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(54) Title: METHODS FOR TREATING PRESSURE INDUCED OPTIC NEUROPATHY, PREVENTING NEURONAL DEGENERATION AND PROMOTING NEURONAL CELL, SURVIVAL VIA ADMINISTRATION OF LINGO-1 ANTAGONISTS AND TRKB AGONISTS

(57) Abstract: This invention relates to methods for promoting neuronal survival and regeneration using LINGO-1 antagonists and TrkB agonists. Additionally, the invention relates to methods for treating pressure induced optic neuropathies using LINGO-1 antagonists. The invention also relates generally to methods for increasing TrkB activity and inhibiting JNK pathway signaling using a LINGO-1 antagonist.



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USE OF LINGO-1 AND TRKB ANTAGONISTS

BACKGROUND OF THE INVENTION

Field of the Invention

[0001] This invention relates to neurology, neurobiology, molecular biology and pharmacology. More particularly, this invention relates to methods for promoting neuronal survival and regeneration using LINGO-1 antagonists and TrkB asdfsafsdagonists. Additionally, the invention relates to methods for treating pressure induced optic neuropathies using LINGO-1 antagonists. The invention also relates generally to methods for increasing TrkB activity and inhibiting JNK pathway signaling using LINGO-1 antagonists.

Background Art

[0002] Optical neuropathies are a group of eye diseases encompassing various clinical presentations and etiologies. Glaucoma is an exemplary optical neuropathy which includes pathological changes in the optic nerve, visible on the optic disk, and corresponding visual field loss, resulting in blindness if untreated. Glaucoma is associated with increased intraocular pressure, but other factors are involved.

[0003] Current therapies for glaucoma are directed at decreasing intraocular pressure. Medical therapy includes topical ophthalmic drops or oral medications that reduce the production or increase the outflow of intraocular fluid. However, these drug therapies for glaucoma are sometimes associated with significant side effects, such as headache, blurred vision, allergic reactions, death from cardiopulmonary complications and potential interactions with other drugs. Surgical therapies also are used, but they also have numerous disadvantages and modest success rates.

[0004] Accordingly, there remains a need for additional treatment methods for pressure induced optical neuropathies, including glaucoma and other conditions characterized by degeneration or death of retinal ganglion cells (RGCs).

BRIEF SUMMARY OF THE INVENTION

[0005] The present invention is based on the discovery that LINGO-1 interacts with and inhibits TrkB in neurons. After ocular hypertension, the TrkB ligand brain-derived

neurotrophic factor (BDNF) is upregulated. BDNF promotes TrkB phosphorylation and activation of a cell survival signaling pathway. However, LINGO-1 is also upregulated after ocular hypertension. The experiments described here use LINGO-1 antagonists to show that LINGO-1 inhibits the phosphorylation and activation of TrkB, thereby inhibiting the cell survival signaling pathway. The experiments described here also show that LINGO-1 promotes activity of the JNK signaling pathway, which is associated with cell death. Antagonists of LINGO-1 and agonists of TrkB, therefore promote cell survival.

[0006] Based on these discoveries, the invention relates generally to methods for inhibiting LINGO-1-TrkB interaction by administration of a LINGO-1 antagonist. The invention also relates to methods for promoting TrkB phosphorylation and TrkB signaling and for inhibiting JNK phosphorylation and JNK signaling by administration of a LINGO-1 antagonist. Furthermore, the invention relates to methods of promoting retinal ganglion cell survival or treating a pressure induced ocular neuropathy by administration of a LINGO-1 antagonist. Additionally, the invention relates to methods for promoting neuronal cell survival by administration of a LINGO-1 antagonist and a TrkB agonist. In certain embodiments, the invention also relates to methods for treating conditions associated with neuronal cell death by administration of LINGO-1 antagonist and a TrkB agonist.

[0007] In certain embodiments, the invention includes a method for inhibiting LINGO-1 and TrkB interaction in a cell comprising contacting a cell which co-expresses LINGO-1 and TrkB with LINGO-1 antagonist. In other embodiments, the invention includes a method for promoting TrkB phosphorylation or TrkB pathway signaling in a cell comprising contacting a cell which coexpresses LINGO-1 and TrkB with a LINGO-1 antagonist. In certain other embodiments, the invention includes a method for promoting TrkB phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.

[0008] In certain embodiments, the invention includes methods for promoting JNK phosphorylation in a cell comprising contacting a cell which expresses LINGO-1 and JNK with a LINGO-1 antagonist. In other embodiments, the invention provides a method for inhibiting JNK phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.

[0009] In certain embodiments, the invention includes methods for promoting survival of a neuron at risk of dying comprising contacting the neuron with an effective amount of a LINGO-1 antagonist and a TrkB agonist.

[0010] In other embodiments, the invention includes methods for promoting survival of retinal ganglion cells in a mammal displaying signs or symptoms of a pressure induced ocular neuropathy comprising administering to a mammal in need of such a treatment an effective amount of a LINGO-1 antagonist and a carrier. The invention also includes a method for treating a disease or disorder associated with neuronal cell death comprising administering to a mammal in need of such treatment an effective amount of a LINGO-1 antagonist and a TrkB agonist. In some embodiments the mammal has been diagnosed with glaucoma.

[0011] In various embodiments of the above methods, the TrkB agonist may be any molecule which increases the ability of TrkB to promote neuronal survival. In certain embodiments, the TrkB agonist is selected from the group consisting of a TrkB agonist compound, a TrkB agonist polypeptide, a TrkB agonist antibody or fragment thereof, a TrkB agonist polynucleotide, a TrkB aptamer, or a combination of two or more TrkB agonists.

[0012] In certain embodiments, the TrkB agonist is a TrkB agonist compound. Certain TrkB agonist compounds of the invention include, but are not limited to, L-783,281 adenosine and CGS 21680. In addition, the TrkB agonist compounds can be small molecules that mimic critical regions of neurotrophins. For example, the small molecule can be a mimetic of a BDNF β -turn loop. Particular examples of small molecule mimetics that may be used according to the invention are disclosed in U.S. Published Application No. 2007/0060526 A1, which is incorporated herein by reference in its entirety.

[0013] In certain embodiments, the TrkB agonist is a TrkB agonist polypeptide. Certain TrkB agonist polypeptides of the invention include, but are not limited to, BDNF, NT-3 and NT-4/5. In certain embodiments, the TrkB agonist polypeptide comprises SEQ ID NO:4. In some embodiments, the TrkB agonist is a fusion polypeptide or conjugate comprising a non-TrkB-agonist heterologous polypeptide or polymer. In some embodiments, the non-TrkB-agonist polypeptide or polymer is selected from the group consisting of polyethylene glycol, a 1-acyl-glycerol derivative, an antibody Ig peptide, a serum albumin peptide, a targeting peptide, a reporter peptide, and a purification-facilitating peptide. In some embodiments, the antibody Ig peptide is a hinge and Fc peptide.

[0014] In alternative embodiments, the TrkB agonist is an antibody or fragment thereof which binds to a TrkB polypeptide. TrkB agonist antibodies for use in the methods of the

present invention include, but are not limited to, 6E2, 7F5, 11E1, 16E11, 17D11, 19E12, 29D7.

- [0015] In other embodiments, the TrkB agonist is TrkB agonist polynucleotide, such as a peptide or protein encoding polynucleotide, or an aptamer.
- [0016] In additional embodiments, the TrkB agonist is a TrkB aptamer. A TrkB aptamer is a small polynucleotide which binds TrkB and promotes the ability of TrkB to increase neuronal survival.
- [0017] In various embodiments of the above methods, the LINGO-1 antagonist may be any molecule which interferes with the ability of LINGO-1 to negatively regulate neuronal survival and/or to bind to TrkB and/or to inhibit or decrease TrkB phosphorylation. In certain embodiments, the LINGO-1 antagonist is selected from the group consisting of a LINGO-1 antagonist polypeptide, a LINGO-1 antagonist antibody or fragment thereof, a LINGO-1 antagonist polynucleotide (*e.g.* RNA interference), a LINGO-1 aptamer, or a combination of two or more LINGO-1 antagonists.
- [0018] In certain embodiments, the LINGO-1 antagonist polypeptide is a soluble LINGO-1 polypeptide. Certain soluble LINGO-1 polypeptides of the invention include, but are not limited to, soluble LINGO-1 polypeptides which comprise or lack one or more of the following domains: (i) a LINGO-1 Leucine-Rich Repeat (LRR) domain, (ii) a LINGO-1 basic region C-terminal to the LRR domain, and (iii) a LINGO-1 immunoglobulin (Ig) domain. In some embodiments, the soluble LINGO-1 polypeptide lacks one or more of a LINGO-1 Ig domain, a LINGO-1 LRR domain, a transmembrane domain, and a cytoplasmic domain. Additional LINGO-1 soluble polypeptides of the invention include polypeptides which lack a transmembrane domain and a cytoplasmic domain. In some embodiments, the soluble LINGO-1 polypeptide comprises a LINGO-1 LRR domain and lacks a LINGO-1 Ig domain, a LINGO-1 basic region, a transmembrane domain, and a cytoplasmic domain. In some embodiments, the soluble LINGO-1 polypeptide comprises amino acid residues 34-532 of SEQ ID NO: 2 or 36-532 of SEQ ID NO:2.
- [0019] In some embodiments, the LINGO-1 antagonist is a fusion polypeptide comprising a non-LINGO-1-antagonist heterologous polypeptide. In some embodiments, the non-LINGO-1-antagonist polypeptide is selected from the group consisting of an antibody Ig polypeptide, a serum albumin polypeptide, a targeting polypeptide, a reporter polypeptide, and a purification-facilitating polypeptide. In some embodiments, the antibody Ig polypeptide is a hinge and Fc polypeptide.

[0020] In alternative embodiments, the LINGO-1 antagonist is an antibody or fragment thereof which binds to a LINGO-1 polypeptide comprising one or more of the following LINGO-1 domains: (i) a LINGO-1 Leucine-Rich Repeat (LRR) domain, (ii) a LINGO-1 basic region C-terminal to the LRR domain, and (iii) a LINGO-1 immunoglobulin (Ig) domain. Additionally, the LINGO-1 antagonist antibody or fragment thereof specifically binds to an epitope within a polypeptide comprising a LINGO-1 polypeptide fragment as described herein. In additional embodiments, the LINGO-1 antagonist antibody or fragment thereof is selected from the group consisting of 201', 3A3, 3A6, 3B5, 1A7, 1D5, 1G7, 2B10, 2C11, 2F3, 3P1B1.1F9, 3P1D10.2C3, 3P1E11.3B7, 3P2C6.3G10.2H7, 3P2C9.2G4, 3P4A6.1D9, 3P4A1.2B9, 3P4C2.2D2, 3P4C5.1D8, 3P4C8.2G9, 6P4F4.1Ds, 6P4F4.1F9, 7P1D5.1G9, 1B6.4, 2C7.2, 2D6.1, 2F7.3, 2H3.2, 3C11.1, 3E3.1, 3H11.2, 3G8.1, 2B8.1, 3B5.230-C12 (Li01), 38-D01 (Li02), 35-E04 (Li03), 36-C09 (Li04), 30-A11 (Li05), 34-F02 (Li06), 29-E07 (Li07), 34-G04 (Li08), 36-A12 (Li09), 28-D02 (Li10), 30-B01 (Li11), 34-B03 (Li12), Li13, Li32, Li33, Li34, 3383 (L1a.1), 3495(L1a.2), 3563 (L1a.3), 3564 (L1a.4), 3565 (L1a.5), 3566 (L1a.6), 3567 (L1a.7), 3568 (L1a.8), 3569 (L1a.9), 3570 (L1a.10), 3571 (L1a.11), 3582 (L1a.12), 1968 (L1a.13), 3011, 3012, 3013, 3418, 3422, 3562, D05, D07, D08, D10 and D11. The LINGO-1 antagonist can also be an antigen-binding fragment of any one of these antibodies or a combination of two or more antibodies or fragments thereof.

[0021] In other embodiments, the LINGO-1 antagonist is a LINGO-1 antagonist polynucleotide such as an antisense polynucleotide, an aptamer, a ribozyme, a small interfering RNA (siRNA), or a small-hairpin RNA (shRNA).

[0022] In additional embodiments, the LINGO-1 antagonist is a LINGO-1 aptamer. A LINGO-1 aptamer is a small polypeptide or a polynucleotide which binds LINGO-1 and interferes with LINGO-1 and TrkB interaction and/or promotes or increases TrkB phosphorylation.

[0023] In some embodiments of the above methods, the TrkB agonist and/or LINGO-1 antagonist is administered by a method comprising (a) introducing into CNS neurons a polynucleotide which encodes the TrkB agonist and/or the LINGO-1 antagonist through operable association with an expression control sequence; and (b) allowing expression of said TrkB agonist and/or LINGO-1 antagonist. In some embodiments the CNS neurons are in a mammal and said introducing comprises (a) administering to said mammal a polynucleotide which encodes a TrkB agonist and/or a LINGO-1 antagonist through operable association with an expression control sequence. In some embodiments, the

cultured host cell is derived from the mammal to be treated. In certain embodiments, the polynucleotide is introduced into the host cell or CNS neuron via transfection, electroporation, viral transduction or direct microinjection.

[0024] In certain embodiments the TrkB agonist and/or the LINGO-1 antagonist is a polynucleotide that can be administered to a mammal, at or near the site of the disease, disorder or injury. In some embodiments, the polynucleotide is administered as an expression vector. In certain embodiments, the vector is a viral vector which is selected from the group consisting of an adenoviral vector, an alphavirus vector, an enterovirus vector, a pestivirus vector, a lentiviral vector, a baculoviral vector, a herpesvirus vector (*e.g.* an Epstein Barr viral vector, or a herpes simplex viral vector) a papovaviral vector, a poxvirus vector (*e.g.* a vaccinia viral vector) and a parvovirus. In some embodiments, the vector is administered by a route selected from the group consisting of topical administration, intraocular administration, and parenteral administration (*e.g.* intravenous, intraarterial, intramuscular, intracardiac, subcutaneous, intradermal, intrathecal, intraperitoneal).

BRIEF DESCRIPTION OF THE DRAWINGS/FIGURES

- [0025] Figures 1A and B – Schematic summarizing the 2-week (FIG 1A) and 4-week (FIG 1B) experimental ocular hypertension models used in the examples.
- [0026] Figure 2 – Schematic showing sections used to quantitate retinal ganglion cells (RGCs) in flat-mounted retinas. RGCs were quantified under an eyepiece grid of 200 x 200 μm^2 along the median line of each quadrant, starting from the optic disc to the border at 500 μm intervals.
- [0027] Figure 3 – Images of aqueous veins prior to (top) and immediately after (bottom) laser photocoagulation.
- [0028] Figures 4A and B – LINGO-1 expression in normal and injured rat retina sections.
- [0029] Figure 5 – Western blot of LINGO-1 in normal and injured rat retina and quantitation of the same.
- [0030] Figure 6 – Measurements of intraocular pressure in left (normal) and right (injured) eyes treated with PBS, control protein, anti-LINGO-1 antibody (1A7) or soluble LINGO-1 (LINGO-1-Fc).
- [0031] Figures 7A and B – Quantitation of the number (FIG 7A) and density (FIG 7B) of surviving RGCs two weeks and four weeks after injury and treated with PBS, control protein, 1A7 or LINGO-1-Fc.

- [0032] Figures 8A and B – Images of microstructures in RGCs (FIG 8A) and cells of the inner plexiform layer (FIG 8B) in normal animals and injured animals treated with PBS, LINGO-1-Fc or 1A7.
- [0033] Figure 9 – Quantitation of surviving RGCs grown in vitro and exposed to control protein, LINGO-1-Fc, BDNF and control protein, or BDNF and LINGO-1-Fc.
- [0034] Figure 10 – Western blot of BDNF in normal retina, in injured retina with no treatment and in injured retina treated with PBS, LINGO-1-Fc or 1A7 and quantitation of the same.
- [0035] Figures 11A and B – Quantitation of the number (FIG 11A) and density (FIG 11B) of RGCs in injured eyes treated with PBS, anti-BDNF antibody, 1A7, 1A7 and anti-BDNF antibody, LINGO-1-Fc or LINGO-1-Fc and anti-BDNF antibody.
- [0036] Figures 12A and B – Western blots of TrkB and LINGO-1 in anti-LINGO-1 immunoprecipitates from 293 cells co-expressing TrkB and LINGO-1 (FIG 12A). Western blots of phospho-TrkB, total TrkB and LINGO-1 in anti-TrkB immunoprecipitates from cells with and without LINGO-1 expression and with and without BDNF stimulation and quantitation of the same (FIG 12B).
- [0037] Figures 13A and B - Western blot of LINGO-1 in anti-TrkB, anti-LINGO-1 and anti-control protein immunoprecipitates from normal and injured retinal lysates (FIG 13A). LINGO-1 and phospho-TrkB immunostaining in retinal sections (FIG 13B).
- [0038] Figures 14A and B – Western blot of phospho-TrkB and total TrkB in normal eyes and injured eyes treated with PBS, LINGO-1-Fc or 1A7 and quantitation of the same (FIG 14A). Western blot of phospho-TrkB and total TrkB in normal eyes and injured eyes treated with BDNF, BDNF and LINGO-1-Fc or BDNF and 1A7 (FIG 14B).
- [0039] Figure 15 – Western blot of phospho-Akt and total Akt in normal and injured eyes with or without LINGO-1-Fc treatment and quantitation of the same.
- [0040] Figure 16 – pAkt immunostaining in retinal sections.
- [0041] Figures 17A and B – Quantitation of the number (FIG. 17A) and density (FIG. 17B) of RGCs in injured eyes treated with PBS, LY294002 (an inhibitor of the PI3K/Akt pathway), LINGO-1-Fc, or LY294002 and LINGO-1-Fc.
- [0042] Figure 18 – Measurement of intraocular pressure in left (normal) and right (injured) eyes in animals treated with LINGO-1-Fc and LY294002 or LY294002 alone.
- [0043] Figure 19 – Western blot of phospho-JNK-2, phospho-JNK-1, total JNK-2 and total JNK-1 in normal and injured eyes treated with PBS, LINGO-1-Fc and 1A7 and quantitation of the same.

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- [0044] Figure 20 - A schematic showing the proposed molecular mechanism. Elevations in intraocular pressure result in increased levels of both LINGO-1 and BDNF. BDNF promotes phosphorylation of TrkB and activation of a cell survival signaling pathway, but the LINGO-1 inhibits this activity. LINGO-1 antagonists, such as a 1A7 or LINGO-1-Fc, promote cell survival by interfering with the ability of LINGO-1 to prevent TrkB phosphorylation in response to BDNF signaling.
- [0045] Figure 21 - Western blot of GTP-RhoA and total RhoA in normal and injured eyes treated with PBS or LINGO-1-Fc and quantitation of the same.

DETAILED DESCRIPTION OF THE INVENTION

Definitions

- [0046] Unless defined otherwise, 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 invention belongs. In case of conflict, the present application including the definitions will control. Unless otherwise required by context, singular terms shall include pluralities and plural terms shall include the singular. All publications, patents and other references mentioned herein are incorporated by reference in their entireties for all purposes as if each individual publication or patent application were specifically and individually indicated to be incorporated by reference.
- [0047] Although methods and materials similar or equivalent to those described herein can be used in practice or testing of the present invention, suitable methods and materials are described below. The materials, methods and examples are illustrative only and are not intended to be limiting. Other features and advantages of the invention will be apparent from the detailed description and from the claims.
- [0048] In order to further define this invention, the following terms and definitions are provided.
- [0049] It is to be noted that the term “a” or “an” entity, refers to one or more of that entity; for example, “an immunoglobulin molecule,” is understood to represent one or more immunoglobulin molecules. As such, the terms “a” (or “an”), “one or more,” and “at least one” can be used interchangeably herein.
- [0050] Throughout this specification and claims, the word “comprise,” or variations such as “comprises” or “comprising,” indicate the inclusion of any recited integer or group of

integers but not the exclusion of any other integer or group of integers. The term "comprising" is inclusive or open-ended and does not exclude additional, unrecited elements or method steps. The phrase "consisting essentially of" indicates the inclusion of the specified materials or steps as well as those which do not materially affect the basic and novel characteristics of the claimed invention. As used herein, the term "consisting" refers only to indicated material or method steps.

[0051] As used herein, a "therapeutically effective amount" refers to an amount effective, at dosages and for periods of time necessary, to achieve a desired therapeutic result. A therapeutic result may be, *e.g.*, lessening of symptoms, prolonged survival, improved mobility, and the like. A therapeutic result need not be a "cure".

[0052] As used herein, the term "treatment" or "treating" refers to the administration of an agent to an animal in order to ameliorate or lessen the symptoms of a disease. Additionally, the terms "treatment" or "treating" refers to the administration of an agent to an animal to prevent the progression of a disease.

[0053] As used herein, a "prophylactically effective amount" refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired prophylactic result. Typically, since a prophylactic dose is used in subjects prior to or at an earlier stage of disease, the prophylactically effective amount will be less than the therapeutically effective amount.

[0054] As used herein, a "polynucleotide" can contain a nucleic acid sequence of a full length cDNA sequence and may include untranslated 5' and 3' sequences, the coding region or fragments or and variants of the nucleic acid sequence. The polynucleotide can be composed of any polyribonucleotide or polydeoxyribonucleotide, which may be unmodified RNA or DNA or modified RNA or DNA. For example, polynucleotides can be composed of single- and double-stranded DNA, DNA that is a mixture of single- and double-stranded regions, single- and double-stranded RNA, and RNA that is mixture of single- and double-stranded regions, hybrid molecules comprising DNA and RNA that may be single-stranded or, more typically, double-stranded or a mixture of single- and double-stranded regions. In addition, the polynucleotides can be composed of triple-stranded regions comprising RNA or DNA or both RNA and DNA. Polynucleotides may also contain one or more modified bases or DNA or RNA backbones modified for stability or for other reasons. "Modified" bases include, for example, tritylated bases and unusual bases such as inosine. A variety of modifications can be made to DNA and

RNA; thus, "polynucleotide" embraces chemically, enzymatically, or metabolically modified forms.

[0055] In the present invention, a "polypeptide" can be composed of amino acids joined to each other by peptide bonds or modified peptide bonds, i.e., peptide isosteres, and may contain amino acids other than the 20 gene-encoded amino acids (e.g. non-naturally occurring amino acids). As used to describe the present invention, the terms "peptide" and "polypeptide" may be used interchangeably. The polypeptides of the present invention may be modified by either natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well as in a voluminous research literature. Modifications can occur anywhere in the polypeptide, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given polypeptide. Also, a given polypeptide may contain many types of modifications. Polypeptides may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic polypeptides may result from posttranslational natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphatidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, *Proteins - Structure And Molecular Properties*, 2nd Ed., T.E. Creighton, W.H. Freeman and Company, New York (1993); *Posttranslational Covalent Modification of Proteins*, B.C. Johnson, Ed., Academic Press, New York, pgs. 1-12 (1983); Seifter *et al.*, *Meth Enzymol* 182:626-646 (1990); Rattan *et al.*, *Ann NY Acad Sci* 663:48-62 (1992).)

[0056] The terms "fragment," "variant," "derivative" and "analog" when referring to a LINGO-1 antagonist or a TrkB agonist of the present invention include any antagonist or

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agonist molecules which retain at least some ability to inhibit LINGO-1 activity, or to promote TrkB activity. LINGO-1 antagonists and TrkB agonists as described herein may include fragment, variant, or derivative molecules therein without limitation, so long as the LINGO-1 antagonist or TrkB agonist still serves its function. LINGO-1 antagonist or TrkB agonist polypeptides of the present invention may include LINGO-1 or TrkB-agonist proteolytic fragments, deletion fragments and in particular, fragments which more easily reach the site of action when delivered to an animal. Polypeptide fragments further include any portion of the polypeptide which comprises an antigenic or immunogenic epitope of the native polypeptide, including linear as well as three-dimensional epitopes. LINGO-1 or TrkB-agonist polypeptides of the present invention may comprise variant LINGO-1 or TrkB-agonist regions, including fragments as described above, and also polypeptides with altered amino acid sequences due to amino acid substitutions, deletions, or insertions. Variants may occur naturally, such as an allelic variant. By an "allelic variant" is intended alternate forms of a gene occupying a given locus on a chromosome of an organism. Genes II, Lewin, B., ed., John Wiley & Sons, New York (1985). Non-naturally occurring variants may be produced using art-known mutagenesis techniques. LINGO-1 or TrkB-agonist polypeptides may comprise conservative or non-conservative amino acid substitutions, deletions or additions. LINGO-1 antagonists or TrkB agonists of the present invention may also include derivative molecules. For example, LINGO-1 or TrkB-agonist polypeptides of the present invention may include LINGO-1 or TrkB-agonist regions which have been altered so as to exhibit additional features not found on the native polypeptide. Examples include fusion proteins and protein conjugates.

[0057] In the present invention, a "polypeptide fragment" refers to a short amino acid sequence of a LINGO-1 or TrkB-agonist polypeptide. Protein fragments may be "free-standing," or comprised within a larger polypeptide of which the fragment forms a part or region. Representative examples of polypeptide fragments of the invention include, for example, fragments comprising about 5 amino acids, about 10 amino acids, about 15 amino acids, about 20 amino acids, about 30 amino acids, about 40 amino acids, about 50 amino acids, about 60 amino acids, about 70 amino acids, about 80 amino acids, about 90 amino acids, and about 100 amino acids or more in length.

[0058] In certain embodiments, LINGO-1 antagonists or TrkB agonists for use in the methods disclosed herein are "antibody" or "immunoglobulin" molecules, or immunospecific fragments thereof, *e.g.*, naturally occurring antibody or immunoglobulin

molecules or engineered antibody molecules or fragments that bind antigen in a manner similar to antibody molecules. The terms "antibody" and "immunoglobulin" are used interchangeably herein. Additionally, immunoglobulin molecules used in the methods of the invention are also described as "immunospecific" or "antigen-specific" or "antigen-binding" molecules and are used interchangeably to refer to antibody molecules and fragments thereof. An antibody or immunoglobulin comprises at least the variable domain of a heavy chain, and normally comprises at least the variable domains of a heavy chain and a light chain. Basic immunoglobulin structures in vertebrate systems are relatively well understood. See, e.g., Harlow *et al.*, *Antibodies: A Laboratory Manual*, (Cold Spring Harbor Laboratory Press, 2nd ed. 1988), incorporated herein by reference.

[0059] As will be discussed in more detail below, the term "immunoglobulin" comprises five broad classes of polypeptides that can be distinguished biochemically. All five classes are clearly within the scope of the present invention, the following discussion will generally be directed to the IgG class of immunoglobulin molecules. With regard to IgG, a standard immunoglobulin molecule comprises two identical light chain polypeptides of molecular weight approximately 23,000 Daltons, and two identical heavy chain polypeptides of molecular weight 53,000-70,000. The four chains are typically joined by disulfide bonds in a "Y" configuration wherein the light chains bracket the heavy chains starting at the mouth of the "Y" and continuing through the variable region.

[0060] Both the light and heavy chains are divided into regions of structural and functional homology. The terms "constant" and "variable" are used functionally. In this regard, it will be appreciated that the variable domains of both the light (V_L) and heavy (V_H) chain portions determine antigen recognition and specificity. Conversely, the constant domains of the light chain (C_L) and the heavy chain (C_{H1} , C_{H2} or C_{H3}) confer important biological properties such as secretion, transplacental mobility, Fc receptor binding, complement binding, and the like. By convention the numbering of the constant region domains increases as they become more distal from the antigen binding site or amino-terminus of the antibody. The N-terminal portion is a variable region and at the C-terminal portion is a constant region; the C_{H3} and C_L domains actually comprise the carboxy-terminus of the heavy and light chain, respectively.

[0061] Light chains are classified as either kappa or lambda (κ , λ). Each heavy chain class may be bound with either a kappa or lambda light chain. In general, the light and heavy chains are covalently bonded to each other, and the "tail" portions of the two heavy chains are bonded to each other by covalent disulfide linkages or non-covalent linkages

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when the immunoglobulins are generated either by hybridomas, B cells or genetically engineered host cells. In the heavy chain, the amino acid sequences run from an N-terminus at the forked ends of the Y configuration to the C-terminus at the bottom of each chain. Those skilled in the art will appreciate that heavy chains are classified as gamma, mu, alpha, delta, or epsilon, (γ , μ , α , δ , ϵ) with some subclasses among them (*e.g.*, $\gamma 1$ - $\gamma 4$). It is the nature of this chain that determines the "class" of the antibody as IgG, IgM, IgA, IgG, or IgE, respectively. The immunoglobulin subclasses (isotypes) *e.g.*, IgG₁, IgG₂, IgG₃, IgG₄, IgA₁, etc. are well characterized and are known to confer functional specialization. Modified versions of each of these classes and isotypes are readily discernable to the skilled artisan in view of the instant disclosure and, accordingly, are within the scope of the instant invention.

[0062] As indicated above, the variable region allows the antibody to selectively recognize and specifically bind epitopes on antigens. That is, the V_L domain and V_H domain of an antibody combine to form the variable region that defines a three dimensional antigen binding site. This quaternary antibody structure forms the antigen binding site present at the end of each arm of the Y. More specifically, the antigen binding site is defined by three complementary determining regions (CDRs) on each of the V_H and V_L chains. In some instances, *e.g.*, certain immunoglobulin molecules derived from camelid species or engineered based on camelid immunoglobulins, a complete immunoglobulin molecule may consist of heavy chains only, with no light chains. See, *e.g.*, Hamers-Casterman *et al.*, *Nature* 363:446-448 (1993).

[0063] In naturally occurring antibodies, the six "complementarity determining regions" or "CDRs" present in each antigen binding domain are short, non-contiguous sequences of amino acids that are specifically positioned to form the antigen binding domain as the antibody assumes its three dimensional configuration in an aqueous environment. The remainder of the amino acids in the antigen binding domains, referred to as "framework" regions, show less inter-molecular variability. The framework regions largely adopt a β -sheet conformation and the CDRs form loops which connect, and in some cases form part of, the β -sheet structure. Thus, framework regions act to form a scaffold that provides for positioning the CDRs in correct orientation by inter-chain, non-covalent interactions. The antigen binding domain formed by the positioned CDRs defines a surface complementary to the epitope on the immunoreactive antigen. This complementary surface promotes the non-covalent binding of the antibody to its cognate epitope. The amino acids comprising the CDRs and the framework regions, respectively, can be readily identified for any given

heavy or light chain variable region by one of ordinary skill in the art, since they have been precisely defined (see, "Sequences of Proteins of Immunological Interest," Kabat, E., *et al.*, U.S. Department of Health and Human Services, (1983); and Chothia and Lesk, *J. Mol. Biol.*, 196:901-917 (1987), which are incorporated herein by reference in their entireties).

[0064] In camelid species, however, the heavy chain variable region, referred to as V_HH, forms the entire CDR. The main differences between camelid V_HH variable regions and those derived from conventional antibodies (V_H) include (a) more hydrophobic amino acids in the light chain contact surface of V_H as compared to the corresponding region in V_HH, (b) a longer CDR3 in V_HH, and (c) the frequent occurrence of a disulfide bond between CDR1 and CDR3 in V_HH.

[0065] In one embodiment, an antigen binding molecule for use in the methods of the invention comprises at least one heavy or light chain CDR of an antibody molecule. In another embodiment, an antigen binding molecule for use in the methods of the invention comprises at least two CDRs from one or more antibody molecules. In another embodiment, an antigen binding molecule for use in the methods of the invention comprises at least three CDRs from one or more antibody molecules. In another embodiment, an antigen binding molecule for use in the methods of the invention comprises at least four CDRs from one or more antibody molecules. In another embodiment, an antigen binding molecule for use in the methods of the invention comprises at least five CDRs from one or more antibody molecules. In another embodiment, an antigen binding molecule for use in the methods of the invention comprises at least six CDRs from one or more antibody molecules. Exemplary antibody molecules comprising at least one CDR that can be included in the subject antigen binding molecules are known in the art and exemplary molecules are described herein.

[0066] Antibodies or immunospecific fragments thereof for use in the methods of the invention include, but are not limited to, polyclonal, monoclonal, multispecific, human, humanized, primatized, or chimeric antibodies, single chain antibodies, epitope-binding fragments, *e.g.*, Fab, Fab' and F(ab')₂, Fd, Fvs, single-chain Fvs (scFv), single-chain antibodies, disulfide-linked Fvs (sdFv), fragments comprising either a VL or VH domain, fragments produced by a Fab expression library, and anti-idiotypic (anti-Id) antibodies (including, *e.g.*, anti-Id antibodies to binding molecules disclosed herein). ScFv molecules are known in the art and are described, *e.g.*, in US patent 5,892,019. Immunoglobulin or antibody molecules of the invention can be of any type (*e.g.*, IgG,

IgE, IgM, IgD, IgA, and IgY), class (e.g., IgG₁, IgG₂, IgG₃, IgG₄, IgA₁ and IgA₂) or subclass of immunoglobulin molecule.

[0067] Antibody fragments, including single-chain antibodies, may comprise the variable region(s) alone or in combination with the entirety or a portion of the following: hinge region, C_H1, C_H2, and C_H3 domains of the heavy chain, or C_L of the light chain. Also included in the invention are antigen-binding fragments also comprising any combination of variable region(s) with a hinge region, C_H1, C_H2, C_H3, or C_L domain. Antibodies or immunospecific fragments thereof for use in the methods disclosed herein may be from any animal origin including birds and mammals. The antibodies can be human, murine, donkey, rabbit, goat, guinea pig, camel, llama, horse, or chicken antibodies. In another embodiment, the variable region may be chondrichthoid in origin (e.g., from sharks). As used herein, "human" antibodies include antibodies having the amino acid sequence of a human immunoglobulin and include antibodies isolated from human immunoglobulin libraries or from animals transgenic for one or more human immunoglobulins and that do not express endogenous immunoglobulins, as described *infra* and, for example, in U.S. Pat. No. 5,939,598 by Kucherlapati *et al.* Such antibodies also include variants that contain one or more amino acid substitutions.

[0068] As used herein, the term "heavy chain portion" includes amino acid sequences derived from an immunoglobulin heavy chain. A polypeptide comprising a heavy chain portion comprises at least one of: a C_H1 domain, a hinge (e.g., upper, middle, and/or lower hinge region) domain, a C_H2 domain, a C_H3 domain, or a variant or fragment thereof. For example, a heavy chain portion may comprise a polypeptide chain comprising a C_H1 domain; a polypeptide chain comprising a C_H1 domain, at least a portion of a hinge domain, and a C_H2 domain; a polypeptide chain comprising a C_H1 domain and a C_H3 domain; a polypeptide chain comprising a C_H1 domain, at least a portion of a hinge domain, and a C_H3 domain, or a polypeptide chain comprising a C_H1 domain, at least a portion of a hinge domain, a C_H2 domain, and a C_H3 domain. The heavy chain portion may also include a polypeptide comprising a polypeptide chain comprising a C_H3 domain. Further, a binding polypeptide for use in the invention may lack at least a portion of a C_H2 domain (e.g., all or part of a C_H2 domain). As set forth above, it will be understood by one of ordinary skill in the art that these domains (e.g., the heavy chain portions) may be modified such that they vary in amino acid sequence from the naturally occurring immunoglobulin molecule.

[0069] In certain LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods disclosed herein, the heavy chain portions of one polypeptide chain of a multimer are identical to those on a second polypeptide chain of the multimer. Alternatively, heavy chain portion-containing monomers for use in the methods of the invention are not identical. For example, each monomer may comprise a different target binding site, forming, for example, a bispecific antibody.

[0070] The heavy chain portions of a binding polypeptide for use in the methods disclosed herein may be derived from different immunoglobulin molecules. For example, a heavy chain portion of a polypeptide may comprise a C_H1 domain derived from an IgG₁ molecule and a hinge region derived from an IgG₃ molecule. In another example, a heavy chain portion can comprise a hinge region derived, in part, from an IgG₁ molecule and, in part, from an IgG₃ molecule. In another example, a heavy chain portion can comprise a chimeric hinge derived, in part, from an IgG₁ molecule and, in part, from an IgG₄ molecule.

[0071] As used herein, the term "light chain portion" includes amino acid sequences derived from an immunoglobulin light chain. The light chain portion can comprise at least one of a V_L or C_L domain.

[0072] An isolated nucleic acid molecule encoding a non-natural variant of a polypeptide derived from an immunoglobulin (*e.g.*, an immunoglobulin heavy chain portion or light chain portion) can be created by introducing one or more nucleotide substitutions, additions or deletions into the nucleotide sequence of the immunoglobulin such that one or more amino acid substitutions, additions or deletions are introduced into the encoded protein. Mutations may be introduced by standard techniques, such as site-directed mutagenesis and PCR-mediated mutagenesis. Conservative amino acid substitutions can be made at one or more non-essential amino acid residues.

[0073] Antibodies or immunospecific fragments thereof for use in the methods disclosed herein may also be described or specified in terms of their binding affinity to a polypeptide of the invention. In some embodiments, binding affinities are those with a dissociation constant or K_d less than 5 x 10⁻² M, 10⁻² M, 5 x 10⁻³ M, 10⁻³ M, 5 x 10⁻⁴ M, 10⁻⁴ M, 5 x 10⁻⁵ M, 10⁻⁵ M, 5 x 10⁻⁶ M, 10⁻⁶ M, 5 x 10⁻⁷ M, 10⁻⁷ M, 5 x 10⁻⁸ M, 10⁻⁸ M, 5 x 10⁻⁹ M, 10⁻⁹ M, 5 x 10⁻¹⁰ M, 10⁻¹⁰ M, 5 x 10⁻¹¹ M, 10⁻¹¹ M, 5 x 10⁻¹² M, 10⁻¹² M, 5 x 10⁻¹³ M, 10⁻¹³ M, 5 x 10⁻¹⁴ M, 10⁻¹⁴ M, 5 x 10⁻¹⁵ M, or 10⁻¹⁵ M.

[0074] Antibodies or immunospecific fragments thereof for use in the methods disclosed herein act as antagonists of LINGO-1 or agonists of TrkB as described herein. For

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example, an antibody for use in the methods of the present invention may function as an antagonist, blocking or inhibiting the suppressive activity of the LINGO-1 polypeptide or as an agonist promoting the activity of TrkB.

