
taca	icgca	aga a	agago	cctci	ce eq	ctgto	ctcc	g ggt	caaat	ga						699
<210 <211 <212 <213 <400)> SH .> LH :> T :> OF :> OF	EQ II ENGTH YPE : RGANI EQUEI	D NO H: 24 PRT ISM: NCE:	6 14 Homo 6	o saj	pien:	3									
Glu 1	Pro	Lys	Ser	Cys 5	Asp	Lys	Thr	His	Thr 10	Cya	Pro	Pro	Сүз	Pro 15	Ala	
Pro	Glu	Leu	Leu 20	Gly	Gly	Pro	Ser	Val 25	Phe	Leu	Phe	Pro	Pro 30	Lys	Pro	
Lys	Asp	Thr 35	Leu	Met	Ile	Ser	Arg 40	Thr	Pro	Glu	Val	Thr 45	Сүз	Val	Val	
Val	Asp 50	Val	Ser	His	Glu	Asp 55	Pro	Glu	Val	Lys	Phe 60	Asn	Trp	Tyr	Val	
Asp 65	Gly	Val	Glu	Val	His 70	Asn	Ala	Lys	Thr	Lys 75	Pro	Arg	Glu	Glu	Gly 80	
Ile	Asn	Tyr	Asn	Ser 85	Thr	Tyr	Arg	Val	Val 90	Ser	Val	Leu	Thr	Val 95	Leu	
His	Gly	Ile	Asn 100	Asp	Trp	Leu	Asn	Gly 105	Lys	Glu	Tyr	Lys	Cys 110	Lys	Val	
Ser	Asn	Lys 115	Ala	Leu	Pro	Ala	Pro 120	Ile	Glu	Lys	Thr	Ile 125	Ser	Lys	Ala	
Lys	Gly 130	Gly	Ile	Asn	Pro	Arg 135	Glu	Pro	Gln	Val	Tyr 140	Thr	Leu	Pro	Pro	
Ser 145	Arg	Aab	Glu	Leu	Thr 150	Lys	Asn	Gln	Val	Ser 155	Leu	Thr	Сүз	Leu	Val 160	
Lys	Gly	Phe	Tyr	Pro 165	Ser	Asp	Ile	Ala	Val 170	Glu	Trp	Glu	Ser	Asn 175	Gly	
Gly	Ile	Asn	Pro 180	Glu	Asn	Asn	Tyr	Lys 185	Thr	Thr	Pro	Pro	Val 190	Leu	Aab	
Ser	Asp	Gly 195	Ser	Phe	Phe	Leu	Tyr 200	Ser	Lys	Leu	Thr	Val 205	Asp	Lys	Ser	
Arg	Trp 210	Gln	Gly	Ile	Asn	Gly 215	Asn	Val	Phe	Ser	Сув 220	Ser	Val	Met	His	
Glu 225	Ala	Leu	His	Asn	His 230	Tyr	Thr	Gly	Ile	Asn 235	Lys	Ser	Leu	Ser	Leu 240	