[0075] As used herein, the term "chimeric antibody" will be held to mean any antibody wherein the immunoreactive region or site is obtained or derived from a first species and the constant region (which may be intact, partial or modified in accordance with the instant invention) is obtained from a second species. In certain embodiments the target binding region or site will be from a non-human source (*e.g.* mouse or primate) and the constant region is human.

[0076] As used herein, the term "engineered antibody" refers to an antibody in which the variable domain in either the heavy or light chain or both is altered by at least partial replacement of one or more CDRs from an antibody of known specificity and, if necessary, by partial framework region replacement and sequence changing. Although the CDRs may be derived from an antibody of the same class or even subclass as the antibody from which the framework regions are derived, it is envisaged that the CDRs will be derived from an antibody of different class or from an antibody from a different species. An engineered antibody in which one or more "donor" CDRs from a non-human antibody of known specificity is grafted into a human heavy or light chain framework region is referred to herein as a "humanized antibody." It may not be necessary to replace all of the CDRs with the complete CDRs from the donor variable region to transfer the antigen binding capacity of one variable domain to another. Rather, it may only be necessary to transfer those residues that are necessary to maintain the activity of the target binding site. Given the explanations set forth in, *e.g.*, U. S. Pat. Nos. 5,585,089, 5,693,761, 5,693,762, and 6,180,370, it will be well within the competence of those skilled in the art, either by carrying out routine experimentation or by trial and error testing to obtain a functional engineered or humanized antibody.

[0077] As used herein, the terms "linked," "fused" or "fusion" are used interchangeably. These terms refer to the joining together of two more elements or components, by whatever means including chemical conjugation or recombinant means. An "in-frame fusion" refers to the joining of two or more open reading frames (ORFs) to form a continuous longer ORF, in a manner that maintains the correct reading frame of the original ORFs. Thus, the resulting recombinant fusion protein is a single protein containing two or more segments that correspond to polypeptides encoded by the original ORFs (which segments are not normally so joined in nature.) Although the

reading frame is thus made continuous throughout the fused segments, the segments may be physically or spatially separated by, for example, in-frame linker sequence.

[0078] In the context of polypeptides, a "linear sequence" or a "sequence" is an order of amino acids in a polypeptide in an amino to carboxyl terminal direction in which residues that neighbor each other in the sequence are contiguous in the primary structure of the polypeptide.

[0079] The term "expression" as used herein refers to a process by which a DNA sequence is used for the production a biochemical, for example, an RNA or polypeptide. The process includes any manifestation of the functional presence of the gene within the cell including, without limitation, gene knockdown as well as both transient expression and stable expression. It includes without limitation transcription of the gene into messenger RNA (mRNA), transfer RNA (tRNA), small hairpin RNA (shRNA), small interfering RNA (siRNA) or any other RNA product and the translation of such mRNA into polypeptide(s). If the final desired product is biochemical, expression includes the creation of that biochemical and any precursors.

[0080] By "subject" or "individual" or "animal" or "patient" or "mammal," is meant any subject, particularly a mammalian subject, for whom diagnosis, prognosis, or therapy is desired. Mammalian subjects include, but are not limited to, humans, domestic animals, farm animals, zoo animals, sport animals, pet animals such as dogs, cats, guinea pigs, hamsters, rabbits, rats, mice; primates such as apes, monkeys, orangutans, and chimpanzees; canids such as foxes and wolves; felids such as lions, and tigers; equids such as horses, donkeys, and zebras; food animals such as cows, pigs, and sheep; ungulates such as deer and giraffes; bears; and so on. In certain embodiments, the mammal is a human subject.

[0081] The term "RNA interference" or "RNAi" refers to the silencing or decreasing of gene expression by siRNAs. It is the process of sequence-specific, post-transcriptional gene silencing in animals and plants, initiated by siRNA that is homologous in its duplex region to the sequence of the silenced gene. The gene may be endogenous or exogenous to the organism, present integrated into a chromosome or present in a transfection vector that is not integrated into the genome. The expression of the gene is either completely or partially inhibited. RNAi may also be considered to inhibit the function of a target RNA; the function of the target RNA may be complete or partial.

LINGO-1 (Sp35/LRRN6)

[0082] Naturally occurring human LINGO-1 is a glycosylated nervous-system-specific protein consisting of 614 amino acids (SEQ ID NO: 2). The human LINGO-1 polypeptide contains an LRR domain consisting of 14 leucine-rich repeats (including N- and C-terminal caps), an Ig domain, a transmembrane region, and a cytoplasmic domain. The cytoplasmic domain contains a canonical tyrosine phosphorylation site. In addition, the naturally occurring LINGO-1 protein contains a signal sequence, a short basic region between the LRRCT and Ig domain, and a transmembrane region between the Ig domain and the cytoplasmic domain. The human LINGO-1 gene contains alternative translation start codons, so that six additional amino acids (MQVSKR; SEQ ID NO:7) may or may not be present at the N-terminus of the LINGO-1 signal sequence. Table 1 lists the LINGO-1 domains and other regions, according to amino acid residue number, based on the sequence of SEQ ID NO:2. As one of skill in the art will appreciate, the beginning and ending residues of the domains listed below may vary depending upon the computer modeling program used or the method used for determining the domain. The LINGO-1 polypeptide is characterized in more detail in PCT Publication No. WO 2004/085648 and U.S. Published Application No. 2006/0009388 A1, which are incorporated herein by reference in their entireties.

Table 1:

Domain or Region	Beginning Residue	Ending Residue
Signal Sequence	1	33 or 35
LRRNT	34 or 36	64
LRR	66	89
LRR	90	113
LRR	114	137
LRR	138	161
LRR	162	185
LRR	186	209
LRR	210	233
LRR	234	257
LRR	258	281
LRR	282	305
LRR	306	329
LRR	330	353
LRRCT	363	414 or 416
Basic	415 or 417	424
Ig	419	493
Connecting sequence	494	551
Transmembrane	552	576
Cytoplasmic	577	614

[0083] Tissue distribution and developmental expression of LINGO-1 have been studied in humans and rats. LINGO-1 biology has been studied in an experimental animal (rat) model. Expression of rat LINGO-1 is localized to nervous-system neurons and brain oligodendrocytes, as determined by northern blot and immuno-histochemical staining. Rat LINGO-1 mRNA expression level is regulated developmentally, peaking shortly after birth, *i.e.*, ca. postnatal day one. In a rat spinal cord transection injury model, LINGO-1 is up-regulated at the injury site, as determined by RT-PCR. In addition, LINGO-1 has been shown to interact with Nogo66 Receptor (Nogo receptor). See, *e.g.*, International Patent Application No. PCT/US2004/00832, PCT Publication No. WO2004/08564.

[0084] LINGO-1 is an additional component of the Nogo Receptor-1-p75-Taj neurotrophin receptor complex. See Mi *et al.*, *Nat Neurosci.* 7:221-228 (2004), which is

incorporated herein by reference. Unlike Nogo receptor 1, LINGO-1 gene expression is increased when adult nerve cells in the spinal cord are exposed to traumatic injuries, suggesting that LINGO-1 has an important biological role for CNS neurological function. Id.

[0085] The nucleotide sequence for the full-length human LINGO-1 molecule is as follows:

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ATGCTGGCGGGGGGCGTGAGGAGCATGCCAGCCCCCTCCTGGCCTGCTGGCAGCCCAT
CCTCCTGCTGGTGCTGGGCTCAGTGCTGTGAGGCTCGGCCACGGGCTGCCCCCCCCGCT
GCGAGTGCTCCGCCCAGGACCGCGCTGTGCTGTGCCACCGCAAGCGCTTTGTGGCAGTC
CCCGAGGGCATCCCCACCGAGACGCGCTGTGACCTAGGCAAGAACCGCATCAAAAC
GCTCAACCAGGACGAGTTCGCCAGCTTCCCGCACCTGGAGGAGCTGGAGCTCAACGAGA
ACATCGTGAGCGCCGTGGAGCCCCGGCGCCTTCAACAACCTCTTCAACCTCCGGACGCTG
GGTCTCCGCAGCAACCGCCTGAAGCTCATCCCGCTAGGCGTCTTCACTGGCCTCAGCAA
CCTGACCAAGCTGGACATCAGCGAGAAACAAGATTGTTATCCTGCTGGACTACATGTTTC
AGGACCTGTACAACCTCAAGTCACTGGAGGTTGGCGACAATGACCTCGTCTACATCTCT
CACCGCGCCTTCAGCGGCCTCAACAGCCTGGAGCAGCTGACGCTGGAGAAATGCAACCT
GACCTCCATCCCCACCGAGGCGCTGTCCACCTGCACGGCCTCATCGTCCTGAGGCTCC
GGCACCTCAACATCAATGCCATCCGGGACTACTCCTTCAAGAGGCTCTACCGACTCAAG
GTCTTGAGATCTCCCACTGGCCCTACTTGGACACCATGACACCCAACCTGCCTCTACGG
CCTCAACCTGACGTCCCTGTCCATCACACACTGCAATCTGACCGCTGTGCCCTACCTGG
CCGTCCGCCACCTAGTCTATCTCCGCTTCCCTCAACCTCTCCTACAACCCCATCAGCACC
ATTGAGGGCTCCATGTTGCATGAGCTGCTCCGGCTGCAGGAGATCCAGCTGGTGGGCGG
GCAGCTGGCCGTGGTGGAGCCCTATGCCTTCCGCGGCCTCAACTACCTGCGCGTGCTCA
ATGTCTCTGGCAACCAGCTGACCACACTGGAGGAATCAGTCTTCCACTCGGTGGGCAAC
CTGGAGACACTCATCCTGGACTCCAACCCGCTGGCCTGCGACTGTCGGCTCCTGTGGGT
GTTCCGGCGCCGCTGGCGGCTCAACTTCAACCGGCAGCAGCCACGTGCGCCACGCCCG
AGTTTGTCCAGGGCAAGGAGTTCAAGGACTTCCCTGATGTGCTACTGCCCAACTACTTC
ACCTGCCGCCGCGCCCGCATCCGGGACCGCAAGGCCAGCAGGTGTTTGTGGACGAGGG
CCACACGGTGACGTTTGTGTGCCGGGCCGATGGCGACCCGCCGCCCATCCTCTGGC
TCTCACCCCGAAAGCACCTGGTCTCAGCCAAGAGCAATGGGCGGCTCACAGTCTTCCCT
GATGGCACGCTGGAGGTGCGCTACGCCCAGGTACAGGACAACGGCACGTACCTGTGCAT
CGCGGCCAACGCGGGCGGCAACGACTCCATGCCCGCCACCTGCATGTGCGCAGCTACT
CGCCCGACTGGCCCCATCAGCCCAACAAGACCTTCGCTTTCATCTCCAACAGCCGGGC
GAGGGAGAGGCCAACAGCACCCGCGCCACTGTGCCTTTCCCCTTCGACATCAAGACCCT
CATCATCGCCACCACCATGGGCTTCATCTCTTTTCTGGGCGTCGTCCTCTTCTGCCTGG
TGCTGCTGTTTCTCTGGAGCCGGGGCAAGGGCAACACAAAGCACACATCGAGATCGAG
TATGTGCCCCGAAAGTCGGACGCAGGCATCAGCTCCGCCGACGCGCCCCGCAAGTTCAA
CATGAAGATGATATGA (SEQ ID NO:1).

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[0086] The polypeptide sequence for the full-length human LINGO-1 polypeptide is as follows:

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MLAGGVRSMPSPLLACWQPIILLVLGSLVSGSATGCPPRCECSAQDRAVLCHRKRFAV
PEGIPTETRLDLGKNRIKTLNQDEFASFPHLEELNENIVSAVEPGAFNNLNLRTL
GLRSNRLKLIPLGVFTGLSNLTKLDISENKIVILLDYMFDLYNLKSLEVGDNDLVYIS
HRAFSGLNSLEQLTLEKCNLTSIPTALSHLHGLIVLRLRLNLINAIIRDYSFKRLYRLK
VLEISHWPYLDTMTPNCLYGLNLTSLSITHCNLTAVPYLAVRHLVYLRFLNLSYNPIST
IEGSMHELLRLQEIQLVGGQLAVEPEYAFRGLNYLRVLNVSGNQLTTEESVFHSGVN
LETILDSNPLACDCRLLWVFRRLNLFNRQQPTCATPEFVQGKEFKDFPDVLLPNYF
TCRRARIRDRKAQQVFVDEGHTVQFVCRADGDPFPAILWLSPRKHLVSAKSNGRLLTVFP

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DGTLEVRYAQVQDNGTYLCIAANAGGNDSPAHHLHVRSYSPDWPHQPNKTFAFISNQPG
 EGEANSTRATVPFPFDIKTLIIATTMGFISFLGVVLFCLVLLFLWSRGKGN TKHNIEIE
 YVPRKSDAGISSADAPRKFNMKMI (SEQ ID NO:2).

TrkB (NTRK2)

[0087] The neurotrophins are a small family of highly homologous growth factors responsible for differentiation, survival and function of neurons. In mammals, the known neurotrophins are nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3), neurotrophin-4 (NT-4), also known as NT-4/5 or NT-5, neurotrophin-6 (NT-6) and neurotrophin-7 (NT-7) (Barbacid, *J. of Neurobiol.* 25:1386-1403 (1994) and Nilsson *et al. FEBS* 424:285-290 (1998)). Neurotrophins bind two receptor types, the p75 neurotrophin receptor (p75NTR) and the three members (in mammals) of the Trk receptor family of tyrosine kinases (TrkA, TrkB and TrkC). Binding of a neurotrophin to a Trk receptor extracellular domain initiates a signal transduction pathway. The binding of the neurotrophin leads to dimerization and autophosphorylation of the receptor. Autophosphorylation of TrkB leads to activation of signaling pathways including mitogen-activated protein kinase (MAPK), phospholipase C- γ (PLC- γ) and phosphatidylinositol-3 kinase (PI3-K) (Yamada, *J. Pharmacol. Sci.* 91:267-270 (2003)).

[0088] The following nucleotide sequence was reported as the mRNA for human TrkB receptor and is accession number NM_006180 in Genbank.

AAGACGGATTCTCAGACAAGGCTTGCAAATGCCCCGCAGCCATCATTTAACTGCACCCGC
 AGAATAGTTACGGTTTGTACCCGACCCTCCCGGATCGCCTAATTTGTCCCTAGTGAGAC
 CCCGAGGCTCTGCCCCGCGCTGGCTTCTTCGTAGCTGGATGCATATCGTGCTCCGGGCAG
 CGCGGGCGCAGGGCACGCGTTTCGCGCACACCCTAGCACACATGAACACGCGCAAGAGCTG
 AACCAAGCACGGTTTCCATTTCAAAAAGGGAGACAGCCTCTACCGCGATTGTAGAAGAGA
 CTGTGGTGTGAATTAGGGACCGGGAGGCGTCGAACGGAGGAACGGTTCATCTTAGAGACT
 AATTTTCTGGAGTTTCTGCCCTGCTCTGCGTCAGCCCTCACGTCACTTCGCCAGCAGTA
 GCAGAGGCGGGCGGGCGGGCTCCCGGAATTGGGTGGAGCAGGAGCCTCGCTGGCTGCTT
 CGCTCGCGCTCTACGCGCTCAGTCCCCGGCGGTAGCAGGAGCCTGGACCCAGGCGCCGCC
 GGCGGGCGTGAGGCGCCGGAGCCCGGCTCGAGGTGCATACCGGACCCCCATTTCGCATCT
 AACAAGGAATCTGCGCCCCAGAGAGTCCCGGGAGCGCCGCCGGTTCGGTGCCCGGCGCGCC
 GGGCCATGCAGCGACGGCCGCCGCGGAGCTCCGAGCAGCGGTAGCGCCCCCTGTAAAGC
 GGTTTCGCTATGCCGGGGCCACTGTGAACCCTGCCGCCTGCCGGAACACTCTTCGCTCCGG
 ACCAGCTCAGCCTCTGATAAGCTGGACTCGGCACGCCCGCAACAAGCACCGAGGAGTTAA
 GAGAGCCGCAAGCGCAGGGAAGGCCTCCCCGCACGGGTGGGGGAAAGCGGCCGGTGCAGC
 GCGGGGACAGGCACTCGGGCTGGCACTGGCTGCTAGGGATGTCGTCCTGGATAAGGTGGC
 ATGGACCCGCCATGGCGCGGCTCTGGGGCTTCTGCTGGCTGGTTGTGGGCTTCTGGAGGG
 CCGCTTTCGCCTGTCCACGTCTGCAAATGCAGTGCCTCTCGGATCTGGTGCAGCGACC
 CTTCTCCTGGCATCGTGGCATTTCGAGATTGGAGCCTAACAGTGTAGATCCTGAGAACA
 TCACCGAAATTTTCATCGCAAACCAGAAAAGGTTAGAAATCATCAACGAAGATGATGTTG
 AAGCTTATGTGGGACTGAGAAATCTGACAATTGTGGATTCTGGATTAAAATTTGTGGCTC

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ATAAAGCATTTCTGAAAAACAGCAACCTGCAGCACATCAATTTTACCCGAAACAAACTGA
CGAGTTTGTCTAGGAAACATTTCCGTACCTTGACTTGTCTGAACTGATCCTGGTGGGCA
ATCCATTTACATGCTCCTGTGACATTATGTGGATCAAGACTCTCCAAGAGGCTAAATCCA
GTCCAGACACTCAGGATTTGTACTGCCTGAATGAAAGCAGCAAGAATATTCCCCTGGCAA
ACCTGCAGATACCCAATTGTGGTTTGCCATCTGCAAATCTGGCCGCACCTAACCTCACTG
TGGAGGAAGGAAAGTCTATCACATTATCCTGTAGTGTGGCAGGTGATCCGGTTCCTAATA
TGTATTGGGATGTTGGTAACCTGGTTTCCAAACATATGAATGAAACAAGCCACACACAGG
GCTCCTTAAGGATAACTAACATTTTCATCCGATGACAGTGGGAAGCAGATCTCTTGTGTGG
CGGAAATCTTGTAGGAGAAGATCAAGATTCTGTCAACCTCACTGTGCATTTTGCACCAA
CTATCACATTTCTCGAATCTCCAACCTCAGACCACCACTGGTGCATTCCATTCACTGTGA
AAGGCAACCCCAAACCAGCGCTTCAGTGGTTCTATAACGGGGCAATATTGAATGAGTCCA
AATACATCTGTACTAAAATACATGTTACCAATCACACGGAGTACCACGGCTGCCTCCAGC
TGGATAATCCCACTCACATGAACAATGGGGACTACACTCTAATAGCCAAGAATGAGTATG
GGAAGGATGAGAAACAGATTTCTGCTCACTTCATGGGCTGGCCTGGAATTGACGATGGTG
CAAACCCAAATTATCCTGATGTAATTTATGAAGATTATGGAAGTGCAGCGAATGACATCG
GGGACACCACGAACAGAAGTAATGAAATCCCTTCCACAGACGTCCTGATAAAACCGGTC
GGGAACATCTCTCGGTCTATGCTGTGGTGGTGATTGCGTCTGTGGTGGGATTTTGCCTTT
TGGTAATGCTGTTTCTGCTTAAGTTGGCAAGACACTCCAAGTTTGGCATGAAAGATTTCT
CATGGTTTGGATTTGGGAAAGTAAAATCAAGACAAGGTGTTGGCCAGCCTCCGTTATCA
GCAATGATGATGACTCTGCCAGCCCACTCCATCACATCTCCAATGGGAGTAACACTCCAT
CTTCTTCGGAAGGTGGCCAGATGCTGTCAATTATTGGAATGACCAAGATCCCTGTCATTG
AAAATCCCCAGTACTTTGGCATCACCAACAGTCAGCTCAAGCCAGACACATTTGTTTCAGC
ACATCAAGCGACATAACATTGTTCTGAAAAGGGAGCTAGGCGAAGGAGCCTTTGGAAAAG
TGTTCCCTAGCTGAATGCTATAACCTCTGTCCCTGAGCAGGACAAGATCTTGGTGGCAGTGA
AGACCCTGAAGGATGCCAGTGACAATGCACGCAAGGACTTCCACCGTGAGGCCGAGCTCC
TGACCAACCTCCAGCATGAGCACATCGTCAAGTTCTATGGCGTCTGCGTGGAGGGCGACC
CCCTCATCATGGTCTTTGAGTACATGAAGCATGGGGACCTCAACAAGTTCTTCAGGGGCAC
ACGGCCCTGATGCCGTGCTGATGGCTGAGGGCAACCCGCCACGGAAGTACGCGAGTCGC
AGATGCTGCATATAGCCCAGCAGATCGCCGCGGGCATGGTCTACCTGGCGTCCCAGCACT
TCGTGCACCGCGATTTGGCCACCAGGAAGTGCCTGGTGGGGAGAACTTGCTGGTGAAAA
TCGGGGACTTTGGGATGTCCCGGGACGTGTACAGCACTGACTACTACAGGGTCGGTGGCC
ACACAATGCTGCCCATTCGCTGGATGCCTCCAGAGAGCATCATGTACAGGAAATTCACGA
CGGAAAGCGACGTCTGGAGCCTGGGGGTGCTGTTGTGGGAGATTTTCACCTATGGCAAAC
AGCCCTGGTACCAGCTGTCAAACAATGAGGTGATAGAGTGTATCACTCAGGGCCGAGTCC
TGCAGCGACCCCGCACGTGCCCCCAGGAGGTGTATGAGCTGATGCTGGGGTGTGTCAGC
GAGAGCCCCACATGAGGAAGAACATCAAGGGCATCCATACCCTCCTTCAGAACTTGGCCA
AGGCATCTCCGGTCTACCTGGACATTCTAGGCTAGGGCCCTTTTCCCCAGACCGATCCTT
CCCAACGTACTCCTCAGACGGGCTGAGAGGATGAACATCTTTTAACTGCCGCTGGAGGCC
ACCAAGCTGCTCTCCTTCACTCTGACAGTATTAACATCAAAGACTCCGAGAAGCTCTCGA
GGGAAGCAGTGTGTACTTCTTCATCCATAGACACAGTATTGACTTCTTTTTTGGCATTATC
TCTTTCTCTCTTCCATCTCCCTTGGTTGTTCTTTTTTCTTTTTTTAAATTTTCTTTTTT
TTTTTTTTTTTCGTCTTCCCTGCTTCACGATTCTTACCCTTTCTTTTGAATCAATCTGGCT
TCTGCATTACTATTAAGCTCTGCATAGACAAAGGCCTTAACAAACGTAATTTGTTATATCA
GCAGACACTCCAGTTTGGCCACCACAATAACAATGCCTTGTTGTATTCTGCCTTTGAT
GTGGATGAAAAAAGGGAAAACAAATATTTCACTTAAACTTTGTCACTTCTGCTGTACAG
ATATCGAGAGTTTCTATGGATTCACTTCTATTTATTTATTATTACTGTTCTTATTGT
TTTTGGATGGCTTAAGCCTGTGTATAAAAAAGAAAACCTTGTGTTCAATCTGTGAAGCCTT
TATCTATGGGAGATTAAAACCAGAGAGAAAGAAGATTTATTATGAACCGCAATATGGGAG
GAACAAAGACAACCACTGGGATCAGCTGGTGTGAGTCCCTACTTAGGAAATACTCAGCAA
CTGTTAGCTGGGAAGAATGTATTCGGCACCTTCCCTGAGGACCTTCTGAGGAGTAAAA
AGACTACTGGCCTCTGTGCCATGGATGATTCTTTTCCCATCACCAGAAATGATAGCGTGC

AGTAGAGA&CAAAGAATGGCTTCCGTGAGACACAAGATGGCGCATAGTGTGCTCGGACACA
GTTTTGTCTTCTGTAGGTTGTGATGATAGCACTGGTTTGTCTTCTCAAGCGCTATCCACAGA
ACCTTTGTCAACTTTCAGTTGAAAAGAGGTGGATTTCATGTCCAGAGCTCATTTTCGGGGTCA
GGTGGGAAAGCCAAGAACTTGGAAAAGATAAGACAAGCTATAAATTCGGAGGCAAGTTTC
TTTTACAATGAACTTTTTCAGATCTCACTTCCCTCCGACCCCTAACTTCCATGCCCACCCG
TCCTTTTAACTGTGCAAGCAAATTTGTGCATGGTCTTCGTCGATTAATACTTGTGTGCA
GACACTACTGCTCCAGACGTCGTTTCCCTGATAGGTAGAGCAGATCCATAAAAAGGTATG
ACTTATACAATTAGGGGAAGCTAATGGAGTTTATTAGCTGAGTATCAATGTCTCTGCGTT
GTACGGTGGTGATGGGTTTTAATGAATATGGACCCTGAAGCCTGGAAATCCTCATCCACG
TCGAACCCACAGGACTGTGGGAAGGGCAGAATCAATCCCTAAGGGAAAGGAAACCTCACC
CTGAGGGCATCACATGCACTCATGTTCAGTGTACACAGGTCAAGTCCCTTGCTCTGGGCT
CTAGTTGGGAGAGTGGTTTTCATTCCAAGTGTACTCCATTGTGAGTATGCTGTTTTTGT
CCTTCACTCCATTCAAAAAGTCAAAAATACAAAATTTGGCACAGCATGCCAACGGGAGGCT
GTGCCCAGACCAAGCACTGGAAGTGTGCTTCTAGGCATAGTCATTGGTTTTTGCAAAAAGA
GGGCTCAAATTTAAATAGAAATTTACAGCTATTTGAATGGTCAGATATACCAAGAAAGAA
AAATATTTCTGTTCTCAAGAAAACCTTGCTACCCTCTGTGAGGGGAATTTTGCTAAACTT
GACATCTTTATAACATGAGCCAGATTGAAAGGGAGTGATTTTCATTCATCTTAGGTCATG
TTATTTTCATATTTGTTTCTGAAGGTGCGATAGCTCTGTTTTAGGTTTTTGCTTGCGCCTGT
TAATTACTGGAACACCTTATTTTTTCATTAAAGGCTTTGAAAGCCAATTCTCAAAAATTCA
AAAGTGCAAATTAACAGAACAAAAGGAAATCCAGTAGCAACTGCAGTCAAGCGAGGGAGT
TGACAAGATAAACCTTACGTCCATTCAAGTTATATGCTGGCCTATGAGAGATGAGAGTTG
GGTCGTTTTGTTCTCTTTGTTGATGATTT (SEQ ID NO:3)

MSSWIRWHGPAMARLWGFCLVVGFWRAAFACPTSCCKCSASRIWCSDPSPGIVAFPRLEP
NSVDPENITEIFIANQKRLEIINEDDVEAYVGLRNLTI VDSGLKFVAHKAFLKNSNLQHI
NFTRNKLTSLSRKHFRHLDLSELI LVGNPFTCSCDIMWI KTLQEAKSSPDTQDLYCLNES
SKNIPLANLQIPNCGLP SANLAAPNLTVEEGKSITLSCSVAGDPVPNMYWDVGNLVSKHM
NETSHTQGSLRITNISSDDSGKQISCVAENLVGEDQDSVNLT VHFAPTITFLESPTSDH
WCIPFTVKG NPKPALQW FYNGAILNESKYICTKIHVTNHT EYHGC LQLDNPTHMNGDYT
LIAKNEYGKDEKQISAHFMGWPGIDDGANPNYPDVIYEDYGT AANDIGD TTNRSNEIPST
DVTDKTGREHLSVYAVVVIASVVGFC LLVMLFLLKLARHSKFGMKDFSWFGFGKVKSRQG
VGPASVISNDDDSASPLHHISNGSNT PPSSEGGPDAVIGMTKIPVIENPQYFGITNSQL
KPDTFVQHIKRHNIVL KRELGE GAFGKVFLAECYNLCPEQDKILVAVKTLKDASDNARKD
FHREAELLTNLQHEHIVKFYGV CVEGDPLIMVFEYMKHGDLNKFLRAHGPD AVLMAEGNP
PTELTQSQMLHIAQQIAAGMVYLASQHFVHRDLATRNCLVGENLLVKIGDFGMSRDVYST
DYRVRGGHTMLPIRWMPPESIMYRKFTTESDVWSLGVVLWEIFTY GKQPWYQLSNNEVIE
CITQGRVLQRPRTC PQEVYELMLGCWQREPHMRKNIKGIHTLLQNLAKASPVYLDILG (S
EO ID NO:4).

[illegible]

GATCTGGCTTCTGTACTCCTATTCAGTGTACATAGACAAAGGCCTTAACAAACCTGATTT
 GTTATATCAGCAGACACTCCAGTTTGCCCAACCACTAACAATGCCTTGTTGTATTCCT
 GCCTTTGATGTGGATGAAAAAAGGGAAAAAATAATCAAACATCTGACTTAAACCGTC
 ACTTCCGATGTACAGACACGGGGCGTTTCTATGGATTCACTTCTATCTATCTATTTATTT
 ATTTATCTATTTATTTATTTCTCTTCTTTGTTGTTTTCCGGTGGTTTTAGCCTGTGTATG
 AGAAGGGAAAGTCATGTACAGTCTGGGAAAACCTTTATCTGTGGGAAATGGAAACCAGAAG
 GGGAAAGAAGCTTTACCATAAAGCACAGCAGGAGTGAGACACAGAAAAGCCATTGGATCA
 GCCAGAGTCCGTCCTGCATAGGAAAACCCAGCAGCCATCAGGCTGGAGGATCATGTTCCGG
 CACTGACCCCCGAGGACCTTTCTGAGGAGGACACAGAAATGTTAAACTCTGCATCATGGAC
 ACAGTTTCCGATCACAGATACTGGCCTTCAATGGAAAAAATAAATAAATAAATAAATAAATAA
 GTTCTTGTGAGACCTGGACAGCACGTCCAACATCCAGACATTGTGGTCGGGCACAGTGAC
 AGAGTTGATGCATTTCTCACGGGTATTCTACAGAGCTTTTGTCAAGTCCAATGGAAGGA
 GGTAGATTCTTGTTTCAGATATGATTTCTGGGAAAAACCGAGTCTTGACAAAGACAGGAGA
 CACCCTCAGTTGGGAGGCAAGTTTCTCTTACCTTGGACTTTCTCACACAGCAATTCTCAC
 CCCCACCCCTCCACTCTCACCTGTCTTGTAAGTGTGCAACAAAAGTGTGCATGGTCTT
 TGTCAGTTGATACCTTTGTGCACCTCTGTGCAGAACTGCTGTCTGTCCCGGCTGTGGTA
 CCCGATCAGTGGGGTAGATCCACGAAAGGTCTCATTTTAGGCCGCTTTGGGAAGGTAACC
 AGATCGGTAGCTGGAAGCACTCTCCAGTAGGTGGCGAAGGGTGAGTGGGTCTGCTGAAGC
 CTGCATATCTTCACCCACCTCAAACCCACCGGGCTGCACAGGGGACAGGCACAGGCCACC
 CCTGAGGGACAGGGAAGCTCTCTTGGGATACCACCTGAGTTTACATTCAGTGTGCTCAGG
 TCAAGTCTCTCGCTCGGGGCTCTGTTTCGGGGAGAATGGTTTCATTCCAACGCACTCATT
 ATCAGGATTCTGTTTTT (SEQ ID NO:5)

[0091] The following polypeptide sequence was reported as the rat TrkB polypeptide sequence (encoded by nucleotides 665-3130 of SEQ ID NO:5) and has the accession number NP_036863 in Genbank.

MSPWPRWHGPAMARLWGLCLLVLGFWRASLACPMSCCKSTTRIWCTEPPGIVAFPRLEP
 NSIDPENITEILIANQKRLEI INEDDVEAYVGLKNLTIVDSGLKFVAYKAFLKNGNLRHI
 NFTRNKLTSLRRHFRHLDLSDLI LTGNPFTCSCDIMWLKTLQETKSSPDTQDLYCLNES
 SKNTPLANLQIPNCGLP SARLAAPNLTVEEGKSVTISCSVGGDPLPTLYWDVGNLVSKHM
 NETSHTQGSLRITNISSDDSGKQISCVAENLVGEDQDSVNLTVHFAPTITFLESPTSDHH
 WCIPFTVRGNPKPALQWFYNGAILNESKYICTKIHVNTNHTHEYHGCLQLDNPTHMNNGDYT
 LMAKNEYGKDERQISAHFMGRPGVDYETNPNYPEVLYEDWTTPTDIDGTTNKSNEIPSTD
 VADQTNREHLSVYAVVVIASVVGFCLLVMLLLLKLARHSKFGMKGPASVISNDDDSASPL
 HHISNGSNTSPSSSEGPD A VIIGMTKIPVIENPQYFGITNSQLKPDFTVQHIKRHNIVLK
 RELGEGAFGKVFLAECYNLCPEQDKILVAVKTLKDASDNARKDFHREAELLTNLQHEHIV
 KFYGVCEGDPLIMVFEYMKHGD LNKFLRAHGPD AVLMAEGNPPTELTQSQMLHIAQQIA
 AGMVYLASQHFVHRDLATRNCLVGENLLVKIGDFGMSRDVYSTDYRVGGHTMLPIRWMP
 PESIMYRKFTTESDVWSLGVVLWEIFTY GKQPWYQLSNNEVIECITQGRVLQRPRTCPOE
 VYELMLGCWQREPHTRKNIKNIHTLLQNLAKASPVYLDILG (SEQ ID NO:6)

[0092] Table 2 lists the TrkB domains and other regions, according to amino acid residue number, based on SEQ ID NO:4. As one of skill in the art will appreciate, the beginning and ending residues of the domains listed below may vary depending upon the computer modeling program used or the method used for determining the domain.

Table 2

Domain or Region	Beginning Residue	Ending Residue
Signal Peptide	1	31
Cysteine-rich 1	32	67
LRR	68	139
Cysteine-rich 2	140	195
Ig-Like 1	214	270
Ig-Like 2	301	365
Transmembrane	434	454
Tyrosine Kinase Catalytic Domain	552	828

Methods Using Antagonists of LINGO-1

[0093] One embodiment of the present invention provides methods for inhibiting LINGO-1 and TrkB interaction in a cell comprising contacting a cell co-expressing LINGO-1 and TrkB with a LINGO-1 antagonist. A LINGO-1 antagonist for the purposes of the present invention may be a LINGO-1 antagonist polypeptide, a LINGO-1 antibody, a LINGO-1 antagonist polynucleotide, a LINGO-1 aptamer or a combination of two or more LINGO-1 antagonists.

[0094] Additional embodiments of the invention include methods for promoting TrkB phosphorylation in a cell, relative to the level of TrkB phosphorylation in the absence of a LINGO-1 antagonist, comprising contacting a cell co-expressing LINGO-1 and TrkB with a LINGO-1 antagonist. Similarly, the invention also provide methods for promoting TrkB signal transduction comprising contacting cells co-expressing LINGO-1 and TrkB with a LINGO-1 antagonist. The invention also includes methods for promoting TrkB phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.

[0095] Further embodiments of the invention include methods for inhibiting JNK phosphorylation in a cell comprising contacting a cell co-expressing LINGO-1 and JNK with a LINGO-1 antagonist as well as methods for inhibiting JNK pathway signal transduction comprising contacting a cell co-expressing LINGO-1 and JNK with a LINGO-1 antagonist. Inhibiting JNK phosphorylation, according to the present invention, includes inhibition of JNK-1 phosphorylation, inhibition of JNK-2 phosphorylation or inhibition of JNK-1 and JNK-2 phosphorylation. Inhibiting JNK

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phosphorylation, according to the present invention, can include decreasing the overall quantity of phosphorylated JNK protein, or decreasing the length of time for which JNK proteins remain phosphorylated. The invention also includes methods for inhibiting JNK phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.

[0096] An additional embodiment of the invention provides a method for promoting survival of retinal ganglion cells in a mammal displaying signs or symptoms of a pressure induced ocular neuropathy comprising administering to a mammal in need of such treatment an effective amount of a LINGO-1 antagonist and a carrier. In one embodiment, the pressure induced ocular neuropathy is glaucoma.

Methods Using Antagonists of LINGO-1 and TrkB Agonists

[0097] In some methods of the present invention both a LINGO-1 antagonist and a TrkB agonist are used. For example, one embodiment of the invention includes a method for promoting survival of a neuron at risk of dying comprising contacting the neuron with an effective amount of a combination of a LINGO-1 antagonist and a TrkB agonist. In embodiments of the invention in which both a LINGO-1 antagonist and a TrkB agonist are used the term "effective amount" refers to the amount of the combination of the LINGO-1 antagonist and the TrkB agonist that is sufficient to produce the desired result. In some instances the amount of a LINGO-1 antagonist required to produce the desired effect may be greater when the LINGO-1 antagonist is used alone than when it is used in combination with a TrkB agonist. Similarly, in some instances, the amount of a TrkB agonist that is required to produce the desired effect may be greater when the TrkB agonist is used alone than when it is used in combination with a LINGO-1 antagonist.

[0098] Another embodiment of the invention provides methods for treating a disease or disorder associated with neuronal cell death comprising administering to an animal in need of such treatment an effective amount of a combination of a LINGO-1 antagonist and a TrkB agonist. In some particular embodiments, the disease or disorder can be a pressure induced optical neuropathy. Further embodiments of the invention include methods for promoting regeneration or survival of CNS neurons in a mammal displaying signs or symptoms of a condition involving neuronal cell death comprising administering to the mammal an effective amount of a combination of a LINGO-1 antagonist and a TrkB agonist. In some embodiments of the invention, the disease or disorder is ALS, Huntington's disease, Alzheimer's disease, Parkinson's disease, diabetic neuropathy,

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stroke or hearing loss. In another particular embodiments, the CNS neurons are sensory neurons such as retinal ganglion cells or hairy cells.

[0099] In embodiments of the invention that involve use a combination of a LINGO-1 antagonist and a TrkB agonist, the LINGO-1 antagonist and TrkB agonist can be administered in a single composition or can be administered separately. In addition, the LINGO-1 antagonist and the TrkB agonist can be administered simultaneously or sequentially.

LINGO-1 Antagonist and TrkB Agonist Compounds

[00100] LINGO-1 antagonists in the methods of the present invention include any chemical or synthetic compound which inhibits or decreases the activity of LINGO-1 compared to the activity of LINGO-1 in the absence of the antagonist compound.

[0100] TrkB agonists in the methods of the present invention include any chemical or synthetic compound which promotes or increases the activity of TrkB or promotes or increases the phosphorylation of TrkB when compared to the state of TrkB in the absence of the agonist compound.

[0101] TrkB agonist compounds include, but are not limited to neurotrophic factor mimetics. TrkB agonist compounds also include, but are not limited to L-783,281 adenosine, CGS 21680, etc. as reviewed in Pollack *et al. Curr. Drug Targ- CNS and Neurol. Disorders* 1:59-80 (2002), which is herein incorporated by reference in its entirety. In some embodiments, the TrkB agonist compounds are selective for TrkB and activate TrkB to a greater extent than TrkA or TrkC. In some embodiments, the TrkB agonist compounds are specific for TrkB and do not activate TrkA or TrkC. In addition, the TrkB agonist compounds can also be small molecules that mimic critical regions of neurotrophins. For example, the small molecule can be a mimetic of a BDNF β -turn loop. Particular examples of small molecule mimetics that may be used according to the invention are disclosed in U.S. Published Application No. 2007/0060526 A1, which is incorporated herein by reference in its entirety.

[0102] One of ordinary skill in the art would know how to screen and test for LINGO-1 antagonist and TrkB agonist compounds which would be useful in the methods of the present invention, for example by screening for compounds that modify neuronal survival using assays described elsewhere herein.

Soluble LINGO-1 Antagonist and TrkB Agonist Polypeptides

Soluble LINGO-1 polypeptides

[0103] LINGO-1 antagonists to be used in the methods of the present invention include those polypeptides which block, inhibit or interfere with the biological function of naturally occurring LINGO-1. Specifically, soluble LINGO-1 polypeptides of the present invention include fragments, variants, or derivatives thereof of a soluble LINGO-1 polypeptide. Table 1 above describes the various domains of the LINGO-1 polypeptide. Soluble LINGO-1 polypeptides lack the transmembrane domain and typically lack the intracellular domain of the LINGO-1 polypeptide. For example, certain soluble LINGO-1 polypeptides lack amino acids 552-576 which comprise the transmembrane domain of LINGO-1 and/or amino acids 577-614 which comprise the intracellular domain of LINGO-1. Additionally, certain soluble LINGO-1 polypeptides comprise the LRR domains, Ig domain, basic region and/or the entire extracellular domain (corresponding to amino acids 34 to 532 of SEQ ID NO: 2) of the LINGO-1 polypeptide. As one of skill in the art would appreciate, the entire extracellular domain of LINGO-1 may comprise additional or fewer amino acids on either the C-terminal or N-terminal end of the extracellular domain polypeptide.

[0104] As such, soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 41 to 525 of SEQ ID NO:2; 40 to 526 of SEQ ID NO:2; 39 to 527 of SEQ ID NO:2; 38 to 528 of SEQ ID NO:2; 37 to 529 of SEQ ID NO:2; 36 to 530 of SEQ ID NO:2; 35 to 531 of SEQ ID NO:2; 34 to 531 of SEQ ID NO:2; 46 to 520 of SEQ ID NO:2; 45 to 521 of SEQ ID NO:2; 44 to 522 of SEQ ID NO:2; 43 to 523 of SEQ ID NO:2; and 42 to 524 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides. LINGO-1 polypeptide antagonists may include any combination of domains as described in Table 1.

[0105] Additional soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 1 to 33 of SEQ ID NO:2; 1 to 35 of SEQ ID NO:2; 34 to 64 of SEQ ID NO:2; 36 to 64 of SEQ ID NO:2; 66 to 89 of SEQ ID NO:2; 90 to 113 of SEQ ID NO:2; 114 to 137 of SEQ ID NO:2; 138 to 161 of SEQ ID NO:2; 162 to 185 of SEQ ID NO:2; 186 to 209 of SEQ ID NO:2; 210 to 233 of SEQ ID NO:2; 234 to 257 of SEQ ID NO:2; 258 to 281 of SEQ ID NO:2; 282 to 305 of SEQ ID NO:2;

306 to 329 of SEQ ID NO:2; 330 to 353 of SEQ ID NO:2; 363 to 416 of SEQ ID NO:2; 417 to 424 of SEQ ID NO:2; 419 to 493 of SEQ ID NO:2; and 494 to 551 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.

[0106] Further soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 1 to 33 of SEQ ID NO:2; 1 to 35 of SEQ ID NO:2; 1 to 64 of SEQ ID NO:2; 1 to 89 of SEQ ID NO:2; 1 to 113 of SEQ ID NO:2; 1 to 137 of SEQ ID NO:2; 1 to 161 of SEQ ID NO:2; 1 to 185 of SEQ ID NO:2; 1 to 209 of SEQ ID NO:2; 1 to 233 of SEQ ID NO:2; 1 to 257 of SEQ ID NO:2; 1 to 281 of SEQ ID NO:2; 1 to 305 of SEQ ID NO:2; 1 to 329 of SEQ ID NO:2; 1 to 353 of SEQ ID NO:2; 1 to 416 of SEQ ID NO:2; 1 to 424 of SEQ ID NO:2; 1 to 493 of SEQ ID NO:2; 1 to 551 of SEQ ID NO:2; 1 to 531 of SEQ ID NO:2 and 1 to 532 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.

[0107] Still further soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 34 to 64 of SEQ ID NO:2; 34 to 89 of SEQ ID NO:2; 34 to 113 of SEQ ID NO:2; 34 to 137 of SEQ ID NO:2; 34 to 161 of SEQ ID NO:2; 34 to 185 of SEQ ID NO:2; 34 to 209 of SEQ ID NO:2; 34 to 233 of SEQ ID NO:2; 34 to 257 of SEQ ID NO:2; 34 to 281 of SEQ ID NO:2; 34 to 305 of SEQ ID NO:2; 34 to 329 of SEQ ID NO:2; 34 to 353 of SEQ ID NO:2; 34 to 416 of SEQ ID NO:2; 34 to 424 of SEQ ID NO:2; 34 to 493 of SEQ ID NO:2; and 34 to 551 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.

[0108] Additional soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 34 to 530 of SEQ ID NO:2; 34 to 531 of SEQ ID NO:2; 34 to 532 of SEQ ID NO:2; 34 to 533 of SEQ ID NO:2; 34 to 534 of SEQ ID NO:2; 34 to 535 of SEQ ID NO:2; 34 to 536 of SEQ ID NO:2; 34 to 537 of SEQ ID NO:2; 34 to 538 of SEQ ID NO:2; 34 to 539 of SEQ ID NO:2; 30 to 532 of SEQ ID NO:2; 31 to 532 of SEQ ID NO:2; 32 to 532 of SEQ ID NO:2; 33 to 532 of SEQ ID NO:2; 34 to 532 of SEQ ID NO:2; 35 to 532 of SEQ ID NO:2; 36 to 532 of SEQ ID NO:2; 30 to 531 of SEQ ID NO:2; 31 to 531 of SEQ ID NO:2; 32 to 531 of SEQ ID NO:2; 33 to 531 of SEQ ID NO:2; 34 to 531 of SEQ ID NO:2; 35 to 531 of SEQ ID NO:2; and 36 to 531 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.

- [0109] Still further soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 36 to 64 of SEQ ID NO:2; 36 to 89 of SEQ ID NO:2; 36 to 113 of SEQ ID NO:2; 36 to 137 of SEQ ID NO:2; 36 to 161 of SEQ ID NO:2; 36 to 185 of SEQ ID NO:2; 36 to 209 of SEQ ID NO:2; 36 to 233 of SEQ ID NO:2; 36 to 257 of SEQ ID NO:2; 36 to 281 of SEQ ID NO:2; 36 to 305 of SEQ ID NO:2; 36 to 329 of SEQ ID NO:2; 36 to 353 of SEQ ID NO:2; 36 to 416 of SEQ ID NO:2; 36 to 424 of SEQ ID NO:2; 36 to 493 of SEQ ID NO:2; and 36 to 551 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.
- [0110] Additional soluble LINGO-1 polypeptides for use in the methods of the present invention include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 36 to 530 of SEQ ID NO:2; 36 to 531 of SEQ ID NO:2; 36 to 532 of SEQ ID NO:2; 36 to 533 of SEQ ID NO:2; 36 to 534 of SEQ ID NO:2; 36 to 535 of SEQ ID NO:2; 36 to 536 of SEQ ID NO:2; 36 to 537 of SEQ ID NO:2; 36 to 538 of SEQ ID NO:2; and 36 to 539 of SEQ ID NO:2; or fragments, variants, or derivatives of such polypeptides.
- [0111] Additional soluble LINGO-1 polypeptides, fragments, variants or derivatives thereof include polypeptides comprising the Ig domain of LINGO-1. For example, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 417 to 493 of SEQ ID NO:2; 417 to 494 of SEQ ID NO:2; 417 to 495 of SEQ ID NO:2; 417 to 496 of SEQ ID NO:2; 417 to 497 of SEQ ID NO:2; 417 to 498 of SEQ ID NO:2; 417 to 499 of SEQ ID NO:2; 417 to 500 of SEQ ID NO:2; 417 to 492 of SEQ ID NO:2; 417 to 491 of SEQ ID NO:2; 412 to 493 of SEQ ID NO:2; 413 to 493 of SEQ ID NO:2; 414 to 493 of SEQ ID NO:2; 415 to 493 of SEQ ID NO:2; 416 to 493 of SEQ ID NO:2; 411 to 493 of SEQ ID NO:2; 410 to 493 of SEQ ID NO:2; 410 to 494 of SEQ ID NO:2; 411 to 494 of SEQ ID NO:2; 412 to 494 of SEQ ID NO:2; 413 to 494 of SEQ ID NO:2; 414 to 494 of SEQ ID NO:2; 415 to 494 of SEQ ID NO:2; 416 to 494 of SEQ ID NO:2; 417 to 494 of SEQ ID NO:2; and 418 to 494 of SEQ ID NO:2 or fragments, variants, or derivatives of such polypeptides.
- [0112] Soluble LINGO-1 polypeptides for use in the methods of the present invention also include combinations of two or more soluble LINGO-1 polypeptides disclosed herein. The two or more soluble LINGO-1 polypeptides for use in the methods of the invention may be fused together to form a single polypeptide comprising multiple

LINGO-1 soluble polypeptides disclosed herein or may be individual soluble LINGO-1 polypeptides comprising a composition for use in the methods of the present invention.

[0113] Various exemplary soluble LINGO-1 polypeptides and methods and materials for obtaining these molecules for practicing the present invention are described below and/or may be found, *e.g.*, in PCT Publication Nos. WO2004/085648, WO 2006/002437, WO 2007/0059793, WO 2007/008547, WO 2007/056161 and WO 2007/064882 and U.S. Published Application No. 2006/017673, which are incorporated herein by reference in their entireties.

TrkB agonist polypeptides

[0114] TrkB agonist polypeptides for use in the methods of the present invention include any polypeptide which can promote, amplify, enhance or increase the activity of TrkB. Such proteins include, but are not limited to, TrkB ligands including neurotrophins such as BDNF, NT-3 and NT-4/5. Additionally, TrkB agonist polypeptides of the present invention also include fragments, variants or derivatives of TrkB ligands as well as chimeric neurotrophin molecules. In some embodiments, the TrkB ligand polypeptides, fragments, variants or derivatives thereof are dimerized or multimerized. TrkB agonist polypeptides of the present invention can act as homodimers or heterodimers. TrkB agonist polypeptides of the present invention also include pan-neurotrophins such as those disclosed in Ibanez *et al.* *EMBO* 12:12:2281-2293 (1993). In particular, TrkB agonist polypeptides of the invention include domains of TrkB ligands that bind to TrkB or variants of such polypeptides. In one particular embodiment of the invention, the TrkB agonist is Brain-Derived Neurotrophic Factor (BDNF), or a fragment, variant or derivative thereof.

[0115] The TrkB agonist polypeptide of the invention may be a hairpin motif of a TrkB ligand as disclosed in U.S. Patent No. 7,205,387. Additionally TrkB ligand polypeptides, fragments, variants or derivatives thereof may be fused to other protein or peptide sequences and/or may be attached to water soluble polymers including polyethylene glycol as disclosed in U.S. Patent No. 5,770,577 or to 1-acyl-glycerol derivatives as disclosed in U.S. Patent No. 6,800,607. Additionally or alternatively, the TrkB ligands, fragments or variants thereof can be altered in order to increase stability upon administration, for example, by using variants with lower isoelectric points as described in U.S. Patent No. 6,723,701. TrkB ligands, as well as fragments and variants thereof, can

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be arranged in tandem and cyclized, for example, to form "mini-neurotrophins", for example B_{AG}, as described in Williams *et al. JBC* 280:5862-5869 (2004).

[0116] Additionally, TrkB agonists polypeptides for use in the methods of the present invention include, but are not limited to TrkB polypeptides, fragments, variants or derivatives thereof. TrkB polypeptides, fragments, variants or derivatives thereof for use in the methods of the present invention also include a TrkB polypeptide comprising, consisting essentially of, or consisting of the full-length TrkB protein fused to an immunoglobulin domain, for example, a TrkB polypeptide comprising, consisting essentially of, or consisting of amino acids 1 to 828 of SEQ ID NO:4; 32 to 828 of SEQ ID NO:4 fused to an IgG domain. Additional TrkB polypeptides, fragments, variants or derivatives thereof described herein may also be fused to an immunoglobulin domain. In some embodiments, the TrkB polypeptides of the present invention are dimerized or multimerized. TrkB polypeptides of the present invention also include TrkB fragments, variants, isoforms or derivatives that do not interact with LINGO-1, have an increased tendency for dimerization or multimerization, have an increased affinity for endogenous or non-endogenous ligands and/or have increased kinase activity compared to the polypeptide of SEQ ID NO:4.

[0117] Additional TrkB agonist polypeptides for use in the methods of the present invention also include a combination of two or more TrkB agonist polypeptides disclosed herein. The two or more TrkB agonist polypeptides for use in the methods of the present invention may be fused together to form a single polypeptide comprising multiple TrkB agonist polypeptides disclosed herein or may be individual TrkB agonist polypeptides comprising a composition for use in the methods of the present invention.

Soluble LINGO-1 antagonist and/or trkB agonist polypeptides

[0118] Soluble LINGO-1 antagonist and TrkB agonist polypeptides for use in the methods of the present invention described herein may be cyclic. Cyclization of the soluble LINGO-1 antagonist or TrkB agonist polypeptides reduces the conformational freedom of linear peptides and results in a more structurally constrained molecule. Many methods of peptide cyclization are known in the art, for example, "backbone to backbone" cyclization by the formation of an amide bond between the N-terminal and the C-terminal amino acid residues of the peptide. The "backbone to backbone" cyclization method includes the formation of disulfide bridges between two ω -thio amino acid residues (*e.g.* cysteine, homocysteine). Certain soluble LINGO-1 antagonist or TrkB agonist peptides of the present invention include modifications on the N- and C- terminus

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of the peptide to form a cyclic LINGO-1 antagonist or TrkB agonist polypeptide. Such modifications include, but are not limited to, cysteine residues, acetylated cysteine residues, cysteine residues with a NH₂ moiety and biotin. Other methods of peptide cyclization are described in Li & Roller, *Curr. Top. Med. Chem.* 3:325-341 (2002), which is incorporated by reference herein in its entirety.

[0119] Soluble LINGO-1 antagonist or TrkB agonist polypeptides described herein may have various alterations such as substitutions, insertions or deletions. For examples, substitutions include, but are not limited to the following substitutions: valine at position 6 of the LINGO-1 polypeptide of SEQ ID NO:2 to methionine; serine at position 294 of the LINGO-1 polypeptide of SEQ ID NO:2 to glycine; valine at position 348 of the LINGO-1 polypeptide of SEQ ID NO:2 to alanine; arginine at position 419 of the LINGO-1 polypeptide to histidine; arginine at position 456 to glutamic acid; and histidine at position 458 of SEQ ID NO:2 to valine.

[0120] Corresponding fragments of soluble LINGO-1 antagonist or TrkB agonist polypeptides at least 70%, 75%, 80%, 85%, 90%, or 95% identical to polypeptides of SEQ ID NO:2 or SEQ ID NO:4 described herein are also contemplated.

[0121] As known in the art, "sequence identity" between two polypeptides is determined by comparing the amino acid sequence of one polypeptide to the sequence of a second polypeptide. When discussed herein, whether any particular polypeptide is at least about 70%, 75%, 80%, 85%, 90% or 95% identical to another polypeptide can be determined using methods and computer programs/software known in the art such as, but not limited to, the BESTFIT program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711). BESTFIT uses the local homology algorithm of Smith and Waterman, *Advances in Applied Mathematics* 2:482-489 (1981), to find the best segment of homology between two sequences. When using BESTFIT or any other sequence alignment program to determine whether a particular sequence is, for example, 95% identical to a reference sequence according to the present invention, the parameters are set, of course, such that the percentage of identity is calculated over the full length of the reference polypeptide sequence and that gaps in homology of up to 5% of the total number of amino acids in the reference sequence are allowed.

[0122] Soluble LINGO-1 antagonist or TrkB-agonist polypeptides for use in the methods of the present invention may include any combination of two or more soluble LINGO-1 antagonist or TrkB agonist polypeptides.

Antibodies or Antigen-binding Fragments Thereof

[0123] In one embodiment, LINGO-1 antagonists or TrkB agonists for use in the methods of the invention include antibodies or antigen-binding fragments, variants or derivatives thereof which are LINGO-1 antagonists or TrkB agonists. For example, binding of certain LINGO-1 or TrkB antibodies to LINGO-1 or TrkB, as expressed by CNS neurons, promotes neuronal cell survival.

[0124] In some embodiments, the LINGO-1 antibody is the antibody 1A7. The 1A7 antibody has been described in International Application PCT/US06/26271, filed July 7, 2006, incorporated herein by reference in its entirety. Sequences of the 1A7 antibody are shown in the table below.

Table 3- 1A7 Antibody Sequences

	Polypeptide Sequence	SEQ ID NO:
VH	QVQLVQSGPELKKPGETVKISCKASGYTFTNYGMNWVKQAPGKGLKWN W INTDTGEPTYTEDFQGRFAFSLETSASTVYLQFNNLKNEDTATYFCAREGY F DYWGQGTTVTVSS	8
VL	QIVLTQSPAISASPGEKVTMTCSASSSVSYMHWYQQKSGTSPKRWIYDT KLASGVPARFSGSGSGTSYSLTISSMEAEDAATYYCQQWSSNPFTFGSGTI EIK	9

[0125] The antibody of the invention can also be an antibody that includes one or more than one CDR of the 1A7 antibody. The following are the sequences for the VH CDR1, CDR2 and CDR3 regions of the 1A7 antibody respectively: NYGMN (SEQ ID NO:10), WINTDTGEPTYTEDFQG (SEQ ID NO:11), and EGVHFDY (SEQ ID NO:12). The following are the sequences for the VL CDR1, CDR2 and CDR3 regions of the 1A7 antibody respectively: SASSSVSYMH (SEQ ID NO:13), DTSKLAS (SEQ ID NO:14), and QQWSSNPFT (SEQ ID NO:15).

[0126] In some embodiments, the LINGO-1 antibody is the monoclonal antibody 3B5.2 (also referred to as 3B5), which can be produced from the hybridoma 2.P3B5.2. The 2.P3B5.2 hybridoma was deposited with the American Type Culture Collection (ATCC) in Manassas, VA on December 27, 2006, and the 3B5 antibody has been described in U.S. Provisional Patent Application No. 60/879,324, filed on January 9, 2007, which is

herein incorporated by reference in its entirety. Sequences of the 3B5 antibody are shown in the table below.

Table 4- 3B5 Antibody Sequences

	Polypeptide Sequence	SEQ ID NO:
VH	QVQLQQPGAELVRPGTSVKLSCRASGYTFTSYWMHWWVKQRPGQGLE WIGVIDPSDSYTNYNQKFRGKATLTVDTSSTAYMQLSSLTSEDSAVY YCARPYYGSHWFFDVWGTGTTVTVSS	16
VL	QIVLTQSPAIMASAPGEKVTMTCSASSRVSYPVHWYQQKSGTSPKRWL YDTSNLASGVPARFGGNGSGTSYSLTISSMEAEDAATYYCQQWSTNP PTFGGGTKLEIK	17

[0127] The antibody of the invention can also be an antibody that includes one or more than one CDR of the 3B5 antibody. The following are the sequences for the VH CDR1, CDR2 and CDR3 regions of the 3B5 antibody respectively: SYWMH (SEQ ID NO:18), VIDPSDSYTNYNQKFRG (SEQ ID NO:19), and PYYGSHWFFDV (SEQ ID NO:20). The following are the sequences for the VL CDR1, CDR2 and CDR3 regions of the 3B5 antibody respectively: SASSRVSYPVH (SEQ ID NO:21), DTSNLAS (SEQ ID NO:22), and QQWSTNPPT (SEQ ID NO:23).

[0128] In some embodiments, the LINGO-1 antibody is the monoclonal antibody LI33. The LI33 antibody has been described in U.S. Provisional Patent Application No. 60/879,324, filed on January 9, 2007, which is herein incorporated by reference in its entirety. Sequences of the LI33 antibody are shown in the table below.

Table 5- LI33 Antibody Sequences

	Polypeptide Sequence	SEQ ID NO:
VH	EVQLLES GGGLVQPGGSLRLSCAASGFTFSIYPMFWVRQAPGKGLEW VSWIGPSGGITKYADSVKGRFTISRDN SKNTLYLQMNSLRAEDTATYY CAREGHNDWYFDLWGRGTLTVSS	24
VL	DIQMTQSPGTLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLI YDASN RATGIPARFSGSGSGTEFTLTIS SLQSEDFAVYYCQQYDKWPL TFGGG TKVEIK	25

- [0129] The antibody of the invention can also be an antibody that includes one or more than one CDR of the LI33 antibody. The following are the sequences for the VH CDR1, CDR2 and CDR3 regions of the LI33 antibody respectively: IYPMF (SEQ ID NO:26), WIGPSGGITKYADSVKG (SEQ ID NO:27), and EGHNDWYFDL (SEQ ID NO:28). The following are the sequences for the VL CDR1, CDR2 and CDR3 regions of the LI33 antibody respectively: RASQSVSSYLA (SEQ ID NO:29), DASNRAT (SEQ ID NO:30), and QQYDKWPLT (SEQ ID NO:31).
- [0130] In another embodiment the LINGO-1 antibody is the monoclonal antibody 7P1D5.1G9, which can be produced from the hydridoma 7.P1D5.1.G9. The 7.P1D5.1.G9 hybridomas was deposited with the American Type Culture Collection (ATCC) in Mannasssas, VA on December 27, 2006, and the 7P1D5.1G9 antibody is described in
- [0131] Certain LINGO-1 antagonist antibodies for use in the methods described herein specifically or preferentially bind to a particular LINGO-1 polypeptide fragment or domain. Such LINGO-1 polypeptide fragments include, but are not limited to, a LINGO-1 polypeptide comprising, consisting essentially of, or consisting of amino acids 34 to 532; 34 to 417, 34 to 425, 34 to 493, 66 to 532, 66 to 417 (LRR domain), 66 to 426, 66 to 493, 66 to 532, 417 to 532, 417 to 425 (the LINGO-1 basic region), 417 to 424 (the LINGO-1 basic region), 417 to 493, 417 to 532, 419 to 493 (the LINGO-1 Ig region), or 425 to 532 of SEQ ID NO:2, or a LINGO-1 variant polypeptide at least 70%, 75%, 80%, 85%, 90%, or 95% identical to amino acids 34 to 532; 34 to 417, 34 to 425, 34 to 493, 66 to 532, 66 to 417, 66 to 426, 66 to 493, 66 to 532, 417 to 532, 417 to 425 (the LINGO-1 basic region), 417 to 493, 417 to 532, 419 to 493 (the LINGO-1 Ig region), or 425 to 532 of SEQ ID NO:2.
- [0132] Additional LINGO-1 peptide fragments to which certain LINGO-1 specific antibodies, or antigen-binding fragments, variants, or derivatives thereof for use in the methods of the present invention bind include, but are not limited to, those fragments comprising, consisting essentially of, or consisting of one or more leucine-rich-repeats (LRR) of LINGO-1. Such fragments, include, for example, fragments comprising, consisting essentially of, or consisting of amino acids 66 to 89, 66 to 113, 66 to 137, 90 to 113, 114 to 137, 138 to 161, 162 to 185, 186 to 209, 210 to 233, 234 to 257, 258 to 281, 282 to 305, 306 to 329, or 330 to 353 of SEQ ID NO:2. Corresponding fragments of a variant LINGO-1 polypeptide at least 70%, 75%, 80%, 85%, 90%, or 95% identical to amino acids 66 to 89, 66 to 113, 90 to 113, 114 to 137, 138 to 161, 162 to 185, 186 to

209, 210 to 233, 234 to 257, 258 to 281, 282 to 305, 306 to 329, or 330 to 353 of SEQ ID NO:2 are also contemplated.

[0133] Additional LINGO-1 peptide fragments to which certain antibodies, or antigen-binding fragments, variants, or derivatives thereof of the present invention bind include, but are not limited to those fragments comprising, consisting essentially of, or consisting of one or more cysteine rich regions flanking the LRR of LINGO-1. Such fragments, include, for example, a fragment comprising, consisting essentially of, or consisting of amino acids 34 to 64 of SEQ ID NO:2 (the N-terminal LRR flanking region (LRRNT)), or a fragment comprising, consisting essentially of, or consisting of amino acids 363 to 416 of SEQ ID NO:2 (the C-terminal LRR flanking region (LRRCT)). Corresponding fragments of a variant LINGO-1 polypeptide at least 70%, 75%, 80%, 85%, 90%, or 95% identical to amino acids 34 to 64 and 363 to 416 of SEQ ID NO:2 are also contemplated.

[0134] In other embodiments, the LINGO-1 antagonists to be used in the methods described herein include an antibody, or antigen-binding fragment, variant, or derivative thereof which specifically or preferentially binds to at least one epitope of LINGO-1, where the epitope comprises, consists essentially of, or consists of at least about four to five amino acids of SEQ ID NO:2, at least seven, at least nine, or between at least about 15 to about 30 amino acids of SEQ ID NO:2. The amino acids of a given epitope of SEQ ID NO:2 as described may be, but need not be, contiguous or linear. In certain embodiments, at least one epitope of LINGO-1 comprises, consists essentially of, or consists of a non-linear epitope formed by the extracellular domain of LINGO-1 as expressed on the surface of a cell or as a soluble fragment, *e.g.*, fused to an IgG Fc region. Thus, in certain embodiments the at least one epitope of LINGO-1 comprises, consists essentially of, or consists of at least 4, at least 5, at least 6, at least 7, at least 8, at least 9, at least 10, at least 15, at least 20, at least 25, between about 15 to about 30, or at least 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, or 100 contiguous or non-contiguous amino acids of SEQ ID NO:2, where non-contiguous amino acids form an epitope through protein folding.

[0135] Certain TrkB agonist antibodies for use in the methods described herein specifically or preferentially bind to a particular TrkB polypeptide fragment or domain. For example, antibodies of the invention can bind epitope(s) in the IgG-1 or IgG-2 regions of TrkB and can bind sequences in the loop region of the IgG-1 and/or IgG-2 domains. Antibodies of the present invention also include those that increase TrkB autophosphorylation. Certain non-limiting examples of TrkB agonist antibodies also

include, but are not limited to, monoclonal antibodies which block binding of ligands such as BDNF to the TrkB receptor, monoclonal antibodies which partially block binding of ligands such as BDNF to the TrkB receptor, monoclonal antibodies that do not block the binding of ligands such as BDNF to the TrkB receptor and monoclonal antibodies that promote or increase the binding of ligands such as BDNF to the TrkB receptor. TrkB agonist antibodies of the present invention also include monoclonal antibodies that do not affect the binding of ligands such as BDNF to the TrkB receptor. Certain non-limiting examples of TrkB agonist antibodies include 6E2, 7F5, 11E1, 16E11, 17D11, 19E12, 29D7, which are described in U.S. Published Patent Application No. 2007/0059304 and in Quin *et al.*, *Jour. of Neuroscience*. 26:9394-9403 (2006), each of which is incorporated herein by reference and antibodies or antigen-binding fragments which bind to the same epitopes as those antibodies. Exemplary TrkB agonist antibodies for use in the methods of the present invention also include isolated antibodies or antigen binding fragments thereof which specifically bind to the same TrkB epitopes as the TrkB agonist antibodies listed above.

[0136] Exemplary antibodies or fragments thereof for use in the methods of the present invention include, but are not limited to, isolated antibodies or antigen binding fragments thereof which specifically binds to the same LINGO-1 epitope as a reference monoclonal antibody selected from the group consisting of 201', 3A3, 3A6, 3B5, 1A7, 1D5, 1G7, 2B10, 2C11, 2F3, 3P1B1.1F9, 3P1D10.2C3, 3P1E11.3B7, 3P2C6.3G10.2H7, 3P2C9.2G4, 3P4A6.1D9, 3P4A1.2B9, 3P4C2.2D2, 3P4C5.1D8, 3P4C8.2G9, 6P4F4.1Ds, 6P4F4.1F9, 7P1D5.1G9, 1B6.4, 2C7.2, 2D6.1, 2F7.3, 2H3.2, 3C11.1, 3E3.1, 3H11.2, 3G8.1, 2B8.1, 3B5.230-C12 (Li01), 38-D01 (Li02), 35-E04 (Li03), 36-C09 (Li04), 30-A11 (Li05), 34-F02 (Li06), 29-E07 (Li07), 34-G04 (Li08), 36-A12 (Li09), 28-D02 (Li10), 30-B01 (Li11), 34-B03 (Li12), Li13, Li32, Li33, Li34, 3383 (L1a.1), 3495(L1a.2), 3563 (L1a.3), 3564 (L1a.4), 3565 (L1a.5), 3566 (L1a.6), 3567 (L1a.7), 3568 (L1a.8), 3569 (L1a.9), 3570 (L1a.10), 3571 (L1a.11), 3582 (L1a.12), 1968 (L1a.13), 3011, 3012, 3013, 3418, 3422, 3562, D05, D07, D08, D10 and D11, as described in the International Application PCT/US06/26271 entitled "Sp35 Antibodies and Uses Thereof" to Mi *et al.*, filed July 7, 2006, incorporated herein by reference in its entirety.

[0137] In other embodiments, the LINGO-1 antagonists and TrkB agonists to be used in the methods of the present invention include LINGO-1 or TrkB antibodies, or antigen-binding fragments, variants, or derivatives thereof which specifically or preferentially bind to at least one epitope of LINGO-1 or TrkB, where the epitope comprises, consists

essentially of, or consists of, in addition to one, two, three, four, five, six or more contiguous or non-contiguous amino acids of SEQ ID NO:2 or SEQ ID NO:4, respectively, as described above, and an additional moiety which modifies the protein, *e.g.*, a carbohydrate moiety may be included such that the LINGO-1 or TrkB antibody binds with higher affinity to modified target protein than it does to an unmodified version of the protein. Alternatively, the LINGO-1 or TrkB antibody does not bind the unmodified version of the target protein at all.

[0138] In certain embodiments, the LINGO-1 antagonist or TrkB agonists to be used in the methods of the present invention include an antibody, or antigen-binding fragment, variant, or derivative thereof of the invention that binds specifically to at least one epitope of LINGO-1 or TrkB or fragment or variant described above, *i.e.*, binds to such an epitope more readily than it would bind to an unrelated, or random epitope; binds preferentially to at least one epitope of LINGO-1 or TrkB or fragment or variant described above, *i.e.*, binds to such an epitope more readily than it would bind to a related, similar, homologous, or analogous epitope; competitively inhibits binding of a reference antibody which itself binds specifically or preferentially to a certain epitope of LINGO-1 or TrkB or fragment or variant described above; or binds to at least one epitope of LINGO-1 or TrkB or fragment or variant described above with an affinity characterized by a dissociation constant K_D of less than about 5×10^{-2} M, about 10^{-2} M, about 5×10^{-3} M, about 10^{-3} M, about 5×10^{-4} M, about 10^{-4} M, about 5×10^{-5} M, about 10^{-5} M, about 5×10^{-6} M, about 10^{-6} M, about 5×10^{-7} M, about 10^{-7} M, about 5×10^{-8} M, about 10^{-8} M, about 5×10^{-9} M, about 10^{-9} M, about 5×10^{-10} M, about 10^{-10} M, about 5×10^{-11} M, about 10^{-11} M, about 5×10^{-12} M, about 10^{-12} M, about 5×10^{-13} M, about 10^{-13} M, about 5×10^{-14} M, about 10^{-14} M, about 5×10^{-15} M, or about 10^{-15} M. In a particular aspect, the antibody or fragment thereof preferentially binds to a human LINGO-1 or TrkB polypeptide or fragment thereof, relative to a murine LINGO-1 or TrkB polypeptide or fragment thereof.

[0139] As used in the context of antibody binding dissociation constants, the term "about" allows for the degree of variation inherent in the methods utilized for measuring antibody affinity. For example, depending on the level of precision of the instrumentation used, standard error based on the number of samples measured, and rounding error, the term "about 10^{-2} M" might include, for example, from 0.05 M to 0.005 M.

- [0140] In specific embodiments, the LINGO-1 antagonist or TrkB agonists for use in the methods of the present invention include an antibody, or antigen-binding fragment, variant, or derivative thereof of the invention binds LINGO-1 or TrkB polypeptides or fragments or variants thereof with an off rate ($k(\text{off})$) of less than or equal to $5 \times 10^{-2} \text{ sec}^{-1}$, 10^{-2} sec^{-1} , $5 \times 10^{-3} \text{ sec}^{-1}$ or 10^{-3} sec^{-1} . Alternatively, an antibody, or antigen-binding fragment, variant, or derivative thereof of the invention binds LINGO-1 or TrkB polypeptides or fragments or variants thereof with an off rate ($k(\text{off})$) of less than or equal to $5 \times 10^{-4} \text{ sec}^{-1}$, 10^{-4} sec^{-1} , $5 \times 10^{-5} \text{ sec}^{-1}$, or 10^{-5} sec^{-1} , $5 \times 10^{-6} \text{ sec}^{-1}$, 10^{-6} sec^{-1} , $5 \times 10^{-7} \text{ sec}^{-1}$ or 10^{-7} sec^{-1} .
- [0141] In other embodiments, the LINGO-1 antagonist or TrkB agonists for use in the methods of the present invention include an antibody, or antigen-binding fragment, variant, or derivative thereof of the invention binds LINGO-1 or TrkB polypeptides or fragments or variants thereof with an on rate ($k(\text{on})$) of greater than or equal to $10^3 \text{ M}^{-1} \text{ sec}^{-1}$, $5 \times 10^3 \text{ M}^{-1} \text{ sec}^{-1}$, $10^4 \text{ M}^{-1} \text{ sec}^{-1}$ or $5 \times 10^4 \text{ M}^{-1} \text{ sec}^{-1}$. Alternatively, an antibody, or antigen-binding fragment, variant, or derivative thereof of the invention binds LINGO-1 or TrkB polypeptides or fragments or variants thereof with an on rate ($k(\text{on})$) greater than or equal to $10^5 \text{ M}^{-1} \text{ sec}^{-1}$, $5 \times 10^5 \text{ M}^{-1} \text{ sec}^{-1}$, $10^6 \text{ M}^{-1} \text{ sec}^{-1}$, or $5 \times 10^6 \text{ M}^{-1} \text{ sec}^{-1}$ or $10^7 \text{ M}^{-1} \text{ sec}^{-1}$.
- [0142] In one embodiment, the LINGO-1 antagonist or TrkB agonist antibody for use in the methods of the invention an antibody molecule or an immunospecific fragment thereof. Unless it is specifically noted, as used herein, a "fragment thereof" in reference to an antibody refers to an immunospecific fragment, *i.e.*, an antigen-specific fragment. In one embodiment, an antibody for use in the methods of the invention is a bispecific binding molecule, binding polypeptide, or antibody, *e.g.*, a bispecific antibody, minibody, domain deleted antibody, or fusion protein having binding specificity for more than one epitope, *e.g.*, more than one antigen or more than one epitope on the same antigen. In one embodiment, a bispecific antibody has at least one binding domain specific for at least one epitope on LINGO-1 or TrkB. A bispecific antibody may be a tetravalent antibody that has two target binding domains specific for an epitope of LINGO-1 or TrkB and two target binding domains specific for a second target. Thus, a tetravalent bispecific antibody may be bivalent for each specificity.
- [0143] Certain methods of the present invention comprise administration of a LINGO-1 antagonist or TrkB agonist antibody, or immunospecific fragment thereof, in which at least a fraction of one or more of the constant region domains has been deleted or

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otherwise altered so as to provide desired biochemical characteristics such as reduced effector functions, the ability to non-covalently dimerize, increased ability to localize to the desired cite, reduced serum half-life, or increased serum half-life when compared with a whole, unaltered antibody of approximately the same immunogenicity. For example, certain antibodies for use in the methods described herein are domain deleted antibodies which comprise a polypeptide chain similar to an immunoglobulin heavy chain, but which lack at least a portion of one or more heavy chain domains. For instance, in certain antibodies, one entire domain of the constant region of the modified antibody will be deleted, for example, all or part of the C_H2 domain will be deleted.

[0144] In certain LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods described herein, the Fc portion may be mutated to decrease effector function using techniques known in the art. For example, the deletion or inactivation (through point mutations or other means) of a constant region domain may reduce Fc receptor binding of the circulating modified antibody thereby increasing localization at the desired cite of action. In other cases it may be that constant region modifications consistent with the instant invention moderate complement binding and thus reduce the serum half life and nonspecific association of a conjugated cytotoxin. Other desirable modifications to the constant region include mutations designed to alter glycosylation, for example to reduce glycosylation, in order to affect the efficacy, safety, antigen binding, Fc effector fuctions, stability and/or antibody product consistency. Further desirable modifications include modifications that alter antibody solubility, for example by modification of residues that are known or predicted to be involved in multimer formation. Yet other modifications of the constant region may be used to modify disulfide linkages or oligosaccharide moieties that allow for enhanced localization due to increased antigen specificity or antibody flexibility. The resulting physiological profile, bioavailability and other biochemical effects of the modifications, such as localization, biodistribution and serum half-life, may easily be measured and quantified using well-known immunological techniques without undue experimentation.