Ser Pro Gly Lys

<210> SEQ ID NO 7 <211> LENGTH: 48 <212> TYPE: DNA <213> ORGANISM: Homo sapiens <400> SEQUENCE: 7 tctagaggtg gtctagtgcc gcgcggcagc ggttcccccg ggttgcag <210> SEQ ID NO 8 <211> LENGTH: 16 <212> TYPE: PRT <213> ORGANISM: Homo sapiens <400> SEQUENCE: 8 Ser Arg Gly Gly Leu Val Pro Arg Gly Ser Gly Ser Pro Gly Leu Gln 10 1 5 15 <210> SEQ ID NO 9 <211> LENGTH: 48 <212> TYPE: PRT <213> ORGANISM: Rattus rattus <400> SEOUENCE: 9 Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Thr Gly Val Thr Gly Gln 1 5 10 15 Ala Ser Ser Leu Leu Ser Gly Arg Leu Leu Glu Gln Lys Ala Glu Ser 20 25 30 Cys His Asn Ser Tyr Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr 35 40 45 <210> SEQ ID NO 10 <211> LENGTH: 48 <212> TYPE: PRT <213> ORGANISM: Bos taurus <400> SEQUENCE: 10 Ser Tyr Cys Glu Thr Trp Arg Thr Asp Ser Arg Ala Ala Thr Gly Gln 1 5 10 15 Ala Ser Ser Leu Leu Ala Gly Arg Leu Leu Glu Gln Lys Ala Ala Gly 20 25 30 Cys His Asn Ala Phe Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr 35 40 45 <210> SEQ ID NO 11 <211> LENGTH: 48 <212> TYPE: PRT <213> ORGANISM: Homo sapiens <400> SEOUENCE: 11 As
n Tyr Cys Glu Ala Trp Arg Thr Ala Asp Thr Ala Val Thr Gly Leu $\$ 1 5 10 15 Ala Ser Pro Leu Ser Thr Gly Lys Ile Leu Asp Gln Lys Ala Tyr Ser 20 25 30 Cys Ala Asn Arg Leu Ile Val Leu Cys Ile Glu Asn Ser Phe Met Thr 35 40 45 <210> SEQ ID NO 12 <211> LENGTH: 552 <212> TYPE: DNA

<213> ORGANISM: Homo sapiens

48

-continued

<400> SEQUENCE: 12	
cacagecace gegaetteea geeggtgete cacetggttg egeteaacag ecceetgtea	60
ggeggeatge gggggeateeg eggggeegae tteeagtget teeageagge gegggeegtg	120
gggetggegg geaeetteeg egeetteetg teetegegee tgeaggaeet gtacageate	180
gtgegeegtg eegaeegege ageegtgeee ategteaace teaaggaega getgetgttt	240
cccagctggg aggetetgtt etcaggetet gagggteege tgaageeegg ggeaegeate	300
ttctcctttg acggcaagga cgtcctgagg caccccacct ggccccagaa gagcgtgtgg	360
catggetegg acceeaaegg gegeaggetg acegagaget aetgtgagae gtggeggaeg	420
gaggeteeet eggeeaeggg eeaggeetee tegetgetgg ggggeagget eetggggeag	480
agtgeegega getgeeatea egeetaeate gtgetetgea ttgagaaeag etteatgaet	540
gcctccaagt ag	552
<210> SEQ ID NO 13 <211> LENGTH: 183 <212> TYPE: PRT <213> ORGANISM: Homo sapiens	
View Segurate. 19	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	
Ser Pro Leu Ser Gly Gly Met Arg Gly Ile Arg Gly Ala Asp Phe Gln 20 25 30	
Cys Phe Gln Gln Ala Arg Ala Val Gly Leu Ala Gly Thr Phe Arg Ala 35 40 45	
Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala 50 55 60	
Asp Arg Ala Ala Val Pro Ile Val Asn Leu Lys Asp Glu Leu Leu Phe 65 70 75 80	
Pro Ser Trp Glu Ala Leu Phe Ser Gly Ser Glu Gly Pro Leu Lys Pro 85 90 95	
Gly Ala Arg Ile Phe Ser Phe Asp Gly Lys Asp Val Leu Arg His Pro 100 105 110	
Thr Trp Pro Gln Lys Ser Val Trp His Gly Ser Asp Pro Asn Gly Arg 115 120 125	
Arg Leu Thr Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Ala Pro Ser130135140	
Ala Thr Gly Gln Ala Ser Ser Leu Leu Gly Gly Arg Leu Leu Gly Gln145150150155	
Ser Ala Ala Ser Cys His His Ala Tyr Ile Val Leu Cys Ile Glu Asn 165 170 175	
Ser Phe Met Thr Ala Ser Lys 180	