[0145] Modified forms of antibodies or immunospecific fragments thereof for use in the methods disclosed herein can be made from whole precursor or parent antibodies using techniques known in the art. Exemplary techniques are discussed in more detail herein.

[0146] In certain embodiments both the variable and constant regions of LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods disclosed herein are fully human. Fully human antibodies can be made using

techniques that are known in the art and as described herein. For example, fully human antibodies against a specific antigen can be prepared by administering the antigen to a transgenic animal which has been modified to produce such antibodies in response to antigenic challenge, but whose endogenous loci have been disabled. Exemplary techniques that can be used to make such antibodies are described in US patents: 6,150,584; 6,458,592; 6,420,140. Other techniques are known in the art. Fully human antibodies can likewise be produced by various display technologies, *e.g.*, phage display or other viral display systems, as described in more detail elsewhere herein.

[0147] LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods disclosed herein can be made or manufactured using techniques that are known in the art. In certain embodiments, antibody molecules or fragments thereof are “recombinantly produced,” *i.e.*, are produced using recombinant DNA technology. Exemplary techniques for making antibody molecules or fragments thereof are discussed in more detail elsewhere herein.

[0148] LINGO-1 antagonist and TrkB agonist antibodies or fragments thereof for use in the methods of the present invention may be generated by any suitable method known in the art.

[0149] Polyclonal antibodies can be produced by various procedures well known in the art. For example, a LINGO-1 or TrkB immunospecific fragment can be administered to various host animals including, but not limited to, rabbits, mice, rats, etc. to induce the production of sera containing polyclonal antibodies specific for the antigen. Various adjuvants may be used to increase the immunological response, depending on the host species, and include but are not limited to, Freund's (complete and incomplete), mineral gels such as aluminum hydroxide, surface active substances such as lysolecithin, pluronic polyols, polyanions, peptides, oil emulsions, keyhole limpet hemocyanins, dinitrophenol, and potentially useful human adjuvants such as BCG (bacille Calmette-Guerin) and *Corynebacterium parvum*. Such adjuvants are also well known in the art.

[0150] Monoclonal antibodies can be prepared using a wide variety of techniques known in the art including the use of hybridoma, recombinant, and phage display technologies, or a combination thereof. For example, monoclonal antibodies can be produced using hybridoma techniques including those known in the art and taught, for example, in Harlow *et al.*, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory Press, 2nd ed. (1988); Hammerling *et al.*, in: *Monoclonal Antibodies and T-Cell Hybridomas* Elsevier, N.Y., 563-681 (1981) (said references incorporated by reference in their

entireties). The term "monoclonal antibody" as used herein is not limited to antibodies produced through hybridoma technology. The term "monoclonal antibody" refers to an antibody that is derived from a single clone, including any eukaryotic, prokaryotic, or phage clone, and not the method by which it is produced. Thus, the term "monoclonal antibody" is not limited to antibodies produced through hybridoma technology. Monoclonal antibodies can be prepared using a wide variety of techniques known in the art including the use of hybridoma and recombinant and phage display technology.

[0151] Using art recognized protocols, in one example, antibodies are raised in mammals by multiple subcutaneous or intraperitoneal injections of the relevant antigen (*e.g.*, purified LINGO-1 or TrkB antigens or cells or cellular extracts comprising such antigens) and an adjuvant. This immunization typically elicits an immune response that comprises production of antigen-reactive antibodies from activated splenocytes or lymphocytes. While the resulting antibodies may be harvested from the serum of the animal to provide polyclonal preparations, it is often desirable to isolate individual lymphocytes from the spleen, lymph nodes or peripheral blood to provide homogenous preparations of monoclonal antibodies (mAbs). The lymphocytes can be obtained, for example, from the spleen.

[0152] In this well known process (Kohler *et al.*, *Nature* 256:495 (1975)) the relatively short-lived, or mortal, lymphocytes from a mammal which has been injected with antigen are fused with an immortal tumor cell line (*e.g.* a myeloma cell line), thus producing hybrid cells or "hybridomas" which are both immortal and capable of producing the genetically coded antibody of the B cell. The resulting hybrids are segregated into single genetic strains by selection, dilution, and regrowth with each individual strain comprising specific genes for the formation of a single antibody. They produce antibodies which are homogeneous against a desired antigen and, in reference to their pure genetic parentage, are termed "monoclonal."

[0153] Typically, hybridoma cells thus prepared are seeded and grown in a suitable culture medium that can contains one or more substances that inhibit the growth or survival of the unfused, parental myeloma cells. Those skilled in the art will appreciate that reagents, cell lines and media for the formation, selection and growth of hybridomas are commercially available from a number of sources and standardized protocols are well established. Generally, culture medium in which the hybridoma cells are growing is assayed for production of monoclonal antibodies against the desired antigen. The binding specificity of the monoclonal antibodies produced by hybridoma cells can be determined

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by *in vitro* assays such as immunoprecipitation, radioimmunoassay (RIA) or enzyme-linked immunoabsorbent assay (ELISA). After hybridoma cells are identified that produce antibodies of the desired specificity, affinity and/or activity, the clones may be subcloned by limiting dilution procedures and grown by standard methods (Goding, *Monoclonal Antibodies: Principles and Practice*, Academic Press, pp 59-103 (1986)). It will further be appreciated that the monoclonal antibodies secreted by the subclones may be separated from culture medium, ascites fluid or serum by conventional purification procedures such as, for example, protein-A, hydroxylapatite chromatography, gel electrophoresis, dialysis or affinity chromatography.

[0154] Antibody fragments that recognize specific epitopes may be generated by known techniques. For example, Fab and F(ab')₂ fragments may be produced by proteolytic cleavage of immunoglobulin molecules, using enzymes such as papain (to produce Fab fragments) or pepsin (to produce F(ab')₂ fragments). F(ab')₂ fragments contain the variable region, the light chain constant region and the C_H1 domain of the heavy chain.

[0155] Those skilled in the art will also appreciate that DNA encoding antibodies or antibody fragments (*e.g.*, antigen binding sites) may also be derived from antibody phage libraries. In a particular, such phage can be utilized to display antigen-binding domains expressed from a repertoire or combinatorial antibody library (*e.g.*, human or murine). Phage expressing an antigen binding domain that binds the antigen of interest can be selected or identified with antigen, *e.g.*, using labeled antigen or antigen bound or captured to a solid surface or bead. Phage used in these methods are typically filamentous phage including fd and M13 binding domains expressed from phage with Fab, Fv or disulfide stabilized Fv antibody domains recombinantly fused to either the phage gene III or gene VIII protein. Exemplary methods are set forth, for example, in EP 368 684 B1; U.S. patent. 5,969,108, Hoogenboom, H.R. and Chames, *Immunol. Today* 21:371 (2000); Nagy *et al. Nat. Med.* 8:801 (2002); Huie *et al., Proc. Natl. Acad. Sci. USA* 98:2682 (2001); Lui *et al., J. Mol. Biol.* 315:1063 (2002), each of which is incorporated herein by reference. Several publications (*e.g.*, Marks *et al., Bio/Technology* 10:779-783 (1992)) have described the production of high affinity human antibodies by chain shuffling, as well as combinatorial infection and *in vivo* recombination as a strategy for constructing large phage libraries. In another embodiment, ribosomal display can be used to replace bacteriophage as the display platform (*see, e.g.*, Hanes *et al., Nat. Biotechnol.* 18:1287 (2000); Wilson *et al., Proc. Natl. Acad. Sci. USA* 98:3750 (2001); or Irving *et al., J. Immunol. Methods* 248:31 (2001)). In yet another embodiment, cell surface libraries can

be screened for antibodies (Boder *et al.*, *Proc. Natl. Acad. Sci. USA* 97:10701 (2000); Daugherty *et al.*, *J. Immunol. Methods* 243:211 (2000)). Such procedures provide alternatives to traditional hybridoma techniques for the isolation and subsequent cloning of monoclonal antibodies.

[0156] In phage display methods, functional antibody domains are displayed on the surface of phage particles which carry the polynucleotide sequences encoding them. In particular, DNA sequences encoding V_H and V_L regions are amplified from animal cDNA libraries (*e.g.*, human or murine cDNA libraries of lymphoid tissues) or synthetic cDNA libraries. In certain embodiments, the DNA encoding the V_H and V_L regions are joined together by an scFv linker by PCR and cloned into a phagemid vector (*e.g.*, p CANTAB 6 or pComb 3 HSS). The vector is electroporated in *E. coli* and the *E. coli* is infected with helper phage. Phage used in these methods are typically filamentous phage including fd and M13 and the V_H or V_L regions are usually recombinantly fused to either the phage gene III or gene VIII. Phage expressing an antigen binding domain that binds to an antigen of interest (*i.e.*, a LINGO-1 or TrkB polypeptide or a fragment thereof) can be selected or identified with antigen, *e.g.*, using labeled antigen or antigen bound or captured to a solid surface or bead.

[0157] Additional examples of phage display methods that can be used to make the antibodies include those disclosed in Brinkman *et al.*, *J. Immunol. Methods* 182:41-50 (1995); Ames *et al.*, *J. Immunol. Methods* 184:177-186 (1995); Kettleborough *et al.*, *Eur. J. Immunol.* 24:952-958 (1994); Persic *et al.*, *Gene* 187:9-18 (1997); Burton *et al.*, *Advances in Immunology* 57:191-280 (1994); PCT Application No. PCT/GB91/01134; PCT publications WO 90/02809; WO 91/10737; WO 92/01047; WO 92/18619; WO 93/11236; WO 95/15982; WO 95/20401; and U.S. Pat. Nos. 5,698,426; 5,223,409; 5,403,484; 5,580,717; 5,427,908; 5,750,753; 5,821,047; 5,571,698; 5,427,908; 5,516,637; 5,780,225; 5,658,727; 5,733,743 and 5,969,108; each of which is incorporated herein by reference in its entirety.

[0158] As described in the above references, after phage selection, the antibody coding regions from the phage can be isolated and used to generate whole antibodies, including human antibodies, or any other desired antigen binding fragment, and expressed in any desired host, including mammalian cells, insect cells, plant cells, yeast, and bacteria. For example, techniques to recombinantly produce Fab, Fab' and F(ab')₂ fragments can also be employed using methods known in the art such as those disclosed in PCT publication WO 92/22324; Mullinax *et al.*, *BioTechniques* 12(6):864-869 (1992); and Sawai *et al.*,

AJRI 34:26-34 (1995); and Better *et al.*, *Science* 240:1041-1043 (1988) (said references incorporated by reference in their entireties).

[0159] In another embodiment, DNA encoding desired monoclonal antibodies for use in the methods of the present invention may be readily isolated and sequenced using conventional procedures (*e.g.*, by using oligonucleotide probes that are capable of binding specifically to genes encoding the heavy and light chains of murine antibodies). The isolated and subcloned hybridoma cells serve as a possible source of such DNA. Once isolated, the DNA may be placed into expression vectors, which are then transfected into prokaryotic or eukaryotic host cells such as *E. coli* cells, simian COS cells, Chinese Hamster Ovary (CHO) cells or myeloma cells that do not otherwise produce immunoglobulins. More particularly, the isolated DNA (which may be synthetic as described herein) may be used to clone constant and variable region sequences for the manufacture antibodies as described in Newman *et al.*, U.S. Pat. No. 5,658,570, filed January 25, 1995, which is incorporated by reference herein. Essentially, this entails extraction of RNA from the selected cells, conversion to cDNA, and amplification by PCR using Ig specific primers. Suitable primers for this purpose are also described in U.S. Pat. No. 5,658,570. As will be discussed in more detail below, transformed cells expressing the desired antibody may be grown up in relatively large quantities to provide clinical and commercial supplies of the immunoglobulin.

[0160] In a specific embodiment, the amino acid sequence of the heavy and/or light chain variable domains may be inspected to identify the sequences of the complementarity determining regions (CDRs) by methods that are well known in the art, *e.g.*, by comparison to known amino acid sequences of other heavy and light chain variable regions to determine the regions of sequence hypervariability. Using routine recombinant DNA techniques, one or more of the CDRs may be inserted within framework regions, *e.g.*, into human framework regions to humanize a non-human antibody. The framework regions may be naturally occurring or consensus framework regions, and may be human framework regions (see, *e.g.*, Chothia *et al.*, *J. Mol. Biol.* 278:457-479 (1998) for a listing of human framework regions). The polynucleotide generated by the combination of the framework regions and CDRs may encode an antibody that specifically binds to at least one epitope of a desired polypeptide, *e.g.*, LINGO-1. One or more amino acid substitutions may be made within the framework regions, and the amino acid substitutions may improve binding of the antibody to its antigen. Additionally, such methods may be used to make amino acid substitutions or deletions of one or more

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variable region cysteine residues participating in an intrachain disulfide bond to generate antibody molecules lacking one or more intrachain disulfide bonds. Other alterations to the polynucleotide are encompassed by the present invention and within the skill of the art.

[0161] In certain embodiments, a LINGO-1 antagonist or TrkB agonist antibody or immunospecific fragment thereof for use in the methods disclosed herein will not elicit a deleterious immune response in the animal to be treated, *e.g.*, in a human. In one embodiment, LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods disclosed herein can be modified to reduce their immunogenicity using art-recognized techniques. For example, antibodies can be humanized, primatized, deimmunized, or chimeric antibodies can be made. These types of antibodies are derived from a non-human antibody, typically a murine or primate antibody, that retains or substantially retains the antigen-binding properties of the parent antibody, but which is less immunogenic in humans. This may be achieved by various methods, including (a) grafting the entire non-human variable domains onto human constant regions to generate chimeric antibodies; (b) grafting at least a part of one or more of the non-human complementarity determining regions (CDRs) into a human framework and constant regions with or without retention of critical framework residues; or (c) transplanting the entire non-human variable domains, but "cloaking" them with a human-like section by replacement of surface residues. Such methods are disclosed in Morrison *et al.*, *Proc. Natl. Acad. Sci.* 81:6851-6855 (1984); Morrison *et al.*, *Adv. Immunol.* 44:65-92 (1988); Verhoeven *et al.*, *Science* 239:1534-1536 (1988); Padlan, *Molec. Immun.* 28:489-498 (1991); Padlan, *Molec. Immun.* 31:169-217 (1994), and U.S. Pat. Nos. 5,585,089, 5,693,761, 5,693,762, and 6,190,370, all of which are hereby incorporated by reference in their entirety.

[0162] De-immunization can also be used to decrease the immunogenicity of an antibody. As used herein, the term "de-immunization" includes alteration of an antibody to modify T cell epitopes (see, *e.g.*, WO9852976A1, WO0034317A2). For example, V_H and V_L sequences from the starting antibody are analyzed and a human T cell epitope "map" from each V region showing the location of epitopes in relation to complementarity-determining regions (CDRs) and other key residues within the sequence. Individual T cell epitopes from the T cell epitope map are analyzed in order to identify alternative amino acid substitutions with a low risk of altering activity of the final antibody. A range of alternative V_H and V_L sequences are designed comprising combinations of amino acid

substitutions and these sequences are subsequently incorporated into a range of binding polypeptides, *e.g.*, LINGO-1 antagonist or TrkB agonist antibodies or immunospecific fragments thereof for use in the methods disclosed herein, which are then tested for function. Typically, between 12 and 24 variant antibodies are generated and tested. Complete heavy and light chain genes comprising modified V and human C regions are then cloned into expression vectors and the subsequent plasmids introduced into cell lines for the production of whole antibody. The antibodies are then compared in appropriate biochemical and biological assays, and the optimal variant is identified.

[0163] A chimeric antibody is a molecule in which different portions of the antibody are derived from different animal species, such as antibodies having a variable region derived from a murine monoclonal antibody and a human immunoglobulin constant region. Methods for producing chimeric antibodies are known in the art. *See, e.g.*, Morrison, *Science* 229:1202 (1985); Oi *et al.*, *BioTechniques* 4:214 (1986); Gillies *et al.*, *J. Immunol. Methods* 125:191-202 (1989); U.S. Pat. Nos. 5,807,715; 4,816,567; and 4,816,397, which are incorporated herein by reference in their entireties. Humanized antibodies are antibody molecules from non-human species antibody that binds the desired antigen having one or more complementarity determining regions (CDRs) from the non-human species and framework regions from a human immunoglobulin molecule. Often, framework residues in the human framework regions will be substituted with the corresponding residue from the CDR donor antibody to alter, and potentially improve, antigen binding. These framework substitutions are identified by methods well known in the art, *e.g.*, by modeling of the interactions of the CDR and framework residues to identify framework residues important for antigen binding and sequence comparison to identify unusual framework residues at particular positions. (See, *e.g.*, Queen *et al.*, U.S. Pat. No. 5,585,089; Riechmann *et al.*, *Nature* 332:323 (1988), which are incorporated herein by reference in their entireties.) Antibodies can be humanized using a variety of techniques known in the art including, for example, CDR-grafting (EP 239,400; PCT publication WO 91/09967; U.S. Pat. Nos. 5,225,539; 5,530,101; and 5,585,089), veneering or resurfacing (EP 592,106; EP 519,596; Padlan, *Molecular Immunology* 28(4/5):489-498 (1991); Studnicka *et al.*, *Protein Engineering* 7(6):805-814 (1994); Roguska. *et al.*, *PNAS* 91:969-973 (1994)), and chain shuffling (U.S. Pat. No. 5,565,332).

[0164] Yet another highly efficient means for generating recombinant antibodies is disclosed by Newman, *Biotechnology* 10: 1455-1460 (1992). Specifically, this technique results in the generation of primatized antibodies that contain monkey variable domains

and human constant sequences. This reference is incorporated by reference in its entirety herein. Moreover, this technique is also described in commonly assigned U.S. Pat. Nos. 5,658,570, 5,693,780 and 5,756,096 each of which is incorporated herein by reference.

[0165] Completely human antibodies are particularly desirable for therapeutic treatment of human patients. Human antibodies can be made by a variety of methods known in the art including phage display methods described above using antibody libraries derived from human immunoglobulin sequences. See also, U.S. Pat. Nos. 4,444,887 and 4,716,111; and PCT publications WO 98/46645, WO 98/50433, WO 98/24893, WO 98/16654, WO 96/34096, WO 96/33735, and WO 91/10741; each of which is incorporated herein by reference in its entirety.

[0166] Human antibodies can also be produced using transgenic mice which are incapable of expressing functional endogenous immunoglobulins, but which can express human immunoglobulin genes. For example, the human heavy and light chain immunoglobulin gene complexes may be introduced randomly or by homologous recombination into mouse embryonic stem cells. Alternatively, the human variable region, constant region, and diversity region may be introduced into mouse embryonic stem cells in addition to the human heavy and light chain genes. The mouse heavy and light chain immunoglobulin genes may be rendered non-functional separately or simultaneously with the introduction of human immunoglobulin loci by homologous recombination. In particular, homozygous deletion of the JH region prevents endogenous antibody production. The modified embryonic stem cells are expanded and microinjected into blastocysts to produce chimeric mice. The chimeric mice are then bred to produce homozygous offspring that express human antibodies. The transgenic mice are immunized in the normal fashion with a selected antigen, *e.g.*, all or a portion of a desired target polypeptide. Monoclonal antibodies directed against the antigen can be obtained from the immunized, transgenic mice using conventional hybridoma technology. The human immunoglobulin transgenes harbored by the transgenic mice rearrange during B-cell differentiation, and subsequently undergo class switching and somatic mutation. Thus, using such a technique, it is possible to produce therapeutically useful IgG, IgA, IgM and IgE antibodies. For an overview of this technology for producing human antibodies, see Lonberg and Huszar *Int. Rev. Immunol.* 13:65-93 (1995). For a detailed discussion of this technology for producing human antibodies and human monoclonal antibodies and protocols for producing such antibodies, see, *e.g.*, PCT publications WO 98/24893; WO 96/34096; WO 96/33735; U.S. Pat. Nos. 5,413,923; 5,625,126; 5,633,425;

5,569,825; 5,661,016; 5,545,806; 5,814,318; and 5,939,598, which are incorporated by reference herein in their entirety. In addition, companies such as Abgenix, Inc. (Freemont, Calif.) and GenPharm (San Jose, Calif.) can be engaged to provide human antibodies directed against a selected antigen using technology similar to that described above.

[0167] Another means of generating human antibodies using SCID mice is disclosed in U.S. Pat. No. 5,811,524 which is incorporated herein by reference. It will be appreciated that the genetic material associated with these human antibodies may also be isolated and manipulated as described herein.

[0168] Completely human antibodies which recognize a selected epitope can be generated using a technique referred to as "guided selection." In this approach a selected non-human monoclonal antibody, *e.g.*, a mouse antibody, is used to guide the selection of a completely human antibody recognizing the same epitope. (Jespers *et al.*, *Bio/Technology* 12:899-903 (1988)). *See also*, U.S. Patent No. 5,565,332.

[0169] Alternatively, techniques described for the production of single chain antibodies (U.S. Pat. No. 4,694,778; Bird, *Science* 242:423-442 (1988); Huston *et al.*, *Proc. Natl. Acad. Sci. USA* 85:5879-5883 (1988); and Ward *et al.*, *Nature* 334:544-554 (1989)) can be adapted to produce single chain antibodies. Single chain antibodies are formed by linking the heavy and light chain fragments of the Fv region via an amino acid bridge, resulting in a single chain antibody. Techniques for the assembly of functional Fv fragments in *E coli* may also be used (Skerra *et al.*, *Science* 242:1038-1041 (1988)). Examples of techniques which can be used to produce single-chain Fvs and antibodies include those described in U.S. Pat. Nos. 4,946,778 and 5,258,498; Huston *et al.*, *Methods in Enzymology* 203:46-88 (1991); Shu *et al.*, *PNAS* 90:7995-7999 (1993); and Skerra *et al.*, *Science* 240:1038-1040 (1988). For some uses, especially including *in vivo* use of antibodies in humans and *in vitro* detection assays, chimeric, humanized, or human antibodies can be used.

[0170] In addition, techniques developed for the production of "chimeric antibodies" (Morrison *et al.*, *Proc. Natl. Acad. Sci.* 81:851-855 (1984); Neuberger *et al.*, *Nature* 312:604-608 (1984); Takeda *et al.*, *Nature* 314:452-454 (1985)) by splicing genes from a mouse antibody molecule of appropriate antigen specificity together with genes from a human antibody molecule of appropriate biological activity can be used. As used herein, a chimeric antibody is a molecule in which different portions are derived from different animal species, such as those having a variable region derived from a murine monoclonal antibody and a human immunoglobulin constant region, *e.g.*, humanized antibodies.

[0171] LINGO-1 antagonist or TrkB agonist antibodies may also be human or substantially human antibodies generated in transgenic animals (e.g., mice) that are incapable of endogenous immunoglobulin production (see e.g., U.S. Pat. Nos. 6,075,181, 5,939,598, 5,591,669 and 5,589,369 each of which is incorporated herein by reference). For example, it has been described that the homozygous deletion of the antibody heavy-chain joining region in chimeric and germ-line mutant mice results in complete inhibition of endogenous antibody production. Transfer of a human immunoglobulin gene array to such germ line mutant mice will result in the production of human antibodies upon antigen challenge.

[0172] In another embodiment, lymphocytes can be selected by micromanipulation and the variable genes isolated. For example, peripheral blood mononuclear cells can be isolated from an immunized mammal and cultured for about 7 days *in vitro*. The cultures can be screened for specific IgGs that meet the screening criteria. Cells from positive wells can be isolated. Individual Ig-producing B cells can be isolated by FACS or by identifying them in a complement-mediated hemolytic plaque assay. Ig-producing B cells can be micromanipulated into a tube and the V_H and V_L genes can be amplified using, e.g., RT-PCR. The V_H and V_L genes can be cloned into an antibody expression vector and transfected into cells (e.g., eukaryotic or prokaryotic cells) for expression.

[0173] Alternatively, antibody-producing cell lines may be selected and cultured using techniques well known to the skilled artisan. Such techniques are described in a variety of laboratory manuals and primary publications. In this respect, techniques suitable for use in the invention as described below are described in *Current Protocols in Immunology*, Coligan *et al.*, Eds., Green Publishing Associates and Wiley-Interscience, John Wiley and Sons, New York (1991) which is herein incorporated by reference in its entirety, including supplements.

[0174] Antibodies for use in the methods disclosed herein can be produced by any method known in the art for the synthesis of antibodies, in particular, by chemical synthesis or by recombinant expression techniques as described herein.

[0175] It will further be appreciated that the scope of this invention further encompasses all alleles, variants and mutations of antigen binding DNA sequences.

[0176] As is well known, RNA may be isolated from the original hybridoma cells or from other transformed cells by standard techniques, such as guanidinium isothiocyanate extraction and precipitation followed by centrifugation or chromatography. Where

desirable, mRNA may be isolated from total RNA by standard techniques such as chromatography on oligo dT cellulose. Suitable techniques are familiar in the art.

[0177] In one embodiment, cDNAs that encode the light and the heavy chains of the antibody for use in the methods of the present invention may be made, either simultaneously or separately, using reverse transcriptase and DNA polymerase in accordance with well known methods. PCR may be initiated by consensus constant region primers or by more specific primers based on the published heavy and light chain DNA and amino acid sequences. As discussed above, PCR also may be used to isolate DNA clones encoding the antibody light and heavy chains. In this case the libraries may be screened by consensus primers or larger homologous probes, such as mouse constant region probes.

[0178] DNA, typically plasmid DNA, may be isolated from the cells using techniques known in the art, restriction mapped and sequenced in accordance with standard, well known techniques set forth in detail, *e.g.*, in the foregoing references relating to recombinant DNA techniques. Of course, the DNA may be synthetic according to the present invention at any point during the isolation process or subsequent analysis.

[0179] Recombinant expression of an antibody, or fragment, derivative or analog thereof, *e.g.*, a heavy or light chain of an antibody which is a LINGO-1 antagonist or TrkB agonist, requires construction of an expression vector containing a polynucleotide that encodes the antibody. Once a polynucleotide encoding an antibody molecule or a heavy or light chain of an antibody, or portion thereof (which may contain the heavy or light chain variable domain), of the invention has been obtained, the vector for the production of the antibody molecule may be produced by recombinant DNA technology using techniques well known in the art. Thus, methods for preparing a protein by expressing a polynucleotide containing an antibody encoding nucleotide sequence are described herein. Methods which are well known to those skilled in the art can be used to construct expression vectors containing antibody coding sequences and appropriate transcriptional and translational control signals. These methods include, for example, *in vitro* recombinant DNA techniques, synthetic techniques, and *in vivo* genetic recombination. The invention, thus, provides replicable vectors comprising a nucleotide sequence encoding an antibody molecule of the invention, or a heavy or light chain thereof, or a heavy or light chain variable domain, operably linked to a promoter. Such vectors may include the nucleotide sequence encoding the constant region of the antibody molecule (see, *e.g.*, PCT Publication WO 86/05807; PCT Publication WO 89/01036; and U.S. Pat.

No. 5,122,464) and the variable domain of the antibody may be cloned into such a vector for expression of the entire heavy or light chain.

[0180] The expression vector is transferred to a host cell by conventional techniques and the transfected cells are then cultured by conventional techniques to produce an antibody for use in the methods described herein. Thus, the invention includes host cells containing a polynucleotide encoding an antibody of the invention, or a heavy or light chain thereof, operably linked to a heterologous promoter. In certain embodiments for the expression of double-chained antibodies, vectors encoding both the heavy and light chains may be co-expressed in the host cell for expression of the entire immunoglobulin molecule, as detailed below.

[0181] A variety of host-expression vector systems may be utilized to express antibody molecules for use in the methods described elsewhere herein.

[0182] The host cell may be co-transfected with two expression vectors of the invention, the first vector encoding a heavy chain derived polypeptide and the second vector encoding a light chain derived polypeptide. The two vectors may contain identical selectable markers which enable equal expression of heavy and light chain polypeptides. Alternatively, a single vector may be used which encodes both heavy and light chain polypeptides. In such situations, the light chain is advantageously placed before the heavy chain to avoid an excess of toxic free heavy chain (Proudfoot, *Nature* 322:52 (1986); Kohler, *Proc. Natl. Acad. Sci. USA* 77:2197 (1980)). The coding sequences for the heavy and light chains may comprise cDNA or genomic DNA.

[0183] Once an antibody molecule of the invention has been recombinantly expressed, it may be purified by any method known in the art for purification of an immunoglobulin molecule, for example, by chromatography (*e.g.*, ion exchange, affinity, particularly by affinity for the specific antigen after Protein A, and sizing column chromatography), centrifugation, differential solubility, or by any other standard technique for the purification of proteins. Alternatively, another method for increasing the affinity of antibodies of the invention is disclosed in US 2002 0123057 A1, which is incorporated herein by reference.

[0184] In one embodiment, a binding molecule or antigen binding molecule for use in the methods of the invention comprises a synthetic constant region wherein one or more domains are partially or entirely deleted ("domain-deleted antibodies"). In certain embodiments compatible modified antibodies will comprise domain deleted constructs or variants wherein the entire C_H2 domain has been removed (Δ C_H2 constructs). For other

embodiments a short connecting peptide may be substituted for the deleted domain to provide flexibility and freedom of movement for the variable region. Those skilled in the art will appreciate that such constructs are particularly useful due to the regulatory properties of the C_H2 domain on the catabolic rate of the antibody.

[0185] In certain embodiments, modified antibodies for use in the methods disclosed herein are minibodies. Minibodies can be made using methods described in the art (*see, e.g.*, US patent 5,837,821 or WO 94/09817A1).

[0186] In another embodiment, modified antibodies for use in the methods disclosed herein are C_H2 domain deleted antibodies which are known in the art. Domain deleted constructs can be derived using a vector (*e.g.*, from Biogen IDEC Incorporated) encoding an IgG₁ human constant domain (*see, e.g.*, WO 02/060955A2 and WO02/096948A2, which are incorporated herein by reference). This exemplary vector was engineered to delete the C_H2 domain and provide a synthetic vector expressing a domain deleted IgG₁ constant region.

[0187] In one embodiment, a LINGO-1 antagonist or TrkB agonist antibody or fragment thereof for use in the methods disclosed herein comprises an immunoglobulin heavy chain having deletion or substitution of a few or even a single amino acid as long as it permits association between the monomeric subunits. Similarly, it may be desirable to simply delete that part of one or more constant region domains that controls the effector function (*e.g.* complement binding) to be modulated. Such partial deletions of the constant regions may improve selected characteristics of the antibody (serum half-life) while leaving other desirable functions associated with the subject constant region domain intact. Moreover, as alluded to above, the constant regions of the disclosed antibodies may be synthetic through the mutation or substitution of one or more amino acids that enhances the profile of the resulting construct. In this respect it may be possible to disrupt the activity provided by a conserved binding site (*e.g.* Fc binding) while substantially maintaining the configuration and immunogenic profile of the modified antibody. Yet other embodiments comprise the addition of one or more amino acids to the constant region to enhance desirable characteristics such as effector function or provide for more cytotoxin or carbohydrate attachment. In such embodiments it may be desirable to insert or replicate specific sequences derived from selected constant region domains.

[0188] The present invention also provides the use of antibodies that comprise, consist essentially of, or consist of, variants (including derivatives) of antibody molecules (*e.g.*,

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the V_H regions and/or V_L regions) described herein, which antibodies or fragments thereof immunospecifically bind to a LINGO-1 or TrkB polypeptide. Standard techniques known to those of skill in the art can be used to introduce mutations in the nucleotide sequence encoding a binding molecule, including, but not limited to, site-directed mutagenesis and PCR-mediated mutagenesis which result in amino acid substitutions. The variants (including derivatives) can encode less than 50 amino acid substitutions, less than 40 amino acid substitutions, less than 30 amino acid substitutions, less than 25 amino acid substitutions, less than 20 amino acid substitutions, less than 15 amino acid substitutions, less than 10 amino acid substitutions, less than 5 amino acid substitutions, less than 4 amino acid substitutions, less than 3 amino acid substitutions, or less than 2 amino acid substitutions relative to the reference V_H region, V_HCDR1, V_HCDR2, V_HCDR3, V_L region, V_LCDR1, V_LCDR2, or V_LCDR3. A "conservative amino acid substitution" is one in which the amino acid residue is replaced with an amino acid residue having a side chain with a similar charge. Families of amino acid residues having side chains with similar charges have been defined in the art. These families include amino acids with basic side chains (*e.g.*, lysine, arginine, histidine), acidic side chains (*e.g.*, aspartic acid, glutamic acid), uncharged polar side chains (*e.g.*, glycine, asparagine, glutamine, serine, threonine, tyrosine, cysteine), nonpolar side chains (*e.g.*, alanine, valine, leucine, isoleucine, proline, phenylalanine, methionine, tryptophan), beta-branched side chains (*e.g.*, threonine, valine, isoleucine) and aromatic side chains (*e.g.*, tyrosine, phenylalanine, tryptophan, histidine). Alternatively, mutations can be introduced randomly along all or part of the coding sequence, such as by saturation mutagenesis, and the resultant mutants can be screened for biological activity to identify mutants that retain activity.

[0189] For example, it is possible to introduce mutations only in framework regions or only in CDR regions of an antibody molecule. Introduced mutations may be silent or neutral missense mutations, *i.e.*, have no, or little, effect on an antibody's ability to bind antigen. These types of mutations may be useful to optimize codon usage, or improve a hybridoma's antibody production. Alternatively, non-neutral missense mutations may alter an antibody's ability to bind antigen. The location of most silent and neutral missense mutations is likely to be in the framework regions, while the location of most non-neutral missense mutations is likely to be in CDR, though this is not an absolute requirement. One of skill in the art would be able to design and test mutant molecules with desired properties such as no alteration in antigen binding activity or alteration in

binding activity (e.g., improvements in antigen binding activity or change in antibody specificity). Following mutagenesis, the encoded protein may routinely be expressed and the functional and/or biological activity of the encoded protein can be determined using techniques described herein or by routinely modifying techniques known in the art.

Additional Antagonists and Agonists and Combinations Thereof for Use in the Methods of the Invention

[0190] In addition to the agonists and antagonists described previously, additional TrkB agonists for use in the methods of the present invention include any polypeptide, antibody, compound or nucleotide which would promote, increase or enhance the activity of TrkB. Such agonist include polypeptides, antibodies, compounds or nucleotides which interfere with or promote the binding of a TrkB ligand such as Brain-Derived Neurotrophic Factor (BDNF), to the TrkB receptor and as such are also considered BDNF agonists. Such molecules include but are not limited to antibodies which disrupt the interact between a ligand and TrkB, peptidomimetic agonists of TrkB and ligand analogs such as those described in 5,770,577, 6,077,829, 6,723,701, and 6,800,607, which are hereby incorporated by reference in their entirety.

[0191] The antagonists and agonists described herein for use in the methods of the present invention may be administered as compositions in various combinations. For example, various TrkB agonists may be used in combination with LINGO-1 antagonists. Compositions for use in the methods of the present invention may also include multiple TrkB agonists and/or LINGO-1 antagonists.

[0192] Additionally, compositions for use in the present invention may include other antagonists or agonists of proteins expressed in the CNS such as Nogo Receptor 1 (NgR1), LINGO-1 (LINGO-1), TAJ or Oligodendrocyte-myelin glycoprotein (OMgp). Antagonists of NgR1 are described in U.S. Publication Nos. 2002/0077295 and 2005/0271655 A1 and International Application Publication Nos. WO 01/51520, WO 03/031462, WO 2004/014311 and WO 2005/016955, which are hereby incorporated by reference in their entirety. Antagonists of LINGO-1 (LINGO-1) may be found in U.S. Publication No. 2006/0009388 A1 and International Publication No. WO 2004/085648, which are hereby incorporated by reference in their entirety. Examples of TAJ antagonists are described in U.S. Publication No. 2006/0058223 A1, which is hereby incorporated by reference in its entirety. OMgp antagonists are described in U.S. Provisional Application Nos. 60/730,357 and 60/735,170 which are hereby incorporated

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by reference in their entireties. Compositions for use in the methods of the present invention may also include any number and combination of TrkB, LINGO-1, NgR1, TAJ and OMgp antagonists.

Aptamers

[0193] In another embodiment, the LINGO-1 antagonist or TrkB agonist for use in the methods of the present invention is an aptamer. An aptamer can be a nucleotide or a polypeptide which has a unique sequence, has the property of binding specifically to a desired target (*e.g.* a polypeptide), and is a specific ligand of a given target. Nucleotide aptamers of the invention include double stranded DNA and single stranded RNA molecules that bind to LINGO-1 or TrkB.

[0194] Nucleic acid aptamers are selected using methods known in the art, for example via the Systematic Evolution of Ligands by Exponential Enrichment (SELEX) process. SELEX is a method for the *in vitro* evolution of nucleic acid molecules with highly specific binding to target molecules as described in *e.g.* U.S. Pat. Nos. 5,475,096, 5,580,737, 5,567,588, 5,707,796, 5,763,177, 6, 011,577, and 6,699,843, incorporated herein by reference in their entirety. Another screening method to identify aptamers is described in U.S. Pat. No. 5,270,163 (also incorporated herein by reference). The SELEX process is based on the capacity of nucleic acids for forming a variety of two- and three-dimensional structures, as well as the chemical versatility available within the nucleotide monomers to act as ligands (form specific binding pairs) with virtually any chemical compound, whether monomeric or polymeric, including other nucleic acid molecules and polypeptides. Molecules of any size or composition can serve as targets.

[0195] The SELEX method involves selection from a mixture of candidate oligonucleotides and step-wise iterations of binding, partitioning and amplification, using the same general selection scheme, to achieve desired binding affinity and selectivity. Starting from a mixture of nucleic acids, which can comprise a segment of randomized sequence, the SELEX method includes steps of contacting the mixture with the target under conditions favorable for binding; partitioning unbound nucleic acids from those nucleic acids which have bound specifically to target molecules; dissociating the nucleic acid-target complexes; amplifying the nucleic acids dissociated from the nucleic acid-target complexes to yield a ligand enriched mixture of nucleic acids. The steps of binding, partitioning, dissociating and amplifying are repeated through as many cycles as desired to yield highly specific high affinity nucleic acid ligands to the target molecule.

- [0196] Nucleotide aptamers may be used, for example, as diagnostic tools or as specific inhibitors to dissect intracellular signaling and transport pathways. See James *Curr. Opin. Pharmacol.* 1:540-546 (2001). The high affinity and specificity of nucleotide aptamers makes them good candidates for drug discovery. For example, aptamer antagonists to the toxin ricin have been isolated and have IC₅₀ values in the nanomolar range. See Hesselberth JR *et al. J Biol Chem* 275:4937-4942 (2000). Nucleotide aptamers may also be used against infectious disease, malignancy and viral surface proteins to reduce cellular infectivity.
- [0197] Nucleotide aptamers for use in the methods of the present invention may be modified (*e.g.*, by modifying the backbone or bases or conjugated to peptides) as described herein for other polynucleotides.
- [0198] Using the protein structure of LINGO-1 or TrkB, screening for aptamers that act on LINGO-1 or TrkB using the SELEX process would allow for the identification of aptamers that inhibit LINGO-1-mediated or promote TrkB-mediated processes (*e.g.* LINGO-1 or TrkB-mediated promotion of cell survival).
- [0199] Polypeptide aptamers for use in the methods of the present invention are random peptides selected for their ability to bind to and block the action of LINGO-1. Polypeptide aptamers may include a short variable peptide domain attached at both ends to a protein scaffold. This double structural constraint greatly increases the binding affinity of the peptide aptamer to levels comparable to an antibody's (nanomolar range). See *e.g.*, Hoppe-Seyler F *et al. J. Mol. Med.* 78(8):426-430 (2000). The length of the short variable peptide is typically about 10 to 20 amino acids, and the scaffold may be any protein which has good solubility and compacity properties. One non-limiting example of a scaffold protein is the bacterial protein Thioredoxin-A. See, *e.g.*, Cohen BA *et al. PNAS* 95(24): 14272-14277 (1998). An additional, non-limiting example, of a polypeptide aptamer for use in the methods of the present invention is a Ligand Regulated Peptide Aptamer (LiRPA). The LiRPA scaffold may be composed of three protein domains: FK506 binding protein (FKBP), FRBP-Rapamycin binding domain (FRB) and glutathione-S-transferase (GST). See, *e.g.*, Binkowski BF *et al., Chem & Biol* 12(7): 847-855 (2005), incorporated herein by reference.
- [0200] Polypeptide aptamers are peptides or small polypeptides that act as dominant inhibitors of protein function. Peptide aptamers specifically bind to target proteins, blocking their functional ability. Kolonin *et al. Proc. Natl. Acad. Sci.* 95: 14,266-14,271 (1998). Peptide aptamers that bind with high affinity and specificity to a target protein

can be isolated by a variety of techniques known in the art. Peptide aptamers can be isolated from random peptide libraries by yeast two-hybrid screens (Xu, C.W., *et al. Proc. Natl. Acad. Sci.* 94:12,473-12,478 (1997)) or by ribosome display (Hanes *et al. Proc. Natl. Acad. Sci.* 94:4937-4942 (1997)). They can also be isolated from phage libraries (Hoogenboom, H.R., *et al. Immunotechnology* 4:1-20 (1998)) or chemically generated peptide libraries. Although the difficult means by which peptide aptamers are synthesized makes their use more complex than polynucleotide aptamers, they have unlimited chemical diversity.

[0201] Peptide aptamers for use in the methods of the present invention may be modified (*e.g.*, conjugated to polymers or fused to proteins) as described for other polypeptides elsewhere herein.

Fusion Proteins and Conjugated Polypeptides, Aptamers, Compounds and Antibodies

[0202] LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds and antibodies for use in the methods disclosed herein may further be recombinantly fused to a heterologous polypeptide at the N- or C-terminus or chemically conjugated (including covalent and non-covalent conjugations) to polypeptides or other compositions. For example, LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds and antibodies may be recombinantly fused or conjugated to molecules useful as labels in detection assays and effector molecules such as heterologous polypeptides, drugs, radionuclides, or toxins. *See, e.g.*, PCT publications WO 92/08495; WO 91/14438; WO 89/12624; U.S. Patent No. 5,314,995; and EP 396,387.

[0203] LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds and antibodies for use in the methods disclosed herein include derivatives that are modified, *i.e.*, by the covalent attachment of any type of molecule such that covalent attachment does not prevent the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound or antibody from inhibiting the biological function of LINGO-1 or TrkB. For example, but not by way of limitation, the LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds and antibodies of the present invention may be modified *e.g.*, by glycosylation, acetylation, pegylation, phosphorylation, phosphorylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, association with a cellular ligand or other protein, etc. Any of numerous chemical modifications may be carried out by known techniques, including, but not limited to

specific: chemical cleavage, acetylation, formylation, metabolic synthesis of tunicamycin, etc. Additionally, the derivative may contain one or more non-classical amino acids.

[0204] LINGO-1 antagonist or TrkB agonist polypeptides, aptamers and antibodies for use in the methods disclosed herein can be composed of amino acids joined to each other by peptide bonds or modified peptide bonds, *i.e.*, peptide isosteres, and may contain amino acids other than the 20 gene-encoded amino acids. LINGO-1 antagonist or TrkB agonist polypeptides, aptamers and antibodies may be modified by natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well as in a voluminous research literature. Modifications can occur anywhere in the LINGO-1 antagonist or TrkB agonist polypeptide or antibody, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini, or on moieties such as carbohydrates. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody. Also, a given LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody may contain many types of modifications. LINGO-1 antagonist or TrkB agonist polypeptides, aptamers or antibodies may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic LINGO-1 antagonist or TrkB agonist polypeptides, aptamers and antibodies may result from posttranslational natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphatidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, *Proteins - Structure And Molecular Properties*, T. E. Creighton, W. H. Freeman and Company, New York 2nd Ed., (1993); *Posttranslational Covalent Modification Of Proteins*, B. C. Johnson, Ed., Academic Press, New York, pgs.

1-12 (1983); Seifter *et al.*, *Meth Enzymol* 182:626-646 (1990); Rattan *et al.*, *Ann NY Acad Sci* 663:48-62 (1992)).

[0205] The present invention also provides for fusion proteins comprising, consisting essentially of, or consisting of a LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody fusion that inhibits or decreases LINGO-1 or increases or promotes TrkB function. In some embodiments, the heterologous polypeptide to which the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody is fused is useful for function or is useful to target the LINGO-1 antagonist or TrkB agonist polypeptide or antibody. In certain embodiments of the invention a soluble LINGO-1 antagonist or TrkB agonist polypeptide, *e.g.*, a LINGO-1 polypeptide comprising the LRR domains, Ig domain, or the entire extracellular domain (corresponding to amino acids 34 to 532 of SEQ ID NO: 2), is fused to a heterologous polypeptide moiety to form a LINGO-1 antagonist fusion polypeptide or a TrkB-agonist polypeptide, such as a BDNF polypeptide is fused to a heterologous polypeptide moiety to form a TrkB agonist fusion polypeptide. LINGO-1 antagonist or TrkB agonist fusion proteins, aptamers and antibodies can be used to accomplish various objectives, *e.g.*, increased serum half-life, improved bioavailability, *in vivo* targeting to a specific organ or tissue type, improved recombinant expression efficiency, improved host cell secretion, ease of purification, and higher avidity. Depending on the objective(s) to be achieved, the heterologous moiety can be inert or biologically active. Also, it can be chosen to be stably fused to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody or to be cleavable, *in vitro* or *in vivo*. Heterologous moieties to accomplish these other objectives are known in the art.

[0206] As an alternative to expression of a LINGO-1 antagonist or TrkB agonist fusion polypeptide, aptamer or antibody, a chosen heterologous moiety can be preformed and chemically conjugated to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody. In most cases, a chosen heterologous moiety will function similarly, whether fused or conjugated to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody. Therefore, in the following discussion of heterologous amino acid sequences, unless otherwise noted, it is to be understood that the heterologous sequence can be joined to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody in the form of a fusion protein or as a chemical conjugate.

[0207] Pharmacologically active polypeptides such as LINGO-1 antagonist or TrkB agonist polypeptides, aptamers or antibodies often exhibit rapid *in vivo* clearance,

necessitating large doses to achieve therapeutically effective concentrations in the body. In addition, polypeptides smaller than about 60 kDa potentially undergo glomerular filtration, which sometimes leads to nephrotoxicity. Fusion or conjugation of relatively small polypeptides such as LINGO-1 antagonist or TrkB agonist polypeptides, aptamers or antibodies can be employed to reduce or avoid the risk of such nephrotoxicity. Various heterologous amino acid sequences, *i.e.*, polypeptide moieties or "carriers," for increasing the *in vivo* stability, *i.e.*, serum half-life, of therapeutic polypeptides are known.

[0208] Due to its long half-life, wide *in vivo* distribution, and lack of enzymatic or immunological function, essentially full-length human serum albumin (HSA), or an HSA fragment, is commonly used as a heterologous moiety. Through application of methods and materials such as those taught in Yeh *et al.*, *Proc. Natl. Acad. Sci. USA* 89:1904-08 (1992) and Syed *et al.*, *Blood* 89:3243-52 (1997), HSA can be used to form a LINGO-1 antagonist or TrkB agonist fusion polypeptide, aptamer, antibody or polypeptide/antibody conjugate that displays pharmacological activity by virtue of the LINGO-1 or TrkB-agonist moiety while displaying significantly increased *in vivo* stability, *e.g.*, 10-fold to 100-fold higher. The C-terminus of the HSA can be fused to the N-terminus of the soluble LINGO-1 or TrkB-agonist moiety. Since HSA is a naturally secreted protein, the HSA signal sequence can be exploited to obtain secretion of the soluble LINGO-1 or TrkB-agonist fusion protein into the cell culture medium when the fusion protein is produced in a eukaryotic, *e.g.*, mammalian, expression system.

[0209] In certain embodiments, LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds, antibodies and antibody fragments thereof for use in the methods of the present invention further comprise a targeting moiety. Targeting moieties include a protein or a peptide which directs localization to a certain part of the body, for example, to the brain or compartments therein. In certain embodiments, LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds, antibodies or antibody fragments thereof for use in the methods of the present invention are attached or fused to a brain targeting moiety. The brain targeting moieties are attached covalently (*e.g.*, direct, translational fusion, or by chemical linkage either directly or through a spacer molecule, which can be optionally cleavable) or non-covalently attached (*e.g.*, through reversible interactions such as avidin, biotin, protein A, IgG, etc.). In other embodiments, the LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds, antibodies or antibody fragments thereof for use in the methods of the present invention are attached to one more brain targeting moieties. In additional embodiments, the brain targeting moiety

is attached to a plurality of LINGO-1 antagonist or TrkB agonist polypeptides, aptamers, compounds, antibodies or antibody fragments thereof for use in the methods of the present invention.

[0210] A brain targeting moiety associated with a LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound, antibody or antibody fragment thereof enhances brain delivery of such a LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound, antibody or antibody fragment thereof. A number of polypeptides have been described which, when fused to a protein or therapeutic agent, delivers the protein or therapeutic agent through the blood brain barrier (BBB). Non-limiting examples include the single domain antibody FC5 (Abulrob *et al. J. Neurochem.* 95, 1201-1214 (2005)); mAB 83-14, a monoclonal antibody to the human insulin receptor (Pardridge *et al. Pharmacol. Res.* 12, 807-816 (1995)); the B2, B6 and B8 peptides binding to the human transferrin receptor (hTfR) (Xia *et al. J. Virol.* 74, 11359-11366 (2000)); the OX26 monoclonal antibody to the transferrin receptor (Pardridge *et al. J. Pharmacol. Exp. Ther.* 259, 66-70 (1991)); and SEQ ID NOs: 1-18 of U.S. Patent No. 6,306,365. The contents of the above references are incorporated herein by reference in their entirety.

[0211] Enhanced brain delivery of a LINGO-1 or TrkB composition is determined by a number of means well established in the art. For example, administering to an animal a radioactively, enzymatically or fluorescently labeled LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound, antibody or antibody fragment thereof linked to a brain targeting moiety; determining brain localization; and comparing localization with an equivalent radioactively labeled LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound, antibody or antibody fragment thereof that is not associated with a brain targeting moiety. Other means of determining enhanced targeting are described in the above references.

[0212] The signal sequence is a polynucleotide that encodes an amino acid sequence that initiates transport of a protein across the membrane of the endoplasmic reticulum. Signal sequences useful for constructing an immunofusin include antibody light chain signal sequences, *e.g.*, antibody 14.18 (Gillies *et al., J. Immunol. Meth.* 125:191-202 (1989)), antibody heavy chain signal sequences, *e.g.*, the MOPC141 antibody heavy chain signal sequence (Sakano *et al., Nature* 286:5774 (1980)). Alternatively, other signal sequences can be used. *See, e.g.*, Watson, *Nucl. Acids Res.* 12:5145 (1984). The signal peptide is usually cleaved in the lumen of the endoplasmic reticulum by signal peptidases. This

results in the secretion of an immunofusin protein containing the Fc region and the soluble LINGO-1 or TrkB-agonist moiety.

[0213] In some embodiments, the DNA sequence may encode a proteolytic cleavage site between the secretion cassette and the LINGO-1 or TrkB-agonist moiety. Such a cleavage site may provide, *e.g.*, for the proteolytic cleavage of the encoded fusion protein, thus separating the Fc domain from the target protein. Useful proteolytic cleavage sites include amino acid sequences recognized by proteolytic enzymes such as trypsin, plasmin, thrombin, factor Xa, or enterokinase K.

[0214] The secretion cassette can be incorporated into a replicable expression vector. Useful vectors include linear nucleic acids, plasmids, phagemids, cosmids and the like. An exemplary expression vector is pdC, in which the transcription of the immunofusin DNA is placed under the control of the enhancer and promoter of the human cytomegalovirus. *See, e.g.*, Lo *et al.*, *Biochim. Biophys. Acta* 1088:712 (1991); and Lo *et al.*, *Protein Engineering* 11:495-500 (1998). An appropriate host cell can be transformed or transfected with a DNA that encodes a soluble LINGO-1 or TrkB-agonist polypeptide and used for the expression and secretion of the soluble LINGO-1 or TrkB-agonist polypeptide. Host cells that are typically used include immortal hybridoma cells, myeloma cells, 293 cells, Chinese hamster ovary (CHO) cells, HeLa cells, and COS cells.

[0215] In one embodiment, a soluble LINGO-1 antagonist or TrkB agonist polypeptide is fused to a hinge and Fc region, *i.e.*, the C-terminal portion of an Ig heavy chain constant region. Potential advantages of a LINGO-1-Fc or TrkB-agonist-Fc fusion include solubility, *in vivo* stability, and multivalency, *e.g.*, dimerization. The Fc region used can be an IgA, IgD, or IgG Fc region (hinge- C_H2- C_H3). Alternatively, it can be an IgE or IgM Fc region (hinge- C_H2- C_H3-C_H4). An IgG Fc region is generally used, *e.g.*, an IgG₁ Fc region or IgG₄ Fc region. In one embodiment, a sequence beginning in the hinge region just upstream of the papain cleavage site which defines IgG Fc chemically (*i.e.* residue 216, taking the first residue of heavy chain constant region to be 114 according to the Kabat system), or analogous sites of other immunoglobulins, is used in the fusion. The precise site at which the fusion is made is not critical; particular sites are well known and may be selected in order to optimize the biological activity, secretion, or binding characteristics of the molecule. Materials and methods for constructing and expressing DNA encoding Fc fusions are known in the art and can be applied to obtain soluble LINGO-1-antagonist or TrkB-agonist fusions without undue experimentation. Some embodiments of the invention employ a LINGO-1-antagonist or TrkB-agonist fusion

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protein such as those described in Capon *et al.*, U.S. Patent Nos. 5,428,130 and 5,565,335.

[0216] Fully intact, wild-type Fc regions display effector functions that normally are unnecessary and undesired in an Fc fusion protein used in the methods of the present invention. Therefore, certain binding sites typically are deleted from the Fc region during the construction of the secretion cassette. For example, since coexpression with the light chain is unnecessary, the binding site for the heavy chain binding protein, Bip (Hendershot *et al.*, *Immunol. Today* 8:111-14 (1987)), is deleted from the C_H2 domain of the Fc region of IgE, such that this site does not interfere with the efficient secretion of the immunofusin. Transmembrane domain sequences, such as those present in IgM, also are generally deleted.

[0217] The IgG₁ Fc region is most often used. Alternatively, the Fc region of the other subclasses of immunoglobulin gamma (gamma-2, gamma-3 and gamma-4) can be used in the secretion cassette. The IgG₁ Fc region of immunoglobulin gamma-1 is generally used in the secretion cassette and includes at least part of the hinge region, the C_H2 region, and the C_H3 region. In some embodiments, the Fc region of immunoglobulin gamma-1 is a C_H2-deleted-Fc, which includes part of the hinge region and the C_H3 region, but not the C_H2 region. A C_H2-deleted-Fc has been described by Gillies *et al.* *Hum. Antibod. Hybridomas* 1:47 (1990). In some embodiments, the Fc region of one of IgA, IgD, IgE, or IgM, is used.

[0218] LINGO-1-antagonist-Fc or TrkB-agonist-Fc fusion proteins can be constructed in several different configurations. In one configuration the C-terminus of the soluble LINGO-1 or TrkB-agonist moiety is fused directly to the N-terminus of the Fc hinge moiety. In a slightly different configuration, a short polypeptide, *e.g.*, 2-10 amino acids, is incorporated into the fusion between the N-terminus of the soluble LINGO-1 or TrkB-agonist moiety and the C-terminus of the Fc moiety. Such a linker provides conformational flexibility, which may improve biological activity in some circumstances. If a sufficient portion of the hinge region is retained in the Fc moiety, the LINGO-1-Fc or TrkB-agonist-Fc fusion will dimerize, thus forming a divalent molecule. A homogeneous population of monomeric Fc fusions will yield monospecific, bivalent dimers. A mixture of two monomeric Fc fusions each having a different specificity will yield bispecific, bivalent dimers.

[0219] Any of a number of cross-linkers that contain a corresponding amino-reactive group and thiol-reactive group can be used to link LINGO-1 antagonist or TrkB agonist

polypeptides to serum albumin. Examples of suitable linkers include amine reactive cross-linkers that insert a thiol-reactive maleimide, *e.g.*, SMCC, AMAS, BMPS, MBS, EMCS, SMPB, SMPH, KMUS, and GMBS. Other suitable linkers insert a thiol-reactive haloacetate group, *e.g.*, SBAP, SIA, SIAB. Linkers that provide a protected or non-protected thiol for reaction with sulfhydryl groups to product a reducible linkage include SPDP, SMPT, SATA, and SATP. Such reagents are commercially available (*e.g.*, Pierce Chemicals).

[0220] Conjugation does not have to involve the N-terminus of a soluble LINGO-1 or TrkB-agonist polypeptide or the thiol moiety on serum albumin. For example, soluble LINGO-1-albumin or TrkB-agonist-albumin fusions can be obtained using genetic engineering techniques, wherein the soluble LINGO-1 or TrkB-agonist moiety is fused to the serum albumin gene at its N-terminus, C-terminus, or both.

[0221] Soluble LINGO-1 or TrkB-agonist polypeptides can be fused at the N- or C-terminus to heterologous peptides in order to facilitate purification or identification of the soluble LINGO-1 or TrkB-agonist moiety. For example, a histidine tag can be fused to a soluble LINGO-1 or TrkB-agonist polypeptide to facilitate purification using commercially available chromatography media. Additionally, an epitope tag enables soluble LINGO-1 or TrkB-agonist fusion polypeptides to be readily purified by affinity purification using an anti-tag antibody or another type of affinity matrix that binds to the epitope tag. Many examples of such purification tags are known in the art and include, but are not limited to, poly-histidine (poly-his) or poly-histidine-glycine (poly-his-gly) tags; the influenza hemagglutinin (HA) tag polypeptide and its antibody 12CA5 (Field *et al.*, *Mol. Cell. Biol.* 8:2159-2165 (1988)); the c-myc tag and the 8F9, 3C7, 6E10, G4, B7 and 9E10 antibodies thereto (Evan *et al.*, *Mol. Cell. Bio.* 5:3610-3616 (1985)); and the Herpes Simplex virus glycoprotein D (gD) tag and its antibody (Paborsky *et al.*, *Protein Engineering* 3(6):547-553 (1990)). Other tag polypeptides include the Flag-peptide (Hopp *et al.*, *BioTechnology* 6:1204-1210 (1988)); the KT3 epitope peptide (Martin *et al.*, *Science* 255:192-194 (1992)); an α -tubulin epitope peptide (Skinner *et al.*, *J. Biol. Chem.* 266:15163-15166 (1991)); and the T7 gene 10 protein peptide tag (Lutz-Freyermuth *et al.*, *Proc. Natl. Acad. Sci. USA*, 87:6393-6397 (1990)). The tag can be any peptide epitope which is recognized by an antibody and does not interfere with the function of the soluble LINGO-1 or TrkB-agonist polypeptide.

[0222] In some embodiments of the invention, a soluble LINGO-1 or TrkB-agonist fusion construct is used to enhance the production of a soluble LINGO-1 or TrkB-agonist moiety

in bacteria. In such constructs a bacterial protein normally expressed and/or secreted at a high level is employed as the N-terminal fusion partner of a soluble LINGO-1 or TrkB-agonist polypeptide. See, e.g., Smith *et al.*, *Gene* 67:31 (1988); Hopp *et al.*, *Biotechnology* 6:1204 (1988); La Vallie *et al.*, *Biotechnology* 11:187 (1993).

[0223] By fusing a soluble LINGO-1 or TrkB-agonist moiety at the amino and carboxy termini of a suitable fusion partner, bivalent or tetravalent forms of a soluble LINGO-1 or TrkB-agonist polypeptide can be obtained. For example, a soluble LINGO-1 or TrkB-agonist moiety can be fused to the amino and carboxy termini of an Ig moiety to produce a bivalent monomeric polypeptide containing two soluble LINGO-1 or TrkB-agonist moieties. Upon dimerization of two of these monomers, by virtue of the Ig moiety, a tetravalent form of a soluble LINGO-1 or TrkB-agonist protein is obtained. Such multivalent forms can be used to achieve increased binding affinity for the target. Multivalent forms of soluble LINGO-1 or TrkB-agonist also can be obtained by placing soluble LINGO-1 or TrkB-agonist moieties in tandem to form concatamers, which can be employed alone or fused to a fusion partner such as Ig or HSA.

Conjugated Polymers (other than polypeptides)

[0224] Some embodiments of the invention involve a soluble LINGO-1 or TrkB-agonist polypeptide, LINGO-1-antagonist or TrkB-agonist aptamer, TrkB agonist compound or antagonistic LINGO-1 or agonistic TrkB antibody wherein one or more polymers are conjugated (covalently linked) to the LINGO-1 or TrkB-agonist polypeptide, compound, aptamer or antibody for use in the methods of the present invention. Examples of polymers suitable for such conjugation include polypeptides (discussed above), aptamers, sugar polymers and polyalkylene glycol chains. Typically, but not necessarily, a polymer is conjugated to the soluble LINGO-1 or TrkB-agonist polypeptide, aptamer, TrkB agonist compound or LINGO-1 or TrkB antibody for the purpose of improving one or more of the following: solubility, stability, or bioavailability.

[0225] The class of polymer generally used for conjugation to a LINGO-1 antagonist polypeptide, compound, aptamer or antibody or to a TrkB agonist polypeptide, compound, aptamer or antibody is a polyalkylene glycol. Polyethylene glycol (PEG) is most frequently used. PEG moieties, e.g., 1, 2, 3, 4 or 5 PEG polymers, can be conjugated to each LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, or antibody, or TrkB agonist compound to increase serum half life, as compared to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound or antibody alone.

PEG moieties are non-antigenic and essentially biologically inert. PEG moieties used in the practice of the invention may be branched or unbranched.

[0226] The number of PEG moieties attached to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer, compound or antibody and the molecular weight of the individual PEG chains can vary. In general, the higher the molecular weight of the polymer, the fewer polymer chains attached to the polypeptide. Usually, the total polymer mass attached to the LINGO-1 antagonist or TrkB agonist polypeptide, compound, aptamer or antibody is from 20 kDa to 40 kDa. Thus, if one polymer chain is attached, the molecular weight of the chain is generally 20-40 kDa. If two chains are attached, the molecular weight of each chain is generally 10-20 kDa. If three chains are attached, the molecular weight is generally 7-14 kDa.

[0227] The polymer, *e.g.*, PEG, can be linked to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody through any suitable, exposed reactive group on the polypeptide. The exposed reactive group(s) can be, *e.g.*, an N-terminal amino group or the epsilon amino group of an internal lysine residue, or both. An activated polymer can react and covalently link at any free amino group on the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody. Free carboxylic groups, suitably activated carbonyl groups, hydroxyl, guanidyl, imidazole, oxidized carbohydrate moieties and mercapto groups of the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody (if available) also can be used as reactive groups for polymer attachment.

[0228] In a conjugation reaction, from about 1.0 to about 10 moles of activated polymer per mole of polypeptide, depending on polypeptide concentration, is typically employed. Usually, the ratio chosen represents a balance between maximizing the reaction while minimizing side reactions (often non-specific) that can impair the desired pharmacological activity of the LINGO-1 antagonist or TrkB agonist polypeptide or antibody. In some embodiments, at least 50% of the biological activity (as demonstrated, *e.g.*, in any of the assays described herein or known in the art) of the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody is retained, and in some embodiments nearly 100% is retained.

[0229] The polymer can be conjugated to the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody using conventional chemistry. For example, a polyalkylene glycol moiety can be coupled to a lysine epsilon amino group of the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody. Linkage to the lysine side chain can be performed with an N-hydroxysuccinimide (NHS) active ester

such as PEG succinimidyl succinate (SS-PEG) and succinimidyl propionate (SPA-PEG). Suitable polyalkylene glycol moieties include, *e.g.*, carboxymethyl-NHS and norleucine-NHS, SC. These reagents are commercially available. Additional amine-reactive PEG linkers can be substituted for the succinimidyl moiety. These include, *e.g.*, isothiocyanates, nitrophenylcarbonates (PNP), epoxides, benzotriazole carbonates, SC-PEG, tresylate, aldehyde, epoxide, carbonylimidazole and PNP carbonate. Conditions are usually optimized to maximize the selectivity and extent of reaction. Such optimization of reaction conditions is within ordinary skill in the art.

[0230] PEGylation can be carried out by any of the PEGylation reactions known in the art. *See, e.g., Focus on Growth Factors* 3:4-10 (1992), and European patent applications EP 0 154 316 and EP 0 401 384. PEGylation may be carried out using an acylation reaction or an alkylation reaction with a reactive polyethylene glycol molecule (or an analogous reactive water-soluble polymer).

[0231] PEGylation by acylation generally involves reacting an active ester derivative of polyethylene glycol. Any reactive PEG molecule can be employed in the PEGylation. PEG esterified to N-hydroxysuccinimide (NHS) is a frequently used activated PEG ester. As used herein, "acylation" includes without limitation the following types of linkages between the therapeutic protein and a water-soluble polymer such as PEG: amide, carbamate, urethane, and the like. *See, e.g., Bioconjugate Chem.* 5:133-140, 1994. Reaction parameters are generally selected to avoid temperature, solvent, and pH conditions that would damage or inactivate the soluble LINGO-1 or TrkB-agonist polypeptide, aptamer or antibody.

[0232] Generally, the connecting linkage is an amide and typically at least 95% of the resulting product is mono-, di- or tri-PEGylated. However, some species with higher degrees of PEGylation may be formed in amounts depending on the specific reaction conditions used. Optionally, purified PEGylated species are separated from the mixture, particularly unreacted species, by conventional purification methods, including, *e.g.*, dialysis, salting-out, ultrafiltration, ion-exchange chromatography, gel filtration chromatography, hydrophobic exchange chromatography, and electrophoresis.

[0233] PEGylation by alkylation generally involves reacting a terminal aldehyde derivative of PEG with LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody in the presence of a reducing agent. In addition, one can manipulate the reaction conditions to favor PEGylation substantially only at the N-terminal amino group of LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody, *i.e.* a mono-

PEGylated protein. In either case of mono-PEGylation or poly-PEGylation, the PEG groups are typically attached to the protein via a -CH₂-NH- group. With particular reference to the -CH₂- group, this type of linkage is known as an "alkyl" linkage.

[0234] Derivatization via reductive alkylation to produce an N-terminally targeted mono-PEGylated product exploits differential reactivity of different types of primary amino groups (lysine versus the N-terminal) available for derivatization. The reaction is performed at a pH that allows one to take advantage of the pK_a differences between the epsilon-amino groups of the lysine residues and that of the N-terminal amino group of the protein. By such selective derivatization, attachment of a water-soluble polymer that contains a reactive group, such as an aldehyde, to a protein is controlled: the conjugation with the polymer takes place predominantly at the N-terminus of the protein and no significant modification of other reactive groups, such as the lysine side chain amino groups, occurs.

[0235] The polymer molecules used in both the acylation and alkylation approaches are selected from among water-soluble polymers. The polymer selected is typically modified to have a single reactive group, such as an active ester for acylation or an aldehyde for alkylation, so that the degree of polymerization may be controlled as provided for in the present methods. An exemplary reactive PEG aldehyde is polyethylene glycol propionaldehyde, which is water stable, or mono C1-C10 alkoxy or aryloxy derivatives thereof (*see, e.g.*, Harris *et al.*, U.S. Pat. No. 5,252,714). The polymer may be branched or unbranched. For the acylation reactions, the polymer(s) selected typically have a single reactive ester group. For reductive alkylation, the polymer(s) selected typically have a single reactive aldehyde group. Generally, the water-soluble polymer will not be selected from naturally occurring glycosyl residues, because these are usually made more conveniently by mammalian recombinant expression systems.

[0236] Methods for preparing a PEGylated LINGO-1 antagonist or TrkB-agonist polypeptide, aptamer or antibody generally includes the steps of (a) reacting a LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody with polyethylene glycol (such as a reactive ester or aldehyde derivative of PEG) under conditions whereby the molecule becomes attached to one or more PEG groups, and (b) obtaining the reaction product(s). In general, the optimal reaction conditions for the acylation reactions will be determined case-by-case based on known parameters and the desired result. For example, a larger ratio of PEG to protein generally leads to a greater the percentage of poly-PEGylated product.

- [0237] Reductive alkylation to produce a substantially homogeneous population of mono-polymer/soluble LINGO-1 antagonist or TrkB-agonist polypeptide, LINGO-1 antagonist or TrkB agonist aptamer or antagonistic LINGO-1 antibody or agonistic TrkB antibody generally includes the steps of: (a) reacting a LINGO-1-antagonist or TrkB-agonist protein or polypeptide with a reactive PEG molecule under reductive alkylation conditions, at a pH suitable to pen-nit selective modification of the N-terminal amino group of the polypeptide or antibody; and (b) obtaining the reaction product(s).
- [0238] For a substantially homogeneous population of mono-polymer/soluble LINGO-1 or TrkB-agonist polypeptide, LINGO-1 antagonist or TrkB agonist aptamer or antagonistic LINGO-1 antibody or agonistic TrkB antibody, the reductive alkylation reaction conditions are those that permit the selective attachment of the water-soluble polymer moiety to the N-terminus of the polypeptide or antibody. Such reaction conditions generally provide for pKa differences between the lysine side chain amino groups and the N-terminal amino group. For purposes of the present invention, the pH is generally in the range of 3-9, typically 3-6.
- [0239] LINGO-1 antagonist or TrkB-agonist polypeptides, aptamers or antibodies can include a tag, *e.g.*, a moiety that can be subsequently released by proteolysis. Thus, the lysine moiety can be selectively modified by first reacting a His-tag modified with a low-molecular-weight linker such as Traut's reagent (Pierce) which will react with both the lysine and N-terminus, and then releasing the His tag. The polypeptide will then contain a free SH group that can be selectively modified with a PEG containing a thiol-reactive head group such as a maleimide group, a vinylsulfone group, a haloacetate group, or a free or protected SH.
- [0240] Traut's reagent can be replaced with any linker that will set up a specific site for PEG attachment. For example, Traut's reagent can be replaced with SPDP, SMPT, SATA, or SATP (Pierce). Similarly, one could react the protein with an amine-reactive linker that inserts a maleimide (for example SMCC, AMAS, BMPS, MBS, EMCS, SMPB, SMPH, KMUS, or GMBS), a haloacetate group (SBAP, SIA, SIAB), or a vinylsulfone group and react the resulting product with a PEG that contains a free SH.
- [0241] In some embodiments, the polyalkylene glycol moiety is coupled to a cysteine group of the LINGO-1 antagonist or TrkB agonist polypeptide, aptamer or antibody for use in the methods of the present invention. Coupling can be effected using, *e.g.*, a maleimide group, a vinylsulfone group, a haloacetate group, or a thiol group.

[0242] Optionally, the LINGO-1 antagonist or TrkB-agonist polypeptide, aptamer or antibody is conjugated to the polyethylene-glycol moiety through a labile bond. The labile bond can be cleaved in, *e.g.*, biochemical hydrolysis, proteolysis, or sulfhydryl cleavage. For example, the bond can be cleaved under *in vivo* (physiological) conditions.

[0243] The reactions may take place by any suitable method used for reacting biologically active materials with inert polymers, generally at about pH 5-8, *e.g.*, pH 5, 6, 7, or 8, if the reactive groups are on the alpha amino group at the N-terminus. Generally the process involves preparing an activated polymer and thereafter reacting the protein with the activated polymer to produce the soluble protein suitable for formulation.

LINGO-1 Antagonist and TrkB Agonist Polynucleotides

LINGO-1 Antagonist Polynucleotides

[0244] LINGO-1 antagonists in the methods of the present invention include a LINGO-1 antagonist polynucleotide which comprises a nucleic acid molecule which specifically binds to a polynucleotide which encodes LINGO-1. The LINGO-1 polynucleotide antagonist generally prevents expression of LINGO-1 (knockdown). LINGO-1 antagonists include, but are not limited to antisense molecules, ribozymes, siRNA, shRNA, RNAi. Typically, such binding molecules are separately administered to the animal (*see*, for example, O'Connor, J. *Neurochem.* 56:560 (1991), but such binding molecules may also be expressed *in vivo* from polynucleotides taken up by a host cell and expressed *in vivo*. *See also* Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988).

[0245] RNAi refers to the expression of an RNA which interferes with the expression of the targeted mRNA. Specifically, the RNAi silences a targeted gene via interacting with the specific mRNA (*e.g.* LINGO-1 or TrkB) through an siRNA (short interfering RNA). The ds RNA complex is then targeted for degradation by the cell. Additional RNAi molecules include short hairpin RNA (shRNA); also short interfering hairpin. The shRNA molecule contains sense and antisense sequences from a target gene connected by a loop. The shRNA is transported from the nucleus into the cytoplasm, it is degraded along with the mRNA. Pol III or U6 promoters can be used to express RNAs for RNAi. In some embodiments of the invention, the shRNA is expressed from a lentiviral vector (*e.g.* pLL3.7).

[0246] RNAi is mediated by double stranded RNA (dsRNA) molecules that have sequence-specific homology to their "target" mRNAs (Caplen *et al.*, *Proc Natl Acad Sci*

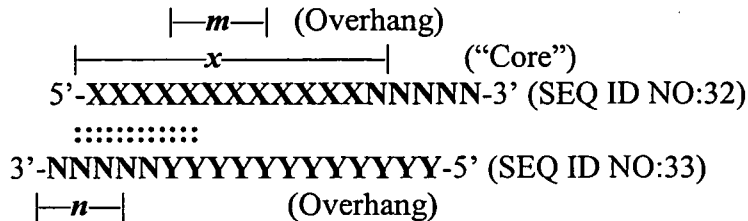
USA 98:9742-9747 (2001)). Biochemical studies in *Drosophila* cell-free lysates indicates that the mediators of RNA-dependent gene silencing are 21-25 nucleotide "small interfering" RNA duplexes (siRNAs). Accordingly, siRNA molecules are advantageously used in the methods of the present invention. The siRNAs are derived from the processing of dsRNA by an RNase known as DICER (Bernstein *et al.*, *Nature* 409:363-366 (2001)). It appears that siRNA duplex products are recruited into a multi-protein siRNA complex termed RISC (RNA Induced Silencing Complex). Without wishing to be bound by any particular theory, it is believed that a RISC is guided to a target mRNA, where the siRNA duplex interacts sequence-specifically to mediate cleavage in a catalytic fashion (Bernstein *et al.*, *Nature* 409:363-366 (2001); Boutla *et al.*, *Curr Biol* 11:1776-1780 (2001)).

[0247] RNAi has been used to analyze gene function and to identify essential genes in mammalian cells (Elbashir *et al.*, *Methods* 26:199-213 (2002); Harborth *et al.*, *J Cell Sci* 114:4557-4565 (2001)), including by way of non-limiting example neurons (Krichevsky *et al.*, *Proc Natl Acad Sci USA* 99:11926-11929 (2002)). RNAi is also being evaluated for therapeutic modalities, such as inhibiting or blocking the infection, replication and/or growth of viruses, including without limitation poliovirus (Gitlin *et al.*, *Nature* 418:379-380 (2002)) and HIV (Capodici *et al.*, *J Immunol* 169:5196-5201 (2002)), and reducing expression of oncogenes (*e.g.*, the bcr-abl gene; Scherr *et al.*, *Blood* 101(4):1566-9 (2002)). RNAi has been used to modulate gene expression in mammalian (mouse) and amphibian (*Xenopus*) embryos (respectively, Calegari *et al.*, *Proc Natl Acad Sci USA* 99:14236-14240 (2002); and Zhou, *et al.*, *Nucleic Acids Res* 30:1664-1669 (2002)), and in postnatal mice (Lewis *et al.*, *Nat Genet* 32:107-108 (2002)), and to reduce transgene expression in adult transgenic mice (McCaffrey *et al.*, *Nature* 418:38-39 (2002)). Methods have been described for determining the efficacy and specificity of siRNAs in cell culture and *in vivo* (see, *e.g.*, Bertrand *et al.*, *Biochem Biophys Res Commun* 296:1000-1004 (2002); Lassus *et al.*, *Sci STKE* 2002(147):PL13 (2002); and Leirdal *et al.*, *Biochem Biophys Res Commun* 295:744-748 (2002)).