The invention claimed is:

1. A composition comprising a polypeptide, wherein the polypeptide is 48, 49, 50, 51, 52 or 53 amino acids in length and comprises the amino acid sequence set forth as amino 60 acids 133-180 of SEQ ID NO: 2, or the amino acid sequence set forth as SEQ ID NO: 2 with at most 5 amino acid substitutions, wherein the polypeptide has anti-fibrotic activity, wherein the polypeptide is optionally amidated, and wherein the polypeptide does not comprise the amino acid sequence 65 set forth as amino acids 1-92 of SEQ ID NO: 2; and

a pharmaceutically acceptable carrier.

2. The composition of claim **1**, wherein the polypeptide consists essentially of

- a) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2; or
- b) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions;
- wherein the carboxy terminus of the polypeptide is optionally amidated.

3. The composition of claim **1**, wherein the polypeptide consists of:

35

(a) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2;

- (b) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions; or
- (c) either (a) or (b), wherein the carboxy terminus of the polypeptides is amidated.

4. The composition of claim **1**, wherein the polypeptide comprises the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13, wherein the carboxy terminus of ¹⁰ the polypeptide is optionally amidated.

5. The composition of claim **1**, wherein the carboxy terminus of the polypeptide is amidated.

6. The composition of claim **1**, wherein the polypeptide 15 comprises amino acids 133-180 of SEQ ID NO: 2.

7. The composition of claim 1, wherein the polypeptide consists of amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions.

8. The composition of claim **1**, comprising a therapeuti- $_{20}$ cally effective amount of the polypeptide.

9. The composition of claim **1**, wherein the polypeptide comprises additional consecutive amino acids of SEQ ID NO: 2.

10. An isolated polypeptide, wherein the polypeptide is 48, $_{25}$ 49, 50, 51, 52 or 53 amino acids in length and comprises the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2, or the amino acid sequence set forth as SEQ ID NO: 2 with at most 5 amino acid substitutions and wherein the polypeptide has anti-fibrotic activity, and wherein the $_{30}$ polypeptide is optionally amidated.

11. The isolated polypeptide of claim **10**, wherein the carboxy terminus of the polypeptide is amidated.

12. The isolated polypeptide of claim **10**, wherein the polypeptide consists essentially of:

- a) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2; or
- b) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions;

wherein the carboxy terminus of the polypeptide is optionally amidated.

13. The isolated polypeptide of claim **10**, wherein the polypeptide consists of:

- (a) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2;
- (b) the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions; or
- (c) either (a) or (b), wherein the carboxy terminus of the polypeptides is amidated.

14. The isolated polypeptide of claim 10, wherein the polypeptide comprises the amino acid sequence set forth as amino acids 133-180 of SEQ ID NO: 13, wherein the carboxy terminus of the polypeptide is optionally amidated.

15. The isolated polypeptide of claim **10**, wherein the polypeptide comprises amino acids 133-180 of SEQ ID NO: 2.

16. The isolated polypeptide of claim **10**, wherein the polypeptide consists of amino acids 133-180 of SEQ ID NO: 2 with at most 5 amino acid substitutions.

17. An isolated polynucleotide encoding the polypeptide of claim 10, operably linked to a heterologous promoter.

18. An expression vector comprising the isolated polynucleotide of claim **17**.

19. An isolated host cell transformed with the expression vector of claim **18**.

20. A method for decreasing lysyl oxidase production by a cell, comprising

- contacting the cell with an effective amount of the composition of claim 1, thereby decreasing the production of lysyl oxidase by the cell.
- **21**. The method of claim **20**, wherein the cell is in vitro.

22. A method for increasing the production of matrix metalloprotease by a cell, comprising

contacting the cell with an effective amount of the composition of claim 1, thereby increasing the production of matrix metalloprotease by the cell.

23. The method of claim 22, wherein the cell is in vitro.

* * * * *