[0248] Molecules that mediate RNAi, including without limitation siRNA, can be produced *in vitro* by chemical synthesis (Hohjoh, *FEBS Lett* 521:195-199 (2002)), hydrolysis of dsRNA (Yang *et al.*, *Proc Natl Acad Sci USA* 99:9942-9947 (2002)), by *in vitro* transcription with T7 RNA polymerase (Donzeet *et al.*, *Nucleic Acids Res* 30:e46, (2002); Yu *et al.*, *Proc Natl Acad Sci USA* 99:6047-6052 (2002)), and by hydrolysis of

double-stranded RNA using a nuclease such as *E. coli* RNase III (Yang *et al.*, *Proc Natl Acad Sci USA* 99:9942-9947 (2002)).

[0249] siRNA molecules may also be formed by annealing two oligonucleotides to each other, typically have the following general structure, which includes both double-stranded and single-stranded portions:



[0250] Wherein N, X and Y are nucleotides; X hydrogen bonds to Y; ":" signifies a hydrogen bond between two bases; x is a natural integer having a value between 1 and about 100; and m and n are whole integers having, independently, values between 0 and about 100. In some embodiments, N, X and Y are independently A, G, C and T or U. Non-naturally occurring bases and nucleotides can be present, particularly in the case of synthetic siRNA (*i.e.*, the product of annealing two oligonucleotides). The double-stranded central section is called the "core" and has base pairs (bp) as units of measurement; the single-stranded portions are overhangs, having nucleotides (nt) as units of measurement. The overhangs shown are 3' overhangs, but molecules with 5' overhangs are also within the scope of the invention. Also within the scope of the invention are siRNA molecules with no overhangs (*i.e.*, $m = 0$ and $n = 0$), and those having an overhang on one side of the core but not the other (*e.g.*, $m = 0$ and $n \geq 1$, or vice-versa).

[0251] Initially, RNAi technology did not appear to be readily applicable to mammalian systems. This is because, in mammals, dsRNA activates dsRNA-activated protein kinase (PKR) resulting in an apoptotic cascade and cell death (Der *et al.*, *Proc. Natl. Acad. Sci. USA* 94:3279-3283 (1997)). In addition, it has long been known that dsRNA activates the interferon cascade in mammalian cells, which can also lead to altered cell physiology (Colby *et al.*, *Annu. Rev. Microbiol.* 25:333 (1971); Kleinschmidt *et al.*, *Annu. Rev. Biochem.* 41:517 (1972); Lampson *et al.*, *Proc. Natl. Acad. Sci. USA* 58:L782 (1967); Lomniczi *et al.*, *J. Gen. Virol.* 8:55 (1970); and Younger *et al.*, *J. Bacteriol.* 92:862 (1966)). However, dsRNA-mediated activation of the PKR and interferon cascades requires dsRNA longer than about 30 base pairs. In contrast, dsRNA less than 30 base

pairs in length has been demonstrated to cause RNAi in mammalian cells (Caplen *et al.*, *Proc. Natl. Acad. Sci. USA* 98:9742-9747 (2001)). Thus, it is expected that undesirable, non-specific effects associated with longer dsRNA molecules can be avoided by preparing short RNA that is substantially free from longer dsRNAs.

[0252] References regarding siRNA: Bernstein *et al.*, *Nature* 409:363-366 (2001); Boutla *et al.*, *Curr Biol* 11:1776-1780 (2001); Cullen, *Nat Immunol.* 3:597-599 (2002); Caplen *et al.*, *Proc Natl Acad Sci USA* 98:9742-9747 (2001); Hamilton *et al.*, *Science* 286:950-952 (1999); Nagase *et al.*, *DNA Res.* 6:63-70 (1999); Napoli *et al.*, *Plant Cell* 2:279-289 (1990); Nicholson *et al.*, *Mamm. Genome* 13:67-73 (2002); Parrish *et al.*, *Mol Cell* 6:1077-1087 (2000); Romano *et al.*, *Mol Microbiol* 6:3343-3353 (1992); Tabara *et al.*, *Cell* 99:123-132 (1999); and Tuschl, *Chembiochem.* 2:239-245 (2001).

[0253] Paddison *et al.* (*Genes & Dev.* 16:948-958 (2002)) have used small RNA molecules folded into hairpins as a means to effect RNAi. Accordingly, such short hairpin RNA (shRNA) molecules are also advantageously used in the methods of the invention. The length of the stem and loop of functional shRNAs varies; stem lengths can range anywhere from about 25 to about 30 nt, and loop size can range between 4 to about 25 nt without affecting silencing activity. While not wishing to be bound by any particular theory, it is believed that these shRNAs resemble the dsRNA products of the DICER RNase and, in any event, have the same capacity for inhibiting expression of a specific gene.

[0254] Antisense technology can be used to control gene expression through antisense DNA or RNA, or through triple-helix formation. Antisense techniques are discussed, for example, in Okano, J. *Neurochem.* 56:560 (1991); Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988). Triple helix formation is discussed in, for instance, Lee *et al.*, *Nucleic Acids Research* 6:3073 (1979); Cooney *et al.*, *Science* 241:456 (1988); and Dervan *et al.*, *Science* 251:1300 (1991). The methods are based on binding of a polynucleotide to a complementary DNA or RNA.

[0255] For example, the 5' coding portion of a polynucleotide that encodes LINGO-1 may be used to design an antisense RNA oligonucleotide of from about 10 to 40 base pairs in length. A DNA oligonucleotide is designed to be complementary to a region of the gene involved in transcription, thereby preventing transcription and the production of the target protein. The antisense RNA oligonucleotide hybridizes to the mRNA *in vivo* and blocks translation of the mRNA molecule into the target polypeptide.

[0256] In one embodiment, antisense nucleic acids specific for the LINGO-1 gene are produced intracellularly by transcription from an exogenous sequence. For example, a vector or a portion thereof, is transcribed, producing an antisense nucleic acid (RNA). Such a vector can remain episomal or become chromosomally integrated, as long as it can be transcribed to produce the desired antisense RNA. Such vectors can be constructed by recombinant DNA technology methods standard in the art. Vectors can be plasmid, viral, or others known in the art, used for replication and expression in vertebrate cells. Expression of the antisense molecule, can be by any promoter known in the art to act in vertebrate, especially human cells, such as those described elsewhere herein.

[0257] Absolute complementarity of an antisense molecule is not required. A sequence complementary to at least a portion of an RNA encoding LINGO-1 means a sequence having sufficient complementarity to be able to hybridize with the RNA, forming a stable duplex; or triplex formation may be assayed. The ability to hybridize will depend on both the degree of complementarity and the length of the antisense nucleic acid. Generally, the larger the hybridizing nucleic acid, the more base mismatches it may contain and still form a stable duplex (or triplex as the case may be). One skilled in the art can ascertain a tolerable degree of mismatch by use of standard procedures to determine the melting point of the hybridized complex.

[0258] Oligonucleotides that are complementary to the 5' end of a messenger RNA, *e.g.*, the 5' untranslated sequence up to and including the AUG initiation codon, should work most efficiently at inhibiting translation. However, sequences complementary to the 3' untranslated sequences of mRNAs have been shown to be effective at inhibiting translation of mRNAs as well. *See, generally, Wagner, R., Nature 372:333-335 (1994).* Thus, oligonucleotides complementary to either the 5'- or 3'- non-translated, non-coding regions could be used in an antisense approach to inhibit translation of LINGO-1. Oligonucleotides complementary to the 5' untranslated region of the mRNA should include the complement of the AUG start codon. Antisense oligonucleotides complementary to mRNA coding regions are less efficient inhibitors of translation but could be used in accordance with the invention. Antisense nucleic acids should be at least six nucleotides in length, and in some embodiments are oligonucleotides ranging from 6 to about 50 nucleotides in length. In specific aspects the oligonucleotide is at least 10 nucleotides, at least 17 nucleotides, at least 25 nucleotides or at least 50 nucleotides.

[0259] In yet another embodiment, an antisense oligonucleotide for use in the methods disclosed herein is an α -anomeric oligonucleotide. An α -anomeric oligonucleotide forms

specific double-stranded hybrids with complementary RNA in which, contrary to the usual situation, the strands run parallel to each other (Gautier *et al.*, *Nucl. Acids Res.* 15:6625-6641(1987)). The oligonucleotide is a 2'-O-methylribonucleotide (Inoue *et al.*, *Nucl. Acids Res.* 15:6131-6148(1987)), or a chimeric RNA-DNA analogue (Inoue *et al.*, *FEBS Lett.* 215:327-330 (1987)).

[0260] Polynucleotide compositions for use in the methods disclosed herein further include catalytic RNA, or a ribozyme (*See, e.g.*, PCT International Publication WO 90/11364, published October 4, 1990; Sarver *et al.*, *Science* 247:1222-1225 (1990)). In some embodiments of the invention, hammerhead ribozymes are used. Hammerhead ribozymes cleave mRNAs at locations dictated by flanking regions that form complementary base pairs with the target mRNA. The sole requirement is that the target mRNA have the following sequence of two bases: 5'-UG-3'. The construction and production of hammerhead ribozymes is well known in the art and is described more fully in Haseloff and Gerlach, *Nature* 334:585-591 (1988). The ribozyme can be engineered so that the cleavage recognition site is located near the 5' end of the target mRNA; *i.e.*, to increase efficiency and minimize the intracellular accumulation of non-functional mRNA transcripts.

[0261] As in the antisense approach, ribozymes for use in the methods disclosed herein can be composed of modified oligonucleotides (*e.g.* for improved stability, targeting, etc.) and may be delivered to cells which express LINGO-1 *in vivo*. DNA constructs encoding the ribozyme may be introduced into the cell in the same manner as described above for the introduction of antisense encoding DNA. One method of delivery involves using a DNA construct "encoding" the ribozyme under the control of a strong constitutive promoter, such as, for example, pol III or pol II promoter, so that transfected cells will produce sufficient quantities of the ribozyme to destroy endogenous LINGO-1 messages and inhibit translation. Since ribozymes, unlike antisense molecules, are catalytic, a lower intracellular concentration is required for efficiency.

TrkB Agonist Polynucleotides

[0262] TrkB agonists for use in the methods of the present invention include a TrkB agonist polynucleotide which comprises a nucleic acid molecule which encodes a TrkB polypeptide, fragment, isoform or variant thereof. The TrkB agonist polynucleotides of the present invention also include nucleic acid molecules which encode a TrkB ligand polypeptide, fragment, isoform or variant thereof. In some embodiments, the TrkB

agonist polynucleotide encodes BDNF. TrkB agonist polynucleotides include any polynucleotide which encodes a TrkB agonist polypeptide of the present invention.

LINGO-1 Antagonists and/or TrkB Agonist Polynucleotides

[0263] Polynucleotides for use in the methods disclosed herein, including aptamers described *supra*, can be DNA or RNA or chimeric mixtures or derivatives or modified versions thereof, single-stranded or double-stranded. The polynucleotide can be modified at the base moiety, sugar moiety, or phosphate backbone, for example, to improve stability of the molecule, hybridization, etc. The polynucleotide may include other appended groups such as peptides (*e.g.*, for targeting host cell receptors *in vivo*), or agents facilitating transport across the cell membrane (see, *e.g.*, Letsinger *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 86:6553-6556 (1989); Lemaitre *et al.*, *Proc. Natl. Acad. Sci.* 84:648-652 (1987)); PCT Publication No. WO88/09810, published December 15, 1988) or the blood-brain barrier (see, *e.g.*, PCT Publication No. WO89/10134, published April 25, 1988), hybridization-triggered cleavage agents. (See, *e.g.*, Krol *et al.*, *BioTechniques* 6:958-976 (1988)) or intercalating agents. (See, *e.g.*, Zon, *Pharm. Res.* 5:539-549(1988)). To this end, the polynucleotide may be conjugated to another molecule, *e.g.*, a peptide, hybridization triggered cross-linking agent, transport agent, hybridization-triggered cleavage agent, etc.

[0264] Polynucleotides, including aptamers, for use in the methods disclosed herein may comprise at least one modified base moiety which is selected from the group including, but not limited to, 5-fluorouracil, 5-bromouracil, 5-chlorouracil, 5-iodouracil, hypoxanthine, xantine, 4-acetylcytosine, 5-(carboxyhydroxymethyl) uracil, 5-carboxymethylaminomethyl-2-thiouridine, 5-carboxymethylaminomethyluracil, dihydrouracil, beta-D-galactosylqueosine, inosine, N-6-isopentenyladenine, 1-methylguanine, 1-methylinosine, 2,2-dimethylguanine, 2-methyladenine, 2-methylguanine, 3-methylcytosine, 5-methylcytosine, N-6-adenine, 7-methylguanine, 5-methylaminomethyluracil, 5-methoxyaminomethyl-2-thiouracil, beta-D-mannosylqueosine, 5'-methoxycarboxymethyluracil, 5-methoxyuracil, 2-methylthio-N-6-isopentenyladenine, uracil-5-oxyacetic acid (v), wybutoxosine, pseudouracil, queosine, 2-thiocytosine, 5-methyl-2-thiouracil, 2-thiouracil, 4-thiouracil, 5-methyluracil, uracil-5-oxyacetic acid methylester, uracil-5-oxyacetic acid (v), 5-methyl-2-thiouracil, 3(3-amino-3-N2-carboxypropyl) uracil, (acp3)w, and 2,6-diaminopurine.

- [0265] Polynucleotides, including aptamers, for use in the methods disclosed herein may also comprise at least one modified sugar moiety selected from the group including, but not limited to, arabinose, 2-fluoroarabinose, xylulose, and hexose.
- [0266] In yet another embodiment, a polynucleotide, including an aptamer, for use in the methods disclosed herein comprises at least one modified phosphate backbone selected from the group including, but not limited to, a phosphorothioate, a phosphorodithioate, a phosphoramidothioate, a phosphoramidate, a phosphordiamidate, a methylphosphonate, an alkyl phosphotriester, and a formacetal or analog thereof.
- [0267] Polynucleotides, including aptamers, for use in the methods of the invention may be synthesized by standard methods known in the art, *e.g.* by use of an automated DNA synthesizer (such as are commercially available from Biosearch, Applied Biosystems, etc.). As examples, phosphorothioate oligonucleotides may be synthesized by the method of Stein *et al.*, *Nucl. Acids Res.* 16:3209 (1988), methylphosphonate oligonucleotides can be prepared by use of controlled pore glass polymer supports (Sarin *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 85:7448-7451 (1988)), etc.

Vectors and Host Cells

- [0268] Host-expression systems represent vehicles by which the coding sequences of interest may be produced and subsequently purified, but also represent cells which may, when transformed or transfected with the appropriate nucleotide coding sequences, express an antibody molecule of the invention *in situ*. These include but are not limited to microorganisms such as bacteria (*e.g.*, *E. coli*, *B. subtilis*) transformed with recombinant bacteriophage DNA, plasmid DNA or cosmid DNA expression vectors containing antibody coding sequences; yeast (*e.g.*, *Saccharomyces*, *Pichia*) transformed with recombinant yeast expression vectors containing antibody coding sequences; insect cell systems infected with recombinant virus expression vectors (*e.g.*, baculovirus) containing antibody coding sequences; plant cell systems infected with recombinant virus expression vectors (*e.g.*, cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or transformed with recombinant plasmid expression vectors (*e.g.*, Ti plasmid) containing antibody coding sequences; or mammalian cell systems (*e.g.*, COS, CHO, BLK, 293, 3T3 cells) harboring recombinant expression constructs containing promoters derived from the genome of mammalian cells (*e.g.*, metallothionein promoter) or from mammalian viruses (*e.g.*, the adenovirus late promoter; the vaccinia virus 7.5K promoter). Bacterial cells such as *Escherichia coli*, or eukaryotic cells (especially for the expression of whole

recombinant antibody molecule) are used for the expression of a recombinant antibody molecule. For example, mammalian cells such as Chinese hamster ovary cells (CHO), in conjunction with a vector such as the major intermediate early gene promoter element from human cytomegalovirus is an effective expression system for antibodies (Foecking *et al.*, *Gene* 45:101 (1986); Cockett *et al.*, *Bio/Technology* 8:2 (1990)).

[0269] In bacterial systems, a number of expression vectors may be advantageously selected depending upon the use intended for the antibody molecule being expressed. For example, when a large quantity of such a protein is to be produced, for the generation of pharmaceutical compositions of an antibody molecule, vectors which direct the expression of high levels of fusion protein products that are readily purified may be desirable. Such vectors include, but are not limited to, the *E. coli* expression vector pUR278 (Ruther *et al.*, *EMBO J.* 2:1791 (1983)), in which the antibody coding sequence may be ligated individually into the vector in frame with the lacZ coding region so that a fusion protein is produced; pIN vectors (Inouye & Inouye, *Nucleic Acids Res.* 13:3101-3109 (1985); Van Heeke & Schuster, *J. Biol. Chem.* 24:5503-5509 (1989)); and the like. pGEX vectors may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption and binding to a matrix glutathione-agarose beads followed by elution in the presence of free glutathione. The pGEX vectors are designed to include thrombin or factor Xa protease cleavage sites so that the cloned target gene product can be released from the GST moiety.

[0270] In an insect system, *Autographa californica* nuclear polyhedrosis virus (AcNPV) is typically used as a vector to express foreign genes. The virus grows in *Spodoptera frugiperda* cells. The antibody coding sequence may be cloned individually into non-essential regions (for example the polyhedrin gene) of the virus and placed under control of an AcNPV promoter (for example the polyhedrin promoter).

[0271] In mammalian host cells, a number of viral-based expression systems may be utilized. In cases where an adenovirus is used as an expression vector, the antibody coding sequence of interest may be ligated to an adenovirus transcription/translation control complex, *e.g.*, the late promoter and tripartite leader sequence. This chimeric gene may then be inserted in the adenovirus genome by *in vitro* or *in vivo* recombination. Insertion in a non-essential region of the viral genome (*e.g.*, region E1 or E3) will result in a recombinant virus that is viable and capable of expressing the antibody molecule in infected hosts. (*e.g.*, see Logan & Shenk, *Proc. Natl. Acad. Sci. USA* 81:355-359 (1984)).

Specific initiation signals may also be required for efficient translation of inserted antibody coding sequences. These signals include the ATG initiation codon and adjacent sequences. Furthermore, the initiation codon must be in phase with the reading frame of the desired coding sequence to ensure translation of the entire insert. These exogenous translational control signals and initiation codons can be of a variety of origins, both natural and synthetic. The efficiency of expression may be enhanced by the inclusion of appropriate transcription enhancer elements, transcription terminators, etc. (*see Bittner et al., Methods in Enzymol. 153:51-544 (1987)*).

[0272] In addition, a host cell strain may be chosen which modulates the expression of the inserted sequences, or modifies and processes the gene product in the specific fashion desired. Such modifications (*e.g.*, glycosylation) and processing (*e.g.*, cleavage) of protein products may be important for the function of the protein. Different host cells have characteristic and specific mechanisms for the post-translational processing and modification of proteins and gene products. Appropriate cell lines or host systems can be chosen to ensure the correct modification and processing of the foreign protein expressed. To this end, eukaryotic host cells which possess the cellular machinery for proper processing of the primary transcript, glycosylation, and phosphorylation of the gene product may be used. Such mammalian host cells include but are not limited to CHO, VERY, BHK, HeLa, COS, MDCK, 293, 3T3, WI38, and in particular, breast cancer cell lines such as, for example, BT483, Hs578T, HTB2, BT20 and T47D, and normal mammary gland cell line such as, for example, CRL7030 and Hs578Bst.

[0273] For long-term, high-yield production of recombinant proteins, stable expression may be used. For example, cell lines which stably express the antibody molecule may be engineered. Rather than using expression vectors which contain viral origins of replication, host cells can be transformed with DNA controlled by appropriate expression control elements (*e.g.*, promoter, enhancer, sequences, transcription terminators, polyadenylation sites, etc.), and a selectable marker. Following the introduction of the foreign DNA, engineered cells may be allowed to grow for 1-2 days in an enriched media, and then are switched to a selective media. The selectable marker in the recombinant plasmid confers resistance to the selection and allows cells to stably integrate the plasmid into their chromosomes and grow to form foci which in turn can be cloned and expanded into cell lines. This method may advantageously be used to engineer cell lines which stably express the antibody molecule.

- [0274] A number of selection systems may be used, including but not limited to the herpes simplex virus thymidine kinase (Wigler *et al.*, *Cell* 11:223 (1977)), hypoxanthine-guanine phosphoribosyltransferase (Szybalska & Szybalski, *Proc. Natl. Acad. Sci. USA* 48:202 (1992)), and adenine phosphoribosyltransferase (Lowy *et al.*, *Cell* 22:817 (1980)) genes, can be employed in tk-, hgp^{rt}- or ap^{rt}-cells, respectively. Also, antimetabolite resistance can be used as the basis of selection for the following genes: dhfr, which confers resistance to methotrexate (Wigler *et al.*, *Natl. Acad. Sci. USA* 77:357 (1980); O'Hare *et al.*, *Proc. Natl. Acad. Sci. USA* 78:1527 (1981)); gpt, which confers resistance to mycophenolic acid (Mulligan & Berg, *Proc. Natl. Acad. Sci. USA* 78:2072 (1981)); neo, which confers resistance to the aminoglycoside G-418 *Clinical Pharmacy* 12:488-505; Wu and Wu, *Biotherapy* 3:87-95 (1991); Tolstoshev, *Ann. Rev. Pharmacol. Toxicol.* 32:573-596 (1993); Mulligan, *Science* 260:926-932 (1993); and Morgan and Anderson, *Ann. Rev. Biochem.* 62:191-217 (1993); *TIB TECH* 11(5):155-215 (May, 1993); and hyg^{ro}, which confers resistance to hygromycin (Santerre *et al.*, *Gene* 30:147 (1984). Methods commonly known in the art of recombinant DNA technology which can be used are described in Ausubel *et al.* (eds.), *Current Protocols in Molecular Biology*, John Wiley & Sons, NY (1993); Kriegler, *Gene Transfer and Expression*, A Laboratory Manual, Stockton Press, NY (1990); and in Chapters 12 and 13, Dracopoli *et al.* (eds), *Current Protocols in Human Genetics*, John Wiley & Sons, NY (1994); Colberre-Garapin *et al.*, *J. Mol. Biol.* 150:1 (1981), which are incorporated by reference herein in their entireties.
- [0275] The expression levels of an antibody molecule can be increased by vector amplification (for a review, see Bebbington and Hentschel, *The use of vectors based on gene amplification for the expression of cloned genes in mammalian cells in DNA cloning*, Academic Press, New York, Vol. 3. (1987)). When a marker in the vector system expressing antibody is amplifiable, increase in the level of inhibitor present in culture of host cell will increase the number of copies of the marker gene. Since the amplified region is associated with the antibody gene, production of the antibody will also increase (Crouse *et al.*, *Mol. Cell. Biol.* 3:257 (1983)).
- [0276] Vectors comprising nucleic acids encoding LINGO-1 antagonist or TrkB agonists may also be used to produce polynucleotides or polypeptides for use in the methods of the invention. The choice of vector and expression control sequences to which such nucleic acids are operably linked depends on the functional properties desired, e.g., protein expression, and the host cell to be transformed.

- [0277] Expression control elements useful for regulating the expression of an operably linked coding sequence are known in the art. Examples include, but are not limited to, inducible promoters, constitutive promoters, secretion signals, and other regulatory elements. When an inducible promoter is used, it can be controlled, *e.g.*, by a change in nutrient status, or a change in temperature, in the host cell medium.
- [0278] The vector can include a prokaryotic replicon, *i.e.*, a DNA sequence having the ability to direct autonomous replication and maintenance of the recombinant DNA molecule extra-chromosomally in a bacterial host cell. Such replicons are well known in the art. In addition, vectors that include a prokaryotic replicon may also include a gene whose expression confers a detectable marker such as a drug resistance. Examples of bacterial drug-resistance genes are those that confer resistance to ampicillin or tetracycline.
- [0279] Vectors that include a prokaryotic replicon can also include a prokaryotic or bacteriophage promoter for directing expression of the coding gene sequences in a bacterial host cell. Promoter sequences compatible with bacterial hosts are typically provided in plasmid vectors containing convenient restriction sites for insertion of a DNA segment to be expressed. Examples of such plasmid vectors are pUC8, pUC9, pBR322 and pBR329 (BioRad), pPL and pKK223 (Pharmacia). Any suitable prokaryotic host can be used to express a recombinant DNA molecule encoding a protein used in the methods of the invention.
- [0280] For the purposes of this invention, numerous expression vector systems may be employed. For example, one class of vector utilizes DNA elements which are derived from animal viruses such as bovine papilloma virus, polyoma virus, adenovirus, vaccinia virus, baculovirus, retroviruses (RSV, MMTV or MOMLV) or SV40 virus. Others involve the use of polycistronic systems with internal ribosome binding sites. Additionally, cells which have integrated the DNA into their chromosomes may be selected by introducing one or more markers which allow selection of transfected host cells. The marker may provide for prototrophy to an auxotrophic host, biocide resistance (*e.g.*, antibiotics) or resistance to heavy metals such as copper. The selectable marker gene can either be directly linked to the DNA sequences to be expressed, or introduced into the same cell by cotransformation. The neomycin phosphotransferase (*neo*) gene is an example of a selectable marker gene (Southern *et al.*, *J. Mol. Anal. Genet.* 1:327-341 (1982)). Additional elements may also be needed for optimal synthesis of mRNA. These

elements may include signal sequences or splice signals, as well as transcriptional promoters, enhancers, and termination signals.

[0281] In one embodiment, an expression vector referred to as NEOSPLA (U.S. patent 6,159,730, incorporated herein by reference) may be used. This vector contains the cytomegalovirus promoter/enhancer, the mouse beta globin major promoter, the SV40 origin of replication, the bovine growth hormone polyadenylation sequence, neomycin phosphotransferase exon 1 and exon 2, the dihydrofolate reductase gene and leader sequence. This vector has been found to result in very high-level expression upon transfection in CHO cells, followed by selection in G418-containing medium and methotrexate amplification. Of course, any expression vector which is capable of eliciting expression in eukaryotic cells may be used in the present invention. Examples of suitable vectors include, but are not limited to, plasmids pcDNA3, pHCMV/Zeo, pCR3.1, pEF1/His, pIND/GS, pRc/HCMV2, pSV40/Zeo2, pTRACER-HCMV, pUB6/V5-His, pVAX1, and pZeoSV2 (available from Invitrogen, San Diego, CA), and plasmid pCI (available from Promega, Madison, WI). Additional eukaryotic cell expression vectors are known in the art and are commercially available. Typically, such vectors contain convenient restriction sites for insertion of the desired DNA segment. Exemplary vectors include pSVL and pKSV-10 (Pharmacia), pBPV-1, pml2d (International Biotechnologies), pTDT1 (ATCC 31255), retroviral expression vector pMIG and pLL3.7, adenovirus shuttle vector pDC315, and AAV vectors. Other exemplary vector systems are disclosed *e.g.*, in U.S. Patent 6,413,777.

[0282] In general, screening large numbers of transformed cells for those which express suitably high levels of the antagonist is routine experimentation which can be carried out, for example, by robotic systems.

[0283] Frequently used regulatory sequences for mammalian host cell expression include viral elements that direct high levels of protein expression in mammalian cells, such as promoters and enhancers derived from retroviral LTRs, cytomegalovirus (CMV) (such as the CMV promoter/enhancer), Simian Virus 40 (SV40) (such as the SV40 promoter/enhancer), adenovirus, (*e.g.*, the adenovirus major late promoter (AdmlP)), polyoma and strong mammalian promoters such as native immunoglobulin and actin promoters. For further description of viral regulatory elements, and sequences thereof, *see, e.g.*, Stinski, U.S. Pat. No. 5,168,062; Bell, U.S. Pat. No. 4,510,245; and Schaffner, U.S. Pat. No. 4,968,615.

- [0284] The recombinant expression vectors may carry sequences that regulate replication of the vector in host cells (*e.g.*, origins of replication) and selectable marker genes. The selectable marker gene facilitates selection of host cells into which the vector has been introduced (*see, e.g.*, Axel, U.S. Pat. Nos. 4,399,216; 4,634,665 and 5,179,017). For example, typically the selectable marker gene confers resistance to a drug, such as G418, hygromycin or methotrexate, on a host cell into which the vector has been introduced. Frequently used selectable marker genes include the dihydrofolate reductase (DHFR) gene (for use in dhfr- host cells with methotrexate selection/amplification) and the neo gene (for G418 selection).
- [0285] Vectors encoding LINGO-1 antagonist or TrkB agonists can be used for transformation of a suitable host cell. Transformation can be by any suitable method. Methods for introduction of exogenous DNA into mammalian cells are well known in the art and include dextran-mediated transfection, calcium phosphate precipitation, polybrene-mediated transfection, protoplast fusion, electroporation, transfection via encapsulation of the polynucleotide(s) in liposomes, and direct microinjection of the DNA into nuclei. In addition, nucleic acid molecules may be introduced into mammalian cells by viral vectors. Mammalian cells may also be transduced by recombinant viruses containing the exogenous DNA which is to be introduced into the mammalian cells.
- [0286] Host cells for expression of a LINGO-1 antagonist or TrkB agonist for use in a method of the invention may be prokaryotic or eukaryotic. Exemplary eukaryotic host cells include, but are not limited to, yeast and mammalian cells, *e.g.*, Chinese hamster ovary (CHO) cells (ATCC Accession No. CCL61), NIH Swiss mouse embryo cells NIH-3T3 (ATCC Accession No. CRL1658), and baby hamster kidney cells (BHK). Other useful eukaryotic host cells include insect cells and plant cells. Exemplary prokaryotic host cells are *E. coli* and *Streptomyces*.
- [0287] Transformation of host cells can be accomplished by conventional methods suited to the vector and host cell employed. For transformation of prokaryotic host cells, electroporation and salt treatment methods can be employed (Cohen *et al.*, *Proc. Natl. Acad. Sci. USA* 69:2110-14 (1972)). For transformation of vertebrate cells, electroporation, cationic lipid or salt treatment methods can be employed. *See, e.g.*, Graham *et al.*, *Virology* 52:456-467 (1973); Wigler *et al.*, *Proc. Natl. Acad. Sci. USA* 76:1373-76 (1979).
- [0288] In certain embodiments, the host cell line used for protein expression is can be of mammalian origin; those skilled in the art are credited with ability to determine particular

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host cell lines which are best suited for the desired gene product to be expressed therein. Exemplary mammalian host cell lines include, but are not limited to, NSO, SP2 cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells (*e.g.*, Hep G2), A549 cells DG44 and DUXB11 (Chinese Hamster Ovary lines, DHFR minus), HELA (human cervical carcinoma), CVI (monkey kidney line), COS (a derivative of CVI with SV40 T antigen), R1610 (Chinese hamster fibroblast) BALBC/3T3 (mouse fibroblast), HAK (hamster kidney line), SP2/O (mouse myeloma), P3x63-Ag3.653 (mouse myeloma), BFA-1c1BPT (bovine endothelial cells), RAJI (human lymphocyte) and 293 (human kidney). Host cells for expression of a LINGO-1 antagonist or TrkB agonist for use in a method of the invention may also be prokaryotic. Exemplary prokaryotic host cells are *E. coli* and *Streptomyces*. Host cell lines are typically available from commercial services, the American Tissue Culture Collection or from published literature.

- [0289] Expression of polypeptides from production cell lines can be enhanced using known techniques. For example, the glutamine synthetase (GS) system is commonly used for enhancing expression under certain conditions. *See, e.g.*, European Patent Nos. 0 216 846, 0 256 055, and 0 323 997 and European Patent Application No. 89303964.4.

Gene Therapy

- [0290] A LINGO-1 and certain TrkB agonist (*e.g.* polynucleotide, polypeptide, antibodies and aptamers) can be produced *in vivo* in a mammal, *e.g.*, a human patient, using a gene-therapy approach to treatment of a disease, disorder or injury associated with neuronal degeneration, death or lack of regeneration. This involves administration of a suitable LINGO-1 antagonist and/or TrkB agonist-encoding nucleic acid operably linked to suitable expression control sequences. Generally, these sequences are incorporated into a viral vector. Suitable viral vectors for such gene therapy include an adenoviral vector, an alphavirus vector, an enterovirus vector, a pestivirus vector, a lentiviral vector, a baculoviral vector, a herpesvirus vector (*e.g.* an Epstein Barr viral vector, or a herpes simplex viral vector) a papovaviral vector, a poxvirus vector (*e.g.* a vaccinia viral vector) and a parvovirus. The viral vector can be a replication-defective viral vector. Adenoviral vectors that have a deletion in their E1 gene or E3 genes are typically used. When an adenoviral vector is used, the vector usually does not have a selectable marker gene.

Pharmaceutical Compositions and Administrative Methods

[0291] The LINGO-1 antagonist or TrkB agonists used in the methods of the invention may be formulated into pharmaceutical compositions for administration to mammals, including humans. The pharmaceutical compositions used in the methods of this invention comprise pharmaceutically acceptable carriers, including, *e.g.*, ion exchangers, alumina, aluminum stearate, lecithin, serum proteins, such as human serum albumin, buffer substances such as phosphates, glycine, sorbic acid, potassium sorbate, partial glyceride mixtures of saturated vegetable fatty acids, water, salts or electrolytes, such as protamine sulfate, disodium hydrogen phosphate, potassium hydrogen phosphate, sodium chloride, zinc salts, colloidal silica, magnesium trisilicate, polyvinyl pyrrolidone, cellulose-based substances, polyethylene glycol, sodium carboxymethylcellulose, polyacrylates, waxes, polyethylene-polyoxypropylene-block polymers, polyethylene glycol and wool fat.

[0292] The compositions used in the methods of the present invention may be administered by any suitable method, *e.g.*, parenterally, intraventricularly, orally, by inhalation spray, topically, rectally, nasally, buccally, vaginally or via an implanted reservoir. The term "parenteral" as used herein includes subcutaneous, intravenous, intramuscular, intra-articular, intra-synovial, intrasternal, intrathecal, intrahepatic, intralesional and intracranial injection or infusion techniques. As described previously, LINGO-1 antagonist or TrkB agonists used in the methods of the invention act in the nervous system to promote survival of neurons. Accordingly, in the methods of the invention, the LINGO-1 antagonist or TrkB agonists are administered in such a way that they cross the blood-brain barrier. This crossing can result from the physico-chemical properties inherent in the LINGO-1 antagonist or TrkB agonist molecule itself, from other components in a pharmaceutical formulation, or from the use of a mechanical device such as a needle, cannula or surgical instruments to breach the blood-brain barrier. Where the LINGO-1 antagonist or TrkB agonist is a molecule that does not inherently cross the blood-brain barrier, *e.g.*, a fusion to a moiety that facilitates the crossing, suitable routes of administration are, *e.g.*, intrathecal or intracranial, *e.g.*, directly into a chronic lesion of MS. Where the LINGO-1 antagonist or TrkB agonist is a molecule that inherently crosses the blood-brain barrier, the route of administration may be by one or more of the various routes described below.

[0293] Sterile injectable forms of the compositions used in the methods of this invention may be aqueous or oleaginous suspension. These suspensions may be formulated

according to techniques known in the art using suitable dispersing or wetting agents and suspending agents. The sterile, injectable preparation may also be a sterile, injectable solution or suspension in a non-toxic parenterally acceptable diluent or solvent, for example as a suspension in 1,3-butanediol. Among the acceptable vehicles and solvents that may be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any bland fixed oil may be employed including synthetic mono- or di-glycerides. Fatty acids, such as oleic acid and its glyceride derivatives are useful in the preparation of injectables, as are natural pharmaceutically acceptable oils, such as olive oil or castor oil, especially in their polyoxyethylated versions. These oil solutions or suspensions may also contain a long-chain alcohol diluent or dispersant, such as carboxymethyl cellulose or similar dispersing agents which are commonly used in the formulation of pharmaceutically acceptable dosage forms including emulsions and suspensions. Other commonly used surfactants, such as Tweens, Spans and other emulsifying agents or bioavailability enhancers which are commonly used in the manufacture of pharmaceutically acceptable solid, liquid, or other dosage forms may also be used for the purposes of formulation.

[0294] Parenteral formulations may be a single bolus dose, an infusion or a loading bolus dose followed with a maintenance dose. These compositions may be administered at specific fixed or variable intervals, *e.g.*, once a day, or on an "as needed" basis.

[0295] Certain pharmaceutical compositions used in the methods of this invention may be orally administered in an acceptable dosage form including, *e.g.*, capsules, tablets, aqueous suspensions or solutions. Certain pharmaceutical compositions also may be administered by nasal aerosol or inhalation. Such compositions may be prepared as solutions in saline, employing benzyl alcohol or other suitable preservatives, absorption promoters to enhance bioavailability, and/or other conventional solubilizing or dispersing agents.

[0296] The amount of a LINGO-1 antagonist and/or TrkB agonist that may be combined with the carrier materials to produce a single dosage form will vary depending upon the host treated, the type of antagonist used and the particular mode of administration. The composition may be administered as a single dose, multiple doses or over an established period of time in an infusion. Dosage regimens also may be adjusted to provide the optimum desired response (*e.g.*, a therapeutic or prophylactic response).

[0297] The methods of the invention use a “therapeutically effective amount” or a “prophylactically effective amount” of a LINGO-1 antagonist or TrkB agonist. Such a therapeutically or prophylactically effective amount may vary according to factors such as the disease state, age, sex, and weight of the individual. A therapeutically or prophylactically effective amount is also one in which any toxic or detrimental effects are outweighed by the therapeutically beneficial effects.

[0298] A specific dosage and treatment regimen for any particular patient will depend upon a variety of factors, including the particular LINGO-1 antagonist or TrkB agonist used, the patient’s age, body weight, general health, sex, and diet, and the time of administration, rate of excretion, drug combination, and the severity of the particular disease being treated. Judgment of such factors by medical caregivers is within the ordinary skill in the art. The amount will also depend on the individual patient to be treated, the route of administration, the type of formulation, the characteristics of the compound used, the severity of the disease, and the desired effect. The amount used can be determined by pharmacological and pharmacokinetic principles well known in the art.

[0299] In the methods of the invention the LINGO-1 antagonist or TrkB agonists are generally administered directly to the nervous system, intracerebroventricularly, or intrathecally, *e.g.* into a chronic lesion of MS. Compositions for administration according to the methods of the invention can be formulated so that a dosage of 0.001 – 10 mg/kg body weight per day of the LINGO-1 antagonist polypeptide is administered. In some embodiments of the invention, the dosage is 0.01 – 1.0 mg/kg body weight per day. In some embodiments, the dosage is 0.001 – 0.5 mg/kg body weight per day.

[0300] For treatment with a LINGO-1 antagonist or TrkB agonist antibody, the dosage can range, *e.g.*, from about 0.0001 to 100 mg/kg, and more usually 0.01 to 5 mg/kg (*e.g.*, 0.02 mg/kg, 0.25 mg/kg, 0.5 mg/kg, 0.75 mg/kg, 1 mg/kg, 2 mg/kg, etc.), of the host body weight. For example, dosages can be 1 mg/kg body weight or 10 mg/kg body weight or within the range of 1-10 mg/kg, optionally at least 1 mg/kg. Doses intermediate in the above ranges are also intended to be within the scope of the invention. Subjects can be administered such doses daily, on alternative days, weekly or according to any other schedule determined by empirical analysis. An exemplary treatment entails administration in multiple dosages over a prolonged period, for example, of at least six months. Additional exemplary treatment regimes entail administration once every two weeks or once a month or once every 3 to 6 months. Exemplary dosage schedules include 1-10 mg/kg or 15 mg/kg on consecutive days, 30 mg/kg on alternate days or 60

mg/kg weekly. In some methods, two or more monoclonal antibodies with different binding specificities are administered simultaneously, in which case the dosage of each antibody administered falls within the ranges indicated.

[0301] In certain embodiments, a subject can be treated with a nucleic acid molecule encoding a LINGO-1 antagonist or TrkB agonist polynucleotide. Doses for nucleic acids range from about 10 ng to 1 g, 100 ng to 100 mg, 1 µg to 10 mg, or 30-300 µg DNA per patient. Doses for infectious viral vectors vary from 10-100, or more, virions per dose.

[0302] In certain embodiments, LINGO-1 antagonists may be administered in an amount effective to block interaction of LINGO-1 and TrkB by at least 5%, 10%, 20%, 25%, 30%, 40%, 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95% or 100% as measured by a competition assay or immunoprecipitation assay as compared to the interaction of LINGO-1 and TrkB in the absence of LINGO-1 antagonists.

[0303] In certain embodiments, LINGO-1 antagonists may be administered in an amount effective to promote phosphorylation of TrkB by at least 5%, 10%, 20%, 25%, 30%, 40%, 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95% or 100% as compared to the amount of phosphorylated TrkB in the absence of LINGO-1 antagonists.

[0304] In certain embodiments, LINGO-1 antagonists may be administered in an amount effective to decrease JNK phosphorylation by 5%, 10%, 20%, 25%, 30%, 40%, 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95% or 100% as compared to the amount of phosphorylated JNK in the absence of LINGO-1 antagonists.

[0305] In certain embodiments, TrkB agonists and LINGO-1 antagonists may be administered in an amount effective to promote survival of a CNS neuron by an increase in the number of surviving neurons of at least 5%, 10%, 20%, 25%, 30%, 40%, 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95%, 100%, 110%, 120%, 130%, 140%, 150%, 160%, 170%, 180%, 190%, 200%, 250%, 300%, 350%, 400%, 450%, 500%, 550%, 600%, 650%, 700%, 750%, 800%, 850%, 900%, 950%, or 1000% as compared to the number of surviving neurons in an untreated CNS neuron or mammal.

[0306] Supplementary active compounds also can be incorporated into the compositions used in the methods of the invention. For example, a soluble LINGO-1 or TrkB-agonist polypeptide or a fusion protein may be coformulated with and/or coadministered with one or more additional therapeutic agents.

[0307] The invention encompasses any suitable delivery method for a LINGO-1 antagonist or TrkB agonist to a selected target tissue, including bolus injection of an

aqueous solution or implantation of a controlled-release system. Use of a controlled-release implant reduces the need for repeat injections.

[0308] The LINGO-1 antagonist or TrkB agonist used in the methods of the invention may be directly infused into the brain. Various implants for direct brain infusion of compounds are known and are effective in the delivery of therapeutic compounds to human patients suffering from neurological disorders. These include chronic infusion into the brain using a pump, stereotactically implanted, temporary interstitial catheters, permanent intracranial catheter implants, and surgically implanted biodegradable implants. *See, e.g., Gill et al., supra; Scharfen et al., "High Activity Iodine-125 Interstitial Implant For Gliomas," Int. J. Radiation Oncology Biol. Phys. 24(4):583-591 (1992); Gaspar et al., "Permanent 125I Implants for Recurrent Malignant Gliomas," Int. J. Radiation Oncology Biol. Phys. 43(5):977-982 (1999); chapter 66, pages 577-580, Bellezza et al., "Stereotactic Interstitial Brachytherapy," in Gildenberg et al., Textbook of Stereotactic and Functional Neurosurgery, McGraw-Hill (1998); and Brem et al., "The Safety of Interstitial Chemotherapy with BCNU-Loaded Polymer Followed by Radiation Therapy in the Treatment of Newly Diagnosed Malignant Gliomas: Phase I Trial," J. Neuro-Oncology 26:111-23 (1995).*

[0309] The compositions may also comprise a LINGO-1 antagonist or TrkB agonist dispersed in a biocompatible carrier material that functions as a suitable delivery or support system for the compounds. Suitable examples of sustained release carriers include semipermeable polymer matrices in the form of shaped articles such as suppositories or capsules. Implantable or microcapsular sustained release matrices include polylactides (U.S. Patent No. 3,773,319; EP 58,481), copolymers of L-glutamic acid and gamma-ethyl-L-glutamate (Sidman et al., *Biopolymers* 22:547-56 (1985)); poly(2-hydroxyethyl-methacrylate), ethylene vinyl acetate (Langer et al., *J. Biomed. Mater. Res.* 15:167-277 (1981); Langer, *Chem. Tech.* 12:98-105 (1982)) or poly-D-(-)-3hydroxybutyric acid (EP 133,988).

[0310] In some embodiments of the invention, a LINGO-1 antagonist or TrkB agonist is administered to a patient by direct infusion into an appropriate region of the brain. *See, e.g., Gill et al., "Direct brain infusion of glial cell line-derived neurotrophic factor in Parkinson disease," Nature Med.* 9:589-95 (2003). Alternative techniques are available and may be applied to administer a LINGO-1 antagonist according to the invention. For example, stereotactic placement of a catheter or implant can be accomplished using the Riechert-Mundinger unit and the ZD (Zamorano-Dujovny) multipurpose localizing unit.

A contrast-enhanced computerized tomography (CT) scan, injecting 120 ml of omnipaque, 350 mg iodine/ml, with 2 mm slice thickness can allow three-dimensional multiplanar treatment planning (STP, Fischer, Freiburg, Germany). This equipment permits planning on the basis of magnetic resonance imaging studies, merging the CT and MRI target information for clear target confirmation.

[0311] The Leksell stereotactic system (Downs Surgical, Inc., Decatur, GA) modified for use with a GE CT scanner (General Electric Company, Milwaukee, WI) as well as the Brown-Roberts-Wells (BRW) stereotactic system (Radionics, Burlington, MA) can be used for this purpose. Thus, on the morning of the implant, the annular base ring of the BRW stereotactic frame can be attached to the patient's skull. Serial CT sections can be obtained at 3 mm intervals through the (target tissue) region with a graphite rod localizer frame clamped to the base plate. A computerized treatment planning program can be run on a VAX 11/780 computer (Digital Equipment Corporation, Maynard, Mass.) using CT coordinates of the graphite rod images to map between CT space and BRW space.

[0312] The methods of treatment of disorders as described herein are typically tested *in vitro*, and then *in vivo* in an acceptable animal model, for the desired therapeutic or prophylactic activity, prior to use in humans. Suitable animal models, including transgenic animals, are well known to those of ordinary skill in the art. For example, *in vitro* assays to demonstrate the survival effect of the LINGO-1 antagonist or TrkB agonists are described herein. Finally, *in vivo* tests can be performed by creating transgenic mice which express the LINGO-1 antagonist or TrkB agonist or by administering the LINGO-1 antagonist or TrkB agonist to mice or rats in models as described herein.

[0313] The practice of the present invention will employ, unless otherwise indicated, conventional techniques of cell biology, cell culture, molecular biology, transgenic biology, microbiology, recombinant DNA, and immunology, which are within the skill of the art. Such techniques are explained fully in the literature. See, for example, Molecular Cloning: A Laboratory Manual (3-Volume Set), J. Sambrook, D. W. Russell, Cold Spring Harbor Laboratory Press (2001); Genes VIII, B. Lewin, Prentice Hall (2003); PCR Primer, C.W. Dieffenbach and G.S. Dveksler, CSHL Press (2003); DNA Cloning, D. N. Glover ed., Volumes I and II (1985); Oligonucleotide Synthesis: Methods and Applications (Methods in Molecular Biology), P. Herdewijn (Ed.), Humana Press (2004); Culture of Animal Cells: A Manual of Basic Technique, 4th edition, R. I. Freshney, Wiley-Liss (2000); Oligonucleotide Synthesis, M. J. Gait (Ed.), (1984); Mullis *et al.* U.S.

Pat. No: 4,683,195; Nucleic Acid Hybridization, B. D. Hames & S. J. Higgins eds. (1984); Nucleic Acid Hybridization, M. L. M. Anderson, Springer (1999); Animal Cell Culture and Technology, 2nd edition, M. Butler, BIOS Scientific Publishers (2004); Immobilized Cells and Enzymes: A Practical Approach (Practical Approach Series), J. Woodward, Irl Pr (1992); Transcription And Translation, B. D. Hames & S. J. Higgins (Eds.) (1984); Culture Of Animal Cells, R. I. Freshney, Alan R. Liss, Inc., (1987); Immobilized Cells And Enzymes, IRL Press, (1986); A Practical Guide To Molecular Cloning, 3rd edition, B. Perbal, John Wiley & Sons Inc. (1988); the treatise, Methods In Enzymology, Academic Press, Inc., N.Y.; Gene Transfer Vectors For Mammalian Cells, J. H. Miller and M. P. Calos eds., Cold Spring Harbor Laboratory (1987); Methods In Enzymology, Vols. 154 and 155, Wu *et al.* (Eds.); Immunochemical Methods In Cell And Molecular Biology, Mayer and Walker, (Eds.), Academic Press, London (1987); Handbook Of Experimental Immunology, Volumes I-IV, D. M. Weir and C. C. Blackwell (Eds.), (1986); Immunology Methods Manual: The Comprehensive Sourcebook of Techniques (4 Volume Set), 1st edition, I. Lefkovits, Academic Press (1997); Manipulating the Mouse Embryo: A Laboratory Manual, 3rd edition, Cold Spring Harbor Laboratory Press (2002); and in Ausubel *et al.*, Current Protocols in Molecular Biology, John Wiley and Sons, Baltimore, Maryland (1989).

- [0314] General principles of antibody engineering are set forth in Antibody Engineering: Methods and Protocols (Methods in Molecular Biology), B.L. Lo (Ed.), Humana Press (2003); Antibody engineering, R. Kontermann and S. Dubel (Eds.), Springer Verlag (2001); Antibody Engineering, 2nd edition, C.A.K. Borrebaeck (Ed.), Oxford Univ. Press (1995). General principles of protein engineering are set forth in Protein Engineering, A Practical Approach, Rickwood, D., *et al.* (Eds.), IRL Press at Oxford Univ. Press, Oxford, Eng. (1995). General principles of antibodies and antibody-hapten binding are set forth in: Antibodies: A Laboratory Manual, E. Harlow and D. Lane, Cold Spring Harbor Laboratory Press (1988); Nisonoff, A., Molecular Immunology, 2nd edition, Sinauer Associates, Sunderland, MA (1984); and Steward, M.W., Antibodies, Their Structure and Function, Chapman and Hall, New York, NY (1984). Additionally, standard methods in immunology known in the art and not specifically described are generally followed as in Current Protocols in Immunology, John Wiley & Sons, New York; Stites *et al.* (Eds.) , Immunochemical Protocols (Methods in Molecular Biology), 2nd edition, J. D. Pound (Ed.), Humana Press (1998), Weir's Handbook of Experimental Immunology, 5th edition, D. M. Weir (Ed.), Blackwell Publishers (1996), Methods in

Cellular Immunology, 2nd edition, R. Fernandez-Botran, CRC Press (2001); Basic and Clinical Immunology, 8th edition, Appleton & Lange, Norwalk, CT (1994) and Mishell and Shiigi (Eds.), Selected Methods in Cellular Immunology, W.H. Freeman and Co., New York (1980).

- [0315] Standard reference works setting forth general principles of immunology include Current Protocols in Immunology, John Wiley & Sons, New York; Klein, J.; Kuby Immunology, 4th edition, R. A. Goldsby, *et al.*, H. Freeman & Co. (2000); Basic and Clinical Immunology, M. Peakman, *et al.*, Churchill Livingstone (1997); Immunology, 6th edition, I. Roitt, *et al.*, Mosby, London (2001); Cellular and Molecular Immunology, 5th edition; A.K. Abbas, A.H. Lichtman, Elsevier - Health Sciences Division (2005); Immunology Methods Manual: The Comprehensive Sourcebook of Techniques (4 Volume Set), 1st edition, I. Lefkovits, Academic Press (1997) Immunology, 5th edition, R.A. Goldsby, *et al.*, W. H. Freeman (2002); Monoclonal Antibodies : Principles and Practice, 3rd Edition , J.W. Goding, Academic Press (1996); Immunology: The Science of Self-Nonself Discrimination, John Wiley & Sons, New York (1982); Kennett, R., *et al.* (Eds.), Monoclonal Antibodies, Hybridoma: A New Dimension in Biological Analyses, Plenum Press, New York (1980); Campbell, A., "Monoclonal Antibody Technology" in Burden, R., *et al.* (Eds.), Laboratory Techniques in Biochemistry and Molecular Biology, Vol. 13, Elsevier, Amsterdam (1984).
- [0316] All of the references cited above, as well as all references cited herein, are incorporated herein by reference in their entireties.

EXAMPLES

MATERIALS AND METHODS

Generation of Recombinant LINGO-1-Fc and anti-LINGO-1 Monoclonal Antibody

- [0317] LINGO-1-Fc (LINGO-1-Fc protein) was prepared as described previously (*Mi et al.*, *Nat. Neurosci.* 7:221-228 (2004)). Residues 1-532 of human LINGO-1 were fused to the hinge and Fc region of human IgG1 and expressed in CHO cells. Human IgG1 (control protein) was purchased from Protos Immunoresearch (San Francisco, CA). The anti-LINGO-1 mab 1A7 was generated in mice immunized with LINGO-1-Fc. The hybridoma cell line was grown in DMEM and the antibody was purified by Protein A Sepharose. MOPC21 Mouse IgG (control protein) was purchased from Protos Immunoresearch (San Francisco, CA).

Ocular Hypertension Model

- [0318] Experiments were carried out according to the National Institutes of Health Guide for the Care and Use of Laboratory Animals (NIH Publications No. 80-23) revised in 1996 and approved by the University of Hong Kong Animal Ethics Committee. Adult female Sprague-Dawley (SD) rats weighing approximately 250 g were used. They were housed 3 per standard laboratory cage and maintained on food and water ad libitum with a 12-h dark/light cycle (7:00 a.m./7:00 p.m.). All operations were carried out in animals anesthetized with intraperitoneal injection of ketamine (80 mg/kg) and xylazine (8 mg/kg). 0.5% alcaine (Alcon-Couvreur, Belgium) was applied to the eyes before all operations and antiseptic eye drops (Tobres [Tobramycin 0.3%], Alcon-Couvreur, Belgium) were used to prevent infection after the treatment. Rimadyl (0.025mg/ml) in drinking water was used to relieve the pain for 7 days after the surgeries.
- [0319] Experimental glaucoma was induced using a chronic hypertension model. SD rats received argon laser photocoagulation of the episcleral and limbal veins in the right eye at a power of 1000 mW, a spot size of 500-100 μ m and a duration of 0.1s (Ji *et al.*, *Eur. J. Neurosci.* 19:265-272 (2004); WoldeMussie *et al.*, *Invest. Ophthalmol.* 42:2849-2855 (2001)). About 90 spots were applied on the three episcleral veins and 70 spots around the limbal vein. A secondary laser surgery was delivered to block the reconnected vascular flow seven days later. The animals' left eyes were used as a contralateral control and were not operated on. Animals were allowed to survive for 2 or 4 weeks post first laser exposure before they were sacrificed. The IOP of right and left eyes were measured using a Tonopen XL Tonometer at different time points after laser operation. An average of ten measurements was obtained for each eye. FG labeling of RGCs was performed four days before sacrifice. Both superior colliculi (SC) were exposed after removing a small piece of skull and cortex, and a piece of Gelfoam (Pharmacia & Upjohn) soaked with FG (6% v/v, Fluorochrome, Denver, CO) was placed on the surface of the SC. FG retrogradely labeled intact RGCs. Twelve animals were treated with PBS and allowed to survive 4 weeks post first laser exposure. In all other experimental groups, ten animals were used. The procedure of glaucoma model is summarized in Figure 1.
- [0320] Following the first laser treatment, animals immediately received an intravitreal injection of 2 μ g LINGO-1-Fc, 2 μ g 1A7 or 2 μ g control protein in PBS. In the 4-week glaucoma model, the proteins were re-injected once a week. Treatments were masked to avoid bias of investigators during counting of RGCs. All animals were euthanized with an overdose of anesthesia.

[0321] At predefined times, rats were sacrificed with an overdose of anesthesia. Both eyes of each animal were enucleated and fixed in 4% paraformaldehyde for 60 minutes. Retinas were prepared as flat-mounts and the FG labeled RGCs were counted under fluorescence microscopy using an ultra-violet filter (excitation wave length = 330-380 nm) as described (Cheung *et al.*, *Mol. Cell. Neurosci.* 25:383-393 (2004); Ji *et al.*, *Eur. J. Neurosci.* 19:265-272 (2004)). The RGCs were quantified under an eyepiece grid of $200 \times 200 \mu\text{m}^2$ along the median line of each quadrant, starting from the optic disc to the border at 500 μm intervals (Figure 2). Eight microscopic fields for each quadrant and a total of 32 per retina for four quadrants were counted, corresponding to approximately 3-3.2% of each retinal area. Percent loss of RGCs was measured to examine the survival effects of different treatments. The data was expressed in terms of relative percentage of RGC loss in the injured eye compared to the contralateral intact eye (% contralateral, mean + sem).

Tissue Processing for Transmission Electron Microscopy

[0322] Two mm segments of the retinas were obtained from normal and 2 week-glaucoma groups treated with PBS, soluble LINGO-1 or 1A7. They were placed in Karnovsky electron microscope (EM) fixative for 2-4 hours at 4°C. After washing in 0.1M phosphate buffer (PB), tissue samples were post-fixed in 1% osmium tetroxide in 0.1M PB, then dehydrated in ethanol and embedded in Epon. Semithin (1 μm) sections were obtained from each of the blocks using a Reichert-Jung ultramicrotome with glass knives made on a LKB knife-maker. The sections were stained with toluidine blue. Ultra-thin sections with silver interference color were obtained, stained with Reynolds lead citrate and uranyl acetate and examined with electron microscope.

Primary RGC Culture

[0323] The eyes of P7 Long-Evans rats were removed and placed in dissociation medium (DM; 90 mM Na_2SO_4 , 30 mM K_2SO_4 , 5.8 mM MgCl_2 , 0.25 mM CaCl_2 , 1 mM HEPES, 0.001% Phenol red) (Furshpan and Potter, 1989). Retinas were removed from the eyes and incubated in 2 ml DM containing 15 U/ml Papain (Worthington, NJ), 1 mM L-cysteine, 0.5 mM EDTA, 0.005% DNase I for 30 min at 37°C. Retinas were rinsed twice in DM, resuspended in 2 ml DM containing 10 mg/ml ovomucoid protease inhibitor, 10 mg/ml Bovine Serum Albumin (both from Worthington) and gently triturated 10-15 times using a 2 ml serological pipette. Dissociated cells were pelleted at 1000 rpm for 5 min,

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resuspended in 2 ml of growth medium modified from Meyer-Franke *et al.* (Meyer-Franke *et al. Neuron* 15:805-819 (1995)) (Neurobasal, 1x B27 from Invitrogen, 5 μ M Forskolin, 60 nM T3, 1mM Pyruvate, 2 mM Glutamine) and counted with a hemocytometer. Cells were plated at a density of 50,000 cells/cm² in BD Biocoat 96-well tissue culture plates coated with poly-D-Lysine and Laminin (BD Biosciences, Bedford, MA) and grown for 3 days in a CO₂ incubator. At the time of plating, cells were treated in quadruplicate wells with human IgG1 (10 μ g/ml), LINGO-1-Fc (10 μ g/ml) alone, human IgG1 and 25 ng/ml BDNF, or LINGO-1 Fc and BDNF.

[0324] Cells were fixed 15 min in methanol at -20°C, rinsed twice in PBS and blocked for 1 h in 5% normal goat serum in PBS for 1 h. Cells were incubated in Thy1.1 antibody (Serotec, clone OX-7, 1:40 in PBS) overnight at 4°C, rinsed 3 times for 5 min in PBS, incubated in goat anti-mouse IgG-conjugated with Alexa 594 (Molecular Probes, Eugene, OR; 1:1000) for 1 h and rinsed 3 times for 5 min in PBS.

[0325] RGCs were identified under epifluorescence microscopy using a Zeiss axiovert inverted microscope. The entire surface of each well was visually scanned under epifluorescence to count surviving RGCs. Only Thy 1.1 positive cells with neuronal morphology and bearing at least one process with a minimum length 3 times the cell body diameter were counted.

Immunohistochemistry for LINGO-1 and p-TrkB

[0326] RGCs were retrogradely labeled with FG at 4 days before sacrifice. The eyes were enucleated at 2 weeks after injury following transcardial perfusion with 0.9% saline and subsequently post-fixed in 4% PFA for 4 h. Ten-micron-thick frozen sections were incubated with mouse anti-LINGO-1 (Biogen), rabbit anti-phosphor-TrkB (Tyr785) (a gift from Dr. B. Sun, Shanghai Institutes of Biological Sciences, Shanghai, China) antibodies (Ji *et al., Nat. Neurosci.* 8:164-172 (2005)), following by treatment with Alexa-labeled secondary antibody. After washing, the sections were mounted with fluorescent mounting medium (DakoCytomation) and analyzed under Carl Zeiss LSM 510 META Confocal microscopy.

Western Blotting

[0327] To measure LINGO-1, BDNF or p-TrkB in the retina, the animals were euthanized at 2 weeks after laser coagulation. For assessing the temporal profile of Akt phosphorylation, animals treated with LINGO-1-Fc or PBS were euthanized at 6 hr, 1 day

and 5 days after laser coagulation. To measure the effects of injury, BDNF, and LINGO-Fc or 1A7 on TrkB phosphorylation, we injected BDNF intravitreally (5 µg/eye, recombinant human BDNF; Regeneron Pharmaceutical, Tarrytown, NY) or BDNF combined with LINGO-1-Fc or 1A7 (2 µg/eye) at the same time of laser coagulation and then euthanized the animals 5 days later. Retinas were dissected and homogenized in lysis buffer (10 mM Tris pH7.4, 150 mM NaCl, 1 mM EDTA, 1 mM EGTA) supplemented with 10% protease inhibitor cocktail and 1% phosphatase inhibitor cocktails from Sigma. Following centrifugation at 13,000 rpm for 30 minutes to remove cell debris, the protein concentration of the supernatant was measured using a Bio-Rad DC protein Assay Kit (Bio-Rad Laboratories, CA, USA). A 40-80 µg aliquot of proteins from individual animals was subjected to 6-12.5% SDS-polyacrylamide gel electrophoresis and transferred onto PVDF membrane. The membranes were blocked with 5% non fat dry milk and 2% bovine serum albumin (BSA) in Tris-buffered saline containing 0.1% Tween 20 (TBST) for 1 h at room temperature. Incubations with mouse anti-LINGO-1 (Biogen), rabbit anti-BDNF (Chemicon), mouse anti-phosphor-Akt (1:1000), rabbit total Akt (1:1000), anti-phospho-JNK (1:1000), total-JNK (Cell Signaling Technology), rabbit anti-phosphor-TrkB (Tyr785) (1:1000) (a gift from Dr. B. Sun, Shanghai Institutes of Biological Sciences, Shanghai, China) (Ji *et al.*, *Nat. Neurosci.* 8:164-172 (2005)) and chicken IgY total TrkB (Promega) (1:100) antibodies were performed overnight at 4°C. After washing, the membranes were incubated with Horseradish Peroxidase-conjugated secondary antibody in 5% non fat dry milk and 2% BSA in TBST for 1 h at room temperature. Immunoreactive proteins were detected using the enhanced chemiluminescence method (ECL, Amersham). Protein loading was controlled using a monoclonal goat antibody against actin (1:1000, C-11, Santa Cruz Biotechnology). The intensity of each band was quantified by densitometric scanning using Labworks gel documentation (UVP, Inc, Upland, CA). All experiments for Western blotting were performed with 3-5 animals in each group and the samples were run on the gels as individual animals. Protein levels were finally expressed as relative values compared to total proteins of normal retinas.

Immunoprecipitations and Western Blotting for LINGO-1 and TrkB

[0328] 293T cells (100 mm dishes) were transfected with HA-tagged full length human LINGO-1, myc-tagged full length human TrkB, or a combination of LINGO-1/TrkB. The cells were harvested after 48 h and lysed in 1 ml RIPA buffer (50 mM Tris, pH 7.2,

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1% Triton X-100, 0.5% sodium deoxycholate, 0.1% SDS, 150 mM NaCl, 10 mM MgCl₂, 5% glycerol) for 30 min at 4°C. After centrifugation at 14,000 x g for 15 min, the supernatants were incubated with ProteinA/G plus-Sepharose beads (Santa Cruz Biotechnology, CA) at 4°C for 1 hr. The pre-cleared lysates were then incubated with an anti-LINGO-1 antibody (Biogen Idec) at 4°C for 1 hr followed by addition of Protein A/G-Sepharose beads for 1 hr. The beads were washed 3 times with 1% Triton buffer (50 mM HEPES, pH 7.5, 150 mM NaCl, 1.5 mM MgCl₂, 1mM EGTA, 1% Triton X-100 and 10% glycerol), boiled in Laemmli sample buffer, subjected to 4-20% SDS-PAGE and analyzed by Western blotting with anti-TrkB antibody (Myc, Roche) or anti-LINGO-1 antibody (HA, Roche). Retinal lysates from normal or 2-week ocular hypertensive rats were also immunoprecipitated with anti-TrkB antibody (Chemicon), an anti-LINGO-1 antibody (Upstate), or a non-specific antibody at 4°C overnight and analyzed by Western blotting with an anti-LINGO-1 antibody (Upstate).

[0329] Neuroscreen-1 cells (a subclonal line of PC12 cells, Cellomics) or Neuroscreen cells with over-expressed stable HA-LINGO1 were infected with TrkB lentiviruses for 2 days. The cells were serum-starved overnight before they were subjected to BDNF for 0 or 30 min in serum-free media. The cell lysates were immunoprecipitated by a pan-Trk antibody (Santa Cruz, sc-139), and then assessed by Western blotting using either an anti-Phospho-Tyr antibodies for Phospho-TrkB or an anti-TrkB antibody for total TrkB (Santa Cruz). The cell lysates were also assessed by Western blotting for LINGO-1 expression (HA, Roche).

Statistics

[0330] Statistical analysis was performed using Student's t test for comparisons between two groups, or by one-way analysis of variance (ANOVA) followed by post-hoc tests (Student-Neuman-Keuls) for comparisons of more than two groups.

EXAMPLE 1

Increased Expression of LINGO-1 in a Rat Glaucoma Model

[0331] The expression of LINGO-1 was examined in a rat glaucoma model. In this model, an argon laser was used to block the outflow of aqueous humor by photocoagulation of the limbal and episcleral drainage vessels, resulting in reliable increase of intraocular pressure (IOP) (See Figure 1).

- [0332] To generate a ocular hypertensive state, the limbal and three episcleral veins were photocoagulated twice at a 7-day interval using an Argon laser. Examination of the aqueous veins prior to and immediately after the laser photocoagulation revealed a marked decrease in venous blood flow, as shown in Figure 3.
- [0333] LINGO-1 expression was examined in normal and injured rat retina sections. Normal retinal ganglion cells (RGCs) expressed low level staining for LINGO-1, but much stronger immunoreactivity for LINGO-1 occurred in RGCs at 2 weeks after laser coagulation. The number of labeled RGCs and the intensity of labeling both increased (See Figure 4). (The term "after laser coagulation" means "after first laser coagulation".)
- [0334] These finding were confirmed by western blotting, which showed that LINGO-1 expression was low in normal retina and increased to 1.6 fold at 2 weeks after the injury ($P < 0.05$) (See Figure 5).

EXAMPLE 2

LINGO-1 Antagonists Act as Neuroprotectants Without Affecting Intraocular Pressure

- [0335] Experimental ocular hypertension can be monitored by measuring changes in pressure or neuronal survival. Therefore, the effect of LINGO-1 antagonists on both intraocular pressure and RGC survival was examined. Animals were subjected to laser treatment, as described *supra* and immediately afterwards received an intravitreal injection of LINGO-1-Fc, 1A7 or control protein.
- [0336] The intraocular pressure (IOP) in both treated and untreated eyes was monitored. IOP in the contralateral left eye of treated animals was about 13 mmHg (See Figure 6) and remained at the same level through the experiment. The IOP of the laser-treated right eye in all four groups increased after the first laser surgery, reached approximately 22 mmHg and remained at this level until sacrifice. Under these experimental conditions, treatment with LINGO-1-Fc and the neutralizing anti-LINGO-1 antibody, mAb 1A7, did not lower IOP (See Figure 6).
- [0337] LINGO-1 antagonists did have an affect on neuronal survival. First, surviving RGCs were quantitated 2 weeks after laser injury. Retinas from both eyes were prepared and suviving RGCs were counted. RGC loss at 2 weeks after laser coagulation in PBS and control protein treatment groups was $13.93 \pm 1.44\%$ and $12.37 \pm 1.84 \%$ respectively (See Figure 7A). Injection of LINGO-1-Fc prevented RGC loss. LINGO-1-Fc treated retinas has only $0.09 \pm 1.47 \%$ RGC loss ($P < 0.001$, compared to PBS and human IgG

control groups). Similarly, mAb 1A7 treatment limited RGC loss to $1.46 \pm 1.32 \%$ ($P < 0.001$ compared to the control groups).

[0338] To investigate the effect of LINGO-1 antagonists on the long-term survival of RGCs, LINGO-1-Fc and the neutralizing LINGO-1 antibody 1A7 were injected intravitreally once a week and the animals were allowed to survive for 4 weeks. The results (see Figure 7A) showed that treatment with LINGO-1-Fc antagonists significantly reduced loss of injured RGCs from $20.09 \pm 1.36 \%$ (PBS control) to $5.98 \pm 0.83 \%$ ($P < 0.001$) for LINGO-1-Fc and to $4.73 \pm 1.72 \%$ ($P < 0.001$) for 1A7 4 weeks after laser coagulation.

[0339] The data on RGC survival are also presented as the mean density (No. of cells/mm²) for each group (See Figure 7B). Previous investigations of the death of RGCs in our rat glaucoma model at 2, 4, 8 and 12 weeks after laser coagulation showed that the loss of RGCs reaches a maximal level after 4 weeks (Li *et al.*, *Invest. Ophthalmol. Vis. Sci.* 47:2951-2958 (2006)). In contrast significant neuroprotection of LINGO-1-Fc and 1A7 was observed 4 weeks after laser coagulation. Unlike ciliary neurotrophic factor (CNTF) treatment (data not shown), neither LINGO-1-Fc or anti-LINGO-1 antibody caused cataracts in the long-term survival experiment.

[0340] Electron microscopy was also used to evaluate the effects of LINGO-1 antagonists. As shown in Figure 8A, at two weeks after ocular hypertension, some of the PBS treated RGCs have an irregular or even cracked nucleus with swollen and melting mitochondrion. Some also show a loss of rough endoplasmic reticulum (ER) and Golgi organelles. In contrast, eyes treated with soluble LINGO-1 and 1A7 keep characteristics of typical retinal ganglion cells with normal ER and Golgi organelles. The organelles of cells in the inner plexiform layer (IPL), where the RGCs, amacrine cells and bipolar cells make connections were also examined. Compared to the LINGO-1 or 1A7 treated animals, which maintained an almost normal axons and dendrites, the cells in the PBS treated animals had less organelles inside of more swollen axons and dendrites (See Figure 8B). In addition, the synapses in PBS-treated animals become narrow but still remain un-separated, while the retinas that had received treatment of LINGO-1-Fc or 1A7 maintained normal characteristics of synapses.

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: EXAMPLE 3

LINGO-1 Antagonists Alone Do Not Promote Survival of Retinal Ganglion Cells *In Vitro*

[0341] To study the effects of LINGO-1-Fc on RGC survival further, an RGC primary culture system was used (Meyer-Franke *et al.*, *Neuron*, 15:805-819 (1995)). Dissociated retinal cultures were grown for 3 days in the presence of control protein or LINGO-1-Fc. Primary RGCs were identified by Thy 1.1 immunostaining, a marker for RGCs, and counted. Unlike the evident neuroprotective activity *in vivo*, under these experimental conditions, LINGO-1-Fc treatment alone did not promote survival of cultured RGCs (See Figure 9).

EXAMPLE 4

LINGO-1 Acts on the BDNF Pathway

[0342] The effect of LINGO-1 antagonists on the survival of RGCs *in vitro* in the presence of brain-derived neurotrophic factor (BDNF) was also examined. In the presence of BDNF, LINGO-1 antagonists were able to promote survival of RGCs. RGCs were grown as in Example 3 and treated with either BDNF and control protein or BDNF and LINGO-1-Fc. In the presence of BDNF, a significantly greater percentage of cell survival in cells treated with LINGO-1-Fc than in cells treated with control protein. This result, in addition to the fact that *in vivo*, animal retinas are exposed to endogenous BDNF, while the RGC cultures did not have any BDNF, demonstrates that LINGO-1-Fc rescues RGCs by modulating their response to BDNF, and/or that LINGO-1-Fc indirectly increases survival by enhancing neurotrophin receptors.

[0343] Therefore, in order to investigate the relationship of LINGO-1-Fc and BDNF in the retinas, BDNF was measured in the retina of ocular hypertensive rats using western blotting. Previous studies showed that normal rat retina express BDNF (Rudzinski *et al.*, *J. Neurobiol.* 58:341-351 (2004)). The results showed a low level of BDNF in the normal retina ($P < 0.05$, Figure 10). However, BDNF levels increased more than 3-fold 2 weeks after laser coagulation ($P < 0.05$), consistent with the previously reported results (Rudzinski *et al.*, *J. Neurobiol.* 58:341-351 (2004)). Thus, laser coagulation activated an endogenous neurotrophin response. However, this response was inadequate to protect against neuronal injury completely, as shown in Figure 7. Treatment with LINGO-1-Fc and 1A7 after ocular hypertension did not change the high levels of BDNF compared to the PBS control group ($P < 0.05$ compared to normal, Figure 10). These results and the *in*

in vitro data indicate that LINGO-1-Fc or 1A7 modulates the response to BDNF and reinforces the neuroprotective activity of endogenous BDNF in the glaucomatous retinas.

[0344] A neutralizing anti-BDNF antibody was also injected in the presence of LINGO-1-Fc and 1A7. In these experiments, 3 μ g of the anti-BDNF antibody (Chemicon) was intravitreally injected to the experimental eye on days 0, 3, 7 and 10 after laser coagulation and/or 2 μ g LINGO-1-Fc or 1A7 was administrated once on day 0. The rats were euthanized at 2 weeks and FG-labeled RGCs were counted.

[0345] Anti-BDNF antibody significantly reversed the protective function of LINGO-1-Fc ($P = 0.002$) or 1A7 ($P = 0.004$) at 2 weeks after laser coagulation. There was no difference in the RGC loss between the PBS and the anti-BDNF antibody group or the anti-BDNF antibody combined with LINGO-1-Fc or 1A7 groups (Figure 11A). The data are also presented as the mean density of RGCs (No. of cells/mm²) (Figure 11B). The proteins in each group have no effect on intraocular pressure. These results further confirmed that LINGO-1-Fc or 1A7 rescued the injured RGCs by reinforcing the increased endogenous BDNF in the retina.

EXAMPLE 5

LINGO-1 Binds to and Negatively Regulates TrkB

[0346] As shown in Example 4, LINGO-1-Fc exerts its neuroprotective activity by modulating response to BDNF. The relationship of LINGO-1 and the BDNF receptor TrkB was examined using immunoprecipitation and immunoblotting methods. Neurotrophins activate two different classes of receptors, the Trk family of receptor tyrosine kinases and p75^{NTR} that in turn activate many downstream signaling pathways. Trk receptors include TrkA, B and C. BDNF activates TrkB (*Huang and Reichardt, 2003, Annu. Rev. Biochem. 72:609-642; Huang and Reichardt, 2001, Annu. Rev. Neurosci. 24:677-736*). It has previously been demonstrated that the dominating Trk receptor in the retina is TrkB (*Cui et al., Invest. Ophthalmol. Vis. Sci. 43:1954-1964 (2002)*).

[0347] To determine whether LINGO-1 interacts directly with TrkB, cell lysates from transfected 293T cells co-expressing TrkB and LINGO-1 were examined. The lysates were immunoprecipitated with anti-LINGO-1 antibody, and an anti-TrkB antibody or anti-LINGO-1 antibody were used for western blotting. Our results indicated that TrkB was immunoprecipitated with anti-LINGO-1 antibody from 293T cells coexpressing both TrkB and LINGO-1 (Figure 12A). This indicates that LINGO-1 binds TrkB. In addition,

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a LINGO-1 expressing cell line was transfected with TrkB lentiviruses for 2 days and the cells were treated with BDNF. Lysates were immunoprecipitated using a pan-Trk antibody and then levels of phosphor-TrkB (p-TrkB) and total TrkB were assessed by western blotting. Total TrkB remained at about the same level in all of the groups (Figure 12B). BDNF stimulation resulted in an increase in phosphorylated TrkB in the absence of LINGO-1 (lanes 3 and 4 of Figure 12B). However, levels of p-TrkB were lower in the presence of LINGO-1 after BDNF stimulation (lanes 7 and 8 of Figure 12B, $P < 0.05$). These results indicate that LINGO-1 binds TrkB and limits TrkB activation after TrkB is bound by neurotrophins. LINGO-1 thus modulates the function of TrkB receptors, acting as a negative regulator of the neuroprotective activity of the BDNF-TrkB system.

[0348] In order to determine whether LINGO-1 and TrkB co-immunoprecipitate from RGCs *in vivo*, the retinal lysates from normal or 2-week ocular hypertension rats were immunoprecipitated with an anti-TrkB antibody, an anti-LINGO-1 antibody or a non-specific antibody, and an anti-LINGO-1 antibody was used for western blotting. LINGO-1 was co-immunoprecipitated from the retinal lysates with anti-TrkB antibody indicating that LINGO-1 interacts with TrkB in normal retina. Furthermore, LINGO-1 constitutively bound to TrkB at 2 weeks after the induction of ocular hypertension (Figure 13). The co-localization of LINGO-1 and TrkB in RGCs was also examined. LINGO-1 and p-TrkB were co-expressed in RGCs retrogradely labeled with FG. Figure 13B shows representative photomicrographs of the co-localization of LINGO-1 and p-TrkB in the retinas 2 weeks after the induction of ocular hypertension and treatment with 1A7.

[0349] To further assess the mechanisms of LINGO-1-Fc and 1A7 in the ocular hypertension model, the effects of LINGO-1-Fc and 1A7 on TrkB activation were assessed with western blotting. Normal rat retinas expressed high levels of total TrkB (Figure 14A). The total TrkB level was no different between the control group and LINGO-1-Fc group or the 1A7 group 2 weeks after the induction of ocular hypertension (Figure 14A). However, LINGO-1-Fc or 1A7 administration up-regulated p-TrkB level significantly more than the control protein 2 weeks after laser treatment (Figure 14A) demonstrating that LINGO-1-Fc and 1A7 treatment permit activation of TrkB in the presence of endogenous levels of BDNF. BDNF was after injected after laser coagulation. Total TrkB levels remained at the same level up to 5 days after laser coagulation of the retina, whether the eyes were treated with BDNF alone or with a combination of BDNF and LINGO-1-Fc or 1A7 (Figure 14B). BDNF treatment alone increased p-TrkB levels, but the change was not statistically significant after 5 days of treatment. However, BDNF in

combination with LINGO-1-Fc or with 1A7 significantly increased p-TrkB levels in the retina (Figure 14B).

[0350] These findings indicate that BDNF stimulation alone produced only limited activation of TrkB receptors. This was true even when exogenous BDNF was added. However treatment with LINGO-1-Fc or 1A7 relieved this limitation. These results are consistent with the *in vitro* results showing that LINGO-1 binds with TrkB and negatively regulates TrkB activation after TrkB is bound by BDNF. LINGO-1-Fc and 1A7 treatment promotes neuroprotective activity by increasing BDNF activation of TrkB receptors.

EXAMPLE 6

LINGO-1 Antagonists Increase Akt Activation After Ocular Hypertension

[0351] BDNF activation of TrkB receptors initiates several downstream signaling pathways including PI-3 kinase (PI3K). Phosphorylation of serine 473 and threonine 308 by PI3K are important survival signals for neurons (Brazil *et al.*, *Cell* 111:293-303 (2002)). As shown in Examples 4 and 5, LINGO-1-Fc and 1A7 regulate the function of BDNF and its cognate receptor TrkB. In addition, the effects of LINGO-1-Fc on the PI3K/Akt signaling pathway were examined by measuring the total Akt and pAkt (the active form of Akt) at various times after laser coagulation surgery. Western blot analysis revealed that total Akt levels remained unchanged up to 5 days post laser coagulation (Figure 15). pAkt levels were low in normal control retinas but increased 5-fold by 6 h after laser treatment and then declined from day 1 to day 5 after laser coagulation. A similar bell shaped-response in pAkt levels has been previously reported after optic nerve transection (Cheung *et al.*, *Mol. Cell Neurosci.* 25:383-393 (2004)). In LINGO-1-Fc treated retinas, the levels of pAkt followed a similar pattern, peaking at 6 h and then declining over 5 days after laser coagulation. However, the levels of pAkt on day 5 were significantly higher in LINGO-1-Fc treated retinas than in control retinas treated with PBS ($P < 0.05$, Figure 15). Thus, treatment with LINGO-1-Fc affects Akt signaling.

[0352] To verify that the changes in pAkt levels occurred in RGCs and not in other cells of the retina, pAkt localization was examined by immunohistochemical study of the retinas. pAkt immunoreactivity only in RGCs retrogradely labeled with FG, and the pAkt localization did not change after laser injury and treatment with LINGO-1-Fc. Figure 16 shows representative photomicrographs of pAkt immunolabelling the retinas at 6 h after laser coagulation and treatment with LINGO-1-Fc.

[0353] To further examine the role of pAkt in the neuroprotective activity of LINGO-1-Fc after injury, the effects of an inhibitor of the PI3K/Akt pathway, LY294002 (LY, Calbiochem) were observed. In these experiments, 10 mM LY294002 was dissolved in 100% dimethylsulfoxide (DMSO; Sigma) and subsequently diluted to 2 mM using sterile PBS. 2 mM LY294002 or 2 μ l vehicle (20% DMSO in PBS) were injected intravitreally on days 0, 3, 7 and 10 days after the first laser photocoagulation was performed on the right eye. Animals were euthanized on day 14 and FG-labeled RGCs were counted.

[0354] While LINGO-1-Fc promoted RGC survival at 2 weeks after ocular hypertension, combining LINGO-1-Fc with LY294002 abolished the neuroprotective effect (Figure 17A). Loss of RGCs increased from 0.09 ± 1.47 % in eyes treated with LINGO-1-Fc to 9.0 ± 1.6 % in eyes treated with LINGO-1-Fc and LY294002 ($P = 0.004$). The data are also presented by the density of RGCs (Figure 17B). Neither LY294002 alone or in combination with LINGO-1-Fc lowered intraocular pressure (Figure 18).

EXAMPLE 7

LINGO-1 Antagonists Decrease c-Jun N-terminal Kinase Activation After Ocular Hypertension

[0355] The JNK pathway can be activated by a variety of cellular stresses. More recently, phosphorylated JNK was detected in human and experimental rat glaucomatous retinas, and the activation of JNK was temporally associated with the death of RGCs (Tezel *et al. Invest. Ophthalmol. Vis. Sci.* 44:3025-3033 (2003)). Recent findings also confirmed that phosphorylated JNK is located in the RGCs after IOP elevation. Total JNK-1 and JNK-2 levels remain unchanged up to 5 days after ocular hypertension (Figure 19). Phosphorylated JNK-1,2 were low in the normal retina, then increased to more than 8-fold after 5 days. A similar response in the p-JNK levels following IOP elevation in a rat glaucoma model has been previously reported by immunohistochemistry. After treatment with LINGO-1-Fc or 1A7, the activation of JNK was almost reduced to the normal level ($P < 0.01$) (Figure 19). These results indicate that LINGO-1 antagonists may rescue injured RGCs by inhibiting the activation of JNK signaling pathway.

EXAMPLE 8

LINGO-1 Antagonists Decrease RhoA Activation After Ocular Hypertension

[0356] Since Rho acts downstream of the NgR1-LINGO-1-p75/TROY complex, and JNK is phosphorylated as a downstream consequence of Rho activation, experiments

were also performed to determine if LINGO-1 antagonists have an effect on the activity of Rho after ocular hypertension. Rho activity can be assessed by evaluating GTP-RhoA levels, where increased levels of GTP-RhoA are associated with increased Rho activity. Low levels of GTP-RhoA were observed in normal retina, and the GTP-RhoA levels increased almost twofold five days after laser coagulation ($p < 0.05$ compared with normal group) (Figure 21). In contrast, LINGO-1-Fc treated retinas significantly reduced the high level of GTP-RhoA to the basal level ($p < 0.05$ compared with the PBS group) (Figure 21). These results suggest that LINGO-1-Fc exerts neuroprotective activity by inhibiting RhoA activation.

- [0357] The present invention is not to be limited in scope by the specific embodiments described which are intended as single illustrations of individual aspects of the invention, and any compositions or methods which are functionally equivalent are within the scope of this invention. Indeed, various modifications of the invention in addition to those shown and described herein will become apparent to those skilled in the art from the foregoing description and accompanying drawings. Such modifications are intended to fall within the scope of the appended claims.
- [0358] All publications and patent applications mentioned in this specification are herein incorporated by reference to the same extent as if each individual publication or patent application was specifically and individually indicated to be incorporated by reference.
- [0359] It is to be appreciated that the Detailed Description section, and not the Summary and Abstract sections, is intended to be used to interpret the claims. The Summary and Abstract sections may set forth one or more but not all exemplary embodiments of the present invention as contemplated by the inventor(s), and thus, are not intended to limit the present invention and the appended claims in any way.

WHAT IS CLAIMED IS:

1. A method for inhibiting LINGO-1 and TrkB interaction in a cell comprising contacting a cell that coexpresses LINGO-1 and TrkB with a LINGO-1 antagonist.
2. A method for promoting TrkB phosphorylation in a cell comprising contacting a cell that coexpresses LINGO-1 and TrkB with a LINGO-1 antagonist.
3. A method for promoting TrkB phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.
4. A method for inhibiting JNK phosphorylation in a cell comprising contacting a cell that coexpresses LINGO-1 and JNK with a LINGO-1 antagonist.
5. A method for inhibiting JNK phosphorylation comprising contacting CNS neurons with a LINGO-1 antagonist.
6. A method for promoting survival of a neuron at risk of dying, comprising contacting the neuron with an effective amount of a LINGO-1 antagonist and a TrkB agonist.
7. A method for promoting survival of retinal ganglion cells in a mammal displaying signs or symptoms of a pressure induced ocular neuropathy comprising administering to a mammal in need of such treatment a therapeutically effective amount of a LINGO-1 antagonist and a carrier.
8. A method for promoting survival of retinal ganglion cells in a mammal displaying signs or symptoms of a pressure induced ocular neuropathy comprising administering to a mammal in need of such treatment a therapeutically effective amount of a combination of a LINGO-1 antagonist and a TrkB agonist.
9. A method for treating a disease or disorder associated with neuronal cell death comprising administering to a mammal in need of such treatment an effective amount of a LINGO-1 antagonist and a TrkB agonist.

10. The method of any one of claims 1-5 further comprising contacting the cell with a TrkB agonist.

11. The method of any one of claims 1-10, wherein said LINGO-1 antagonist comprises a soluble LINGO-1 polypeptide.

12. The method of claim 11, wherein said soluble LINGO-1 polypeptide comprises LINGO-1 region selected from the group consisting of:

- (i) a LINGO Ig domain or a fragment, variant, or derivative thereof,
- (ii) a LINGO LRR domain or a fragment, variant, or derivative thereof,
- (iii) a LINGO cytoplasmic domain or a fragment, variant, or derivative thereof,
- (iv) a LINGO transmembrane domain or a fragment, variant, or derivative thereof,
- (v) a LINGO basic region C-terminal to the LRR domain or a fragment, variant, or derivative thereof, and
- (vi) a combination of at least two of said LINGO-1 domains or a fragments, variants, or derivatives thereof of (i) to (v).

13. The method of claim 11, wherein said soluble LINGO-1 polypeptide lacks a LINGO-1 region selected from the group consisting of:

- (i) a LINGO Ig domain,
- (ii) a LINGO LRR domain,
- (iii) a LINGO cytoplasmic domain,
- (iv) a LINGO transmembrane domain,
- (v) a LINGO basic region C-terminal to the LRR domain and
- (vi) a combination of at least two of said LINGO-1 domains of (i) to (v).

14. The method of claim 13, wherein said soluble LINGO-1 polypeptide lacks a LINGO-1 transmembrane domain and a LINGO-1 cytoplasmic domain.

15. The method of any one of claims 11 to 14, wherein said soluble LINGO-1 polypeptide comprises

- (i) a LINGO-1 LRR domain or a fragment, variant, or derivative thereof,

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- (ii) a LINGO-1 basic region C-terminal to the LRR domain or a fragment, variant, or derivative thereof, and
- (iii) a LINGO-1 immunoglobulin (Ig) domain or a fragment, variant, or derivative thereof.

16. The method of claim 13 or claim 14, wherein said soluble LINGO-1 polypeptide comprises at least a portion of the LINGO-1 LRR domain, but lacks a LINGO-1 Ig domain, LINGO-1 basic region, a LINGO-1 transmembrane domain, and a LINGO-1 cytoplasmic domain.

17. The method of any one of claims 11 to 15, wherein said soluble LINGO-1 polypeptide comprises a polypeptide fragment selected from the group consisting of:

- (i) amino acids 1 to 33 of SEQ ID NO:2;
- (ii) amino acids 1 to 35 of SEQ ID NO:2;
- (iii) amino acids 34 to 64 of SEQ ID NO:2;
- (iv) amino acids 36 to 64 of SEQ ID NO:2;
- (v) amino acids 66 to 89 of SEQ ID NO:2;
- (vi) amino acids 90 to 113 of SEQ ID NO:2;
- (vii) amino acids 114 to 137 of SEQ ID NO:2;
- (viii) amino acids 138 to 161 of SEQ ID NO:2;
- (ix) amino acids 162 to 185 of SEQ ID NO:2;
- (x) amino acids 186 to 209 of SEQ ID NO:2;
- (xi) amino acids 210 to 233 of SEQ ID NO:2;
- (xii) amino acids 234 to 257 of SEQ ID NO:2;
- (xiii) amino acids 258 to 281 of SEQ ID NO:2;
- (xiv) amino acids 282 to 305 of SEQ ID NO:2;
- (xv) amino acids 306 to 329 of SEQ ID NO:2;
- (xvi) amino acids 330 to 353 of SEQ ID NO:2;
- (xvii) amino acids 363 to 416 of SEQ ID NO:2;
- (xviii) amino acids 417 to 424 of SEQ ID NO:2;
- (xix) amino acids 419 to 493 of SEQ ID NO:2;
- (xx) amino acids 494 to 551 of SEQ ID NO:2;
- (xxi) amino acids 1 to 64 of SEQ ID NO:2;
- (xxii) amino acids 1 to 89 of SEQ ID NO:2;

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- (xxiii) amino acids 1 to 113 of SEQ ID NO:2;
- (xxiv) amino acids 1 to 137 of SEQ ID NO:2;
- (xxv) amino acids 1 to 161 of SEQ ID NO:2;
- (xxvi) amino acids 1 to 185 of SEQ ID NO:2;
- (xxvii) amino acids 1 to 209 of SEQ ID NO:2;
- (xxviii) amino acids 1 to 233 of SEQ ID NO:2;
- (xxix) amino acids 1 to 257 of SEQ ID NO:2;
- (xxxx) amino acids 1 to 281 of SEQ ID NO:2;
- (xxxxi) amino acids 1 to 305 of SEQ ID NO:2;
- (xxxxii) amino acids 1 to 329 of SEQ ID NO:2;
- (xxxxiii) amino acids 1 to 353 of SEQ ID NO:2;
- (xxxxiv) amino acids 1 to 416 of SEQ ID NO:2;
- (xxxxv) amino acids 1 to 424 of SEQ ID NO:2;
- (xxxxvi) amino acids 1 to 493 of SEQ ID NO:2;
- (xxxxvii) amino acids 1 to 551 of SEQ ID NO:2;
- (xxxxviii) amino acids 1 to 531 of SEQ ID NO:2;
- (iL) amino acids 1 to 532 of SEQ ID NO:2;
- (Li) amino acids 34 to 89 of SEQ ID NO:2;
- (Lii) amino acids 34 to 113 of SEQ ID NO:2;
- (Liii) amino acids 34 to 137 of SEQ ID NO:2;
- (Liv) amino acids 34 to 161 of SEQ ID NO:2;
- (Lv) amino acids 34 to 185 of SEQ ID NO:2;
- (Lvi) amino acids 34 to 209 of SEQ ID NO:2;
- (Lvii) amino acids 34 to 233 of SEQ ID NO:2;
- (Lviii) amino acids 34 to 257 of SEQ ID NO:2;
- (Lix) amino acids 34 to 281 of SEQ ID NO:2;
- (Lx) amino acids 34 to 305 of SEQ ID NO:2;
- (Lxi) amino acids 34 to 329 of SEQ ID NO:2;
- (Lxii) amino acids 34 to 353 of SEQ ID NO:2;
- (Lxiii) amino acids 34 to 416 of SEQ ID NO:2;
- (Lxiv) amino acids 34 to 424 of SEQ ID NO:2;
- (Lxv) amino acids 34 to 493 of SEQ ID NO:2;
- (Lxvi) amino acids 34 to 551 of SEQ ID NO:2;
- (Lxxi) amino acids 34 to 530 of SEQ ID NO:2;

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- (Lxxii) amino acids 34 to 531 of SEQ ID NO:2;
- (Lxxiii) amino acids 34 to 532 of SEQ ID NO:2;
- (Lxxiv) amino acids 34 to 533 of SEQ ID NO:2;
- (Lxxv) amino acids 34 to 534 of SEQ ID NO:2;
- (Lxxvi) amino acids 34 to 535 of SEQ ID NO:2;
- (Lxxvii) amino acids 34 to 536 of SEQ ID NO:2;
- (Lxxviii) amino acids 34 to 537 of SEQ ID NO:2;
- (Lxxix) amino acids 34 to 538 of SEQ ID NO:2;
- (Lxxx) amino acids 34 to 539 of SEQ ID NO:2;
- (Lxxxi) amino acids 30 to 532 of SEQ ID NO:2;
- (Lxxxii) amino acids 31 to 532 of SEQ ID NO:2;
- (Lxxxiii) amino acids 32 to 532 of SEQ ID NO:2;
- (Lxxxiv) amino acids 33 to 532 of SEQ ID NO:2;
- (Lxxxv) amino acids 34 to 532 of SEQ ID NO:2;
- (Lxxxvi) amino acids 35 to 532 of SEQ ID NO:2;
- (Lxxxvii) amino acids 36 to 532 of SEQ ID NO:2;
- (Lxxxviii) amino acids 30 to 531 of SEQ ID NO:2;
- (Lxxxix) amino acids 31 to 531 of SEQ ID NO:2;
- (Lxxxx) amino acids 32 to 531 of SEQ ID NO:2;
- (Lxxxxi) amino acids 33 to 531 of SEQ ID NO:2;
- (Lxxxxii) amino acids 34 to 531 of SEQ ID NO:2;
- (Lxxxxiii) amino acids 35 to 531 of SEQ ID NO:2;
- (Lxxxxiv) amino acids 36 to 531 of SEQ ID NO:2;
- (Lxxxxv) amino acids 36 to 89 of SEQ ID NO:2;
- (Lxxxxvi) amino acids 36 to 113 of SEQ ID NO:2;
- (Lxxxxvii) amino acids 36 to 137 of SEQ ID NO:2;
- (Lxxxxviii) amino acids 36 to 161 of SEQ ID NO:2;
- (Lxxxxvc) amino acids 36 to 185 of SEQ ID NO:2;
- (c) amino acids 36 to 209 of SEQ ID NO:2;
- (ci) amino acids 36 to 233 of SEQ ID NO:2;
- (cii) amino acids 36 to 257 of SEQ ID NO:2;
- (ciii) amino acids 36 to 281 of SEQ ID NO:2;
- (civ) amino acids 36 to 305 of SEQ ID NO:2;
- (cv) amino acids 36 to 329 of SEQ ID NO:2;

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- (cvi) amino acids 36 to 353 of SEQ ID NO:2;
- (cvii) amino acids 36 to 416 of SEQ ID NO:2;
- (cviii) amino acids 36 to 424 of SEQ ID NO:2;
- (cix) amino acids 36 to 493 of SEQ ID NO:2;
- (cx) amino acids 36 to 551 of SEQ ID NO:2;
- (cxi) amino acids 36 to 530 of SEQ ID NO:2;
- (cxii) amino acids 36 to 531 of SEQ ID NO:2;
- (cxiii) amino acids 36 to 532 of SEQ ID NO:2;
- (cxiv) amino acids 36 to 533 of SEQ ID NO:2;
- (cxv) amino acids 36 to 534 of SEQ ID NO:2;
- (cxvi) amino acids 36 to 535 of SEQ ID NO:2;
- (cxvii) amino acids 36 to 536 of SEQ ID NO:2;
- (cxviii) amino acids 36 to 537 of SEQ ID NO:2;
- (cxix) amino acids 36 to 538 of SEQ ID NO:2;
- (cxx) amino acids 36 to 539 of SEQ ID NO:2;
- (cxxi) variants or derivatives of any of said polypeptide fragments, and
- (cxxii) a combination of at least two of any of said polypeptide fragments

or variants or derivatives thereof.

18. The method of claim 17, wherein said soluble LINGO-1 polypeptide comprises amino acids 34-532 of SEQ ID NO: 2 or amino acids 36-532 of SEQ ID NO:2.

19. The method of any one of claims 11-18 wherein said soluble LINGO-1 polypeptide comprises a LINGO-1 Ig domain or fragment, variant, or derivative thereof.

20. The method of claim 19, wherein said soluble LINGO-1 polypeptide comprises amino acids 417-493 of SEQ ID NO:2.

21. The method of any one of claims 11 to 20, wherein said soluble LINGO-1 polypeptide further comprises a non-LINGO-1 moiety.

22. The method of claim 21, wherein said non-LINGO-1 moiety is a heterologous polypeptide fused to said soluble LINGO-1 polypeptide.

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23. The method of claim 22, wherein heterologous polypeptide is selected from the group consisting of an immunoglobulin fragment, serum albumin, a targeting protein, a reporter protein, and a purification-facilitating protein.

24. The method of claim 23, wherein said heterologous polypeptide is an immunoglobulin fragment.

25. The method of claim 24, wherein said immunoglobulin fragment comprises a hinge and Fc region.

26. The method of any one of claims 21-25, wherein said soluble LINGO-1 polypeptide is conjugated to a polymer.

27. The method of claim 26, wherein the polymer is selected from the group consisting of a polyalkylene glycol, a sugar polymer, and a polypeptide.

28. The method of claim 27, wherein the polymer is a polyalkylene glycol.

29. The method of claim 28, wherein the polyalkylene glycol is polyethylene glycol (PEG).

30. The method of claim 26, wherein said soluble LINGO-1 polypeptide is conjugated to 1, 2, 3 or 4 polymers.

31. The method of claim 30, wherein the total molecular weight of the polymers is from 5,000 Da to 100,000 Da.

32. The method of any one of claims 1-10, wherein said LINGO-1 antagonist comprises a LINGO-1 antibody, or fragment thereof.

33. The method of claim 32, wherein said LINGO-1 antibody or fragment thereof specifically binds to an epitope consisting essentially of a polypeptide fragment selected from the group consisting of:

(i) amino acids 66 to 89 of SEQ ID NO:2;

- (ii) amino acids 66 to 113 of SEQ ID NO:2;
- (iii) amino acids 66 to 137 of SEQ ID NO:2;
- (iv) amino acids 90 to 113 of SEQ ID NO:2;
- (v) amino acids 114 to 137 of SEQ ID NO:2;
- (vi) amino acids 138 to 161 of SEQ ID NO:2;
- (vii) amino acids 162 to 185 of SEQ ID NO:2;
- (viii) amino acids 186 to 209 of SEQ ID NO:2;
- (ix) amino acids 210 to 233 of SEQ ID NO:2;
- (x) amino acids 234 to 257 of SEQ ID NO:2;
- (xi) amino acids 258 to 281 of SEQ ID NO:2;
- (xii) amino acids 282 to 305 of SEQ ID NO:2;
- (xiii) amino acids 306 to 329 of SEQ ID NO:2;
- (xiv) amino acids 330 to 353 of SEQ ID NO:2;
- (xv) amino acids 34 to 64 of SEQ ID NO:2;
- (xvi) amino acids 363 to 416 of SEQ ID NO:2;
- (xvii) variants or derivatives of any of said polypeptide fragments; and
- (xviii) a combination of two or more of any of said polypeptide fragments or variants or derivatives thereof.

34. The method of claim 32, wherein said LINGO-1 antibody is selected from the group consisting of: 201', 3A3, 3A6, 3B5, 1A7, 1D5, 1G7, 2B10, 2C11, 2F3, 3P1B1.1F9, 3P1D10.2C3, 3P1E11.3B7, 3P2C6.3G10.2H7, 3P2C9.2G4, 3P4A6.1D9, 3P4A1.2B9, 3P4C2.2D2, 3P4C5.1D8, 3P4C8.2G9, 6P4F4.1Ds, 6P4F4.1F9, 7P1D5.1G9, 1B6.4, 2C7.2, 2D6.1, 2F7.3, 2H3.2, 3C11.1, 3E3.1, 3H11.2, 3G8.1, 2B8.1, 3B5.230-C12 (Li01), 38-D01 (Li02), 35-E04 (Li03), 36-C09 (Li04), 30-A11 (Li05), 34-F02 (Li06), 29-E07 (Li07), 34-G04 (Li08), 36-A12 (Li09), 28-D02 (Li10), 30-B01 (Li11), 34-B03 (Li12), Li13, Li32, Li33, Li34, 3383 (L1a.1), 3495(L1a.2), 3563 (L1a.3), 3564 (L1a.4), 3565 (L1a.5), 3566 (L1a.6), 3567 (L1a.7), 3568 (L1a.8), 3569 (L1a.9), 3570 (L1a.10), 3571 (L1a.11), 3582 (L1a.12), 1968 (L1a.13), 3011, 3012, 3013, 3418, 3422, 3562, D05, D07, D08, D10 and D11.

35. The method of claim 34 wherein said LINGO-1 antibody is 1A7.

36. The method of claim 34 wherein said LINGO-1 antibody is 3B5.

37. The method of any one of claims 1-10, wherein said LINGO-1 antagonist comprises a LINGO-1 antagonist polynucleotide.

38. The method of claim 37, wherein said LINGO-1 antagonist polynucleotide is selected from the group consisting of:

- (i) an antisense polynucleotide;
- (ii) a ribozyme;
- (iii) a small interfering RNA (siRNA); and
- (iv) a small-hairpin RNA (shRNA).

39. The method of claim 38, wherein said LINGO-1 antagonist polynucleotide is an antisense polynucleotide comprising at least 10 bases complementary to the coding portion of the LINGO-1 mRNA.

40. The method of claim 39, wherein said LINGO-1 antagonist polynucleotide is a ribozyme.

41. The method of claim 38, wherein said LINGO-1 antagonist polynucleotide is a siRNA.

42. The method of claim 38 wherein said LINGO-1 antagonist polynucleotide is a shRNA.

43. The method of claim 39, wherein said shRNA comprises the nucleotide sequence:
UGAUCGUCAU CCUGCUAGAC UUCAAGAGAG UCUAGCAGGA UGACGAUCUU
UUUUC (SEQ ID NO:34).

44. The method of any one of claims 1-10, wherein said LINGO-1 antagonist is an aptamer.

45. The method of any one of claims 1-6 wherein the contacting comprises (a) introducing into said cell or said CNS neurons a polynucleotide which encodes said LINGO-1 antagonist through operable association with an expression control sequence, and (b) allowing expression of said LINGO-1 antagonist.

46. The method of claim 45, wherein said polynucleotide is introduced into said cell by a method selected from the group consisting of:

- (a) transfection;
- (b) electroporation;
- (c) transduction; and
- (d) direct microinjection.

47. The method of any one of claims 44-46, wherein said polynucleotide is administered as an expression vector.

48. The method of claim 47, wherein said expression vector is a viral vector.

49. The method of claim 48, wherein the viral vector is selected from the group consisting of an adenoviral vector, an alphavirus vector, an enterovirus vector, a pestivirus vector, a lentiviral vector, a baculoviral vector, a herpesvirus vector a papovaviral vector, a poxvirus vector and a parvovirus.

50. The method of claim 49, wherein said herpesvirus vector is selected from the group consisting of a herpes simplex virus vector and an Epstein Barr virus vector.

51. The method of claim 49, wherein said poxvirus vector is a vaccinia virus vector.

52. The method of claim 49, wherein said lentivirus vector is pLL3.7.

53. The method of claim 49, wherein said parvovirus vector is adeno-associated virus (AAV).

54. The method of any one of claims 1-6 or 10 further comprising contacting the cell, the neuron or the CNS neurons with an additional antagonist selected from the group consisting of:

- (i) an NGF antagonist;
- (ii) an NgR1 antagonist;
- (iii) a TAJ antagonist; and

(iv) an OMgp antagonist.

55. The method of any one of claims 7-9 further comprising administering an additional antagonist selected from the group consisting of:

- (i) an NGF antagonist;
- (ii) an NgR1 antagonist;
- (iii) a TAJ antagonist; and
- (iv) an OMgp antagonist.

56. The method of any one of claims 6, 8, 9 or 10 wherein said TrkB agonist is a TrkB agonist compound.

57. The method of claim 56 wherein the TrkB agonist compound is selected from the group consisting of L-783,281, adenosine and CGS 21680.

58. The method of any one of claims 6, 8, 9 or 10 wherein said TrkB agonist is a TrkB-agonist polypeptide.

59. The method of claim 58 wherein the TrkB agonist polypeptide is selected from the group consisting of a TrkB ligand, a fragment of a TrkB ligand, a variant of a TrkB ligand, a TrkB polypeptide, a fragment of a TrkB polypeptide or a variant of a TrkB polypeptide.

60. The method of claim 59 wherein the TrkB agonist polypeptide is a BDNF polypeptide.

61. The method of any one of claims 58-60 wherein said TrkB agonist polypeptide further comprises a non-TrkB-agonist moiety.

62. The method of claim 61, wherein said non-TrkB-agonist moiety is a heterologous polypeptide fused to said TrkB-agonist polypeptide.

63. The method of any one of claims 61 or 62, wherein said non-TrkB-agonist moiety is selected from the group consisting of an immunoglobulin fragment, serum albumin, a targeting protein, a reporter protein, and a purification-facilitating protein.

64. The method of claim 63, wherein said non-TrkB-agonist moiety is an immunoglobulin fragment.

65. The method of claim 63, wherein said immunoglobulin fragment comprises a hinge and Fc region.

66. The method of claim 61, wherein said TrkB agonist polypeptide is conjugated to a polymer.

67. The method of claim 66, wherein the polymer is selected from the group consisting of a polyalkylene glycol, a sugar polymer, and a polypeptide.

68. The method of claim 67, wherein the polymer is polyethylene glycol (PEG).

69. The method of claim 61, wherein said TrkB agonist polypeptide is conjugated to a 1-acyl-glycerol derivative.

70. The method of claim 66, wherein said TrkB agonist polypeptide is conjugated to 1, 2, 3 or 4 polymers.

71. The method of claim 70, wherein the total molecular weight of the polymers is from 5,000 Da to 100,000 Da.

72. The method of any one of claims 6, 8, 9 or 10 wherein said TrkB agonist is a TrkB agonist antibody or fragment thereof.

73. The method of claim 72, wherein said TrkB agonist antibody or fragment thereof is selected from the group consisting of: 6E2, 7F5, 11E1, 16E11, 17D11, 19E12, 29D7 or a TrkB monoclonal antibody.

74. The method of any one of claims 6, 8, 9 or 10 wherein said TrkB agonist is a TrkB agonist polynucleotide.

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75. The method of claim 74 wherein the TrkB agonist polynucleotide encodes a neurotrophin or a fragment or variant thereof.

76. The method of claim 75, wherein the neurotrophin is BDNF.

77. The method of any one of claims 6, 8, 9 or 10 wherein said TrkB agonist is a TrkB agonist aptamer.

78. The method of any one of claims 1-6 or 10, wherein said cell or neuron or said CNS neurons are in a mammal, and wherein said contacting comprises administering an effective amount of said LINGO-1 antagonist or said LINGO-1 antagonist and TrkB agonist to a mammal in need thereof.

79. The method of any one of claims 7-10 or 78, wherein said mammal has been diagnosed with a disease, disorder, or injury involving neurodegeneration.

80. The method of claim 79, wherein said disease, disorder, or injury is glaucoma.

81. The method of any one of claims 7-10 or 78 wherein at least one of said TrkB agonist or LINGO-1 antagonist is administered by bolus injection or chronic infusion.

82. The method of any one of claims 7-10 or 78 wherein at least one of said TrkB agonist or LINGO-1 antagonist is administered directly into the central nervous system.

83. The method of any one of claims 6, 8, 9, 10 or 78 wherein said TrkB agonist is selected from the group consisting of a TrkB agonist polynucleotide or a polynucleotide encoding a TrkB agonist polypeptide or TrkB agonist aptamer.

84. The method of claim 83, wherein the contacting or administering comprises (a) introducing into a cell the TrkB agonist polynucleotide or a polynucleotide which encodes said TrkB agonist and (b) allowing expression of said TrkB agonist.

85. The method of claim 84, wherein said polynucleotide is introduced into said cell by a method selected from the group consisting of:

- 123 -

- (a) transfection;
- (b) electroporation;
- (c) transduction; and
- (d) direct microinjection.

86. The method of any one of claims 84-85, wherein said cell is in a mammal and wherein said introducing comprises (a) administering to said mammal said TrkB agonist polynucleotide or said polynucleotide which encodes said TrkB agonist and (b) allowing expression of said TrkB agonist.

87. The method of claim 83, wherein said TrkB agonist polynucleotide or said polynucleotide which encodes said TrkB agonist is administered in a cultured host cell capable of expressing said polynucleotide.

88. The method of any one of claims 84 to 87, wherein said polynucleotide is administered as an expression vector .

89. The method of claim 88, wherein said expression vector is a viral vector.

90. The method of any one of claims 86 to 89 wherein said polynucleotide is introduced into said mammal at or near the site of the disease, disorder or injury.

91. The method of claim 87, wherein said cultured host cell is made by a method comprising (a) transforming or transfecting a recipient host cell with the TrkB agonist polynucleotide or polynucleotide encoding a TrkB agonist of claim 83 or the vector of claim 88 or claim 89, and (b) culturing said transformed or transfected host cell.

92. The method of any one of claims 87 to 91, wherein said host cell is derived from the mammal to be treated.

93. The method of claim 89, wherein the viral vector is selected from the group consisting of an adenoviral vector, an alphavirus vector, an enterovirus vector, a pestivirus vector, a lentiviral vector, a baculoviral vector, a herpesvirus vector a papovaviral vector, a poxvirus vector and a parvovirus.

94. The method of claim 93, wherein said herpesvirus vector is selected from the group consisting of a herpes simplex virus vector and an Epstein Barr virus vector.

95. The method of claim 93, wherein said poxvirus vector is a vaccinia virus vector.

96. The method of claim 93, wherein said lentivirus vector is pLL3.7.

97. The method of claim 93, wherein said parvovirus vector is adeno-associated virus (AAV).

98. The method any one of claims 8-10 or 78, wherein said TrkB agonist is administered by a route selected from the group consisting of topical administration, intraocular administration, intravitreal administration, parenteral administration, intrathecal administration, subdural administration, subcutaneous administration or via a capsule implant.

99. The method of claim 98 wherein at least one of the LINGO-1 antagonist or TrkB agonist is administered directly into the eye.

100. The method of claim 98, wherein at least one of the LINGO-1 antagonist or TrkB agonist is administered via a capsule implant.

101. The method of any one of claims 1-6 or 78, wherein the cell or neuron or the CNS neurons are sensory neurons.

102. The method of claim 101, wherein the sensory neuron in a retinal ganglion cell (RGC).

103. The method of any one of claims 7-10 or 78, wherein the mammal suffers from an optical neuropathy.

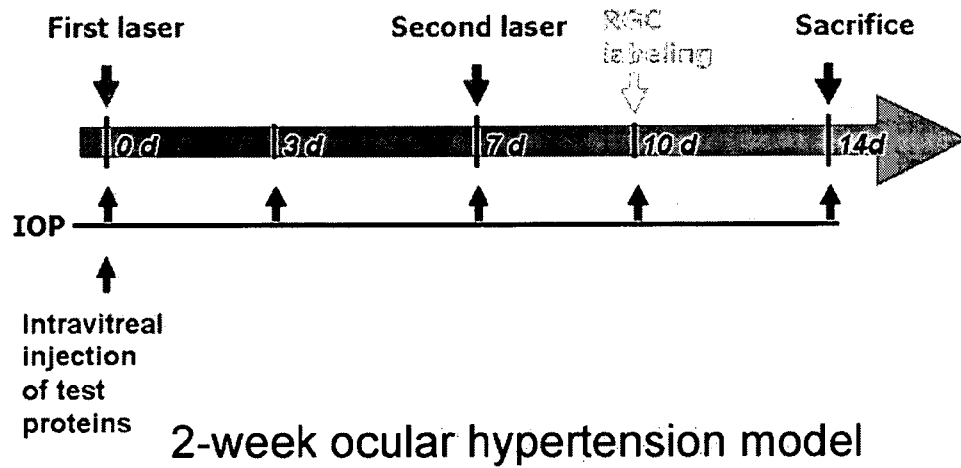
104. The method of claim 103, wherein said optical neuropathy is glaucoma.

105. The method of claim 101, wherein the sensory neuron is a hairy cell.

106. The method of claim 9 or 78, wherein the mammal suffers from a disease or disorder selected from the group consisting of ALS, Huntington's disease, Alzheimer's disease, Parkinson's disease, diabetic neuropathy, stroke and hearing loss.

107. The method of claim 34 wherein said LINGO-1 antibody is Li33.

A



B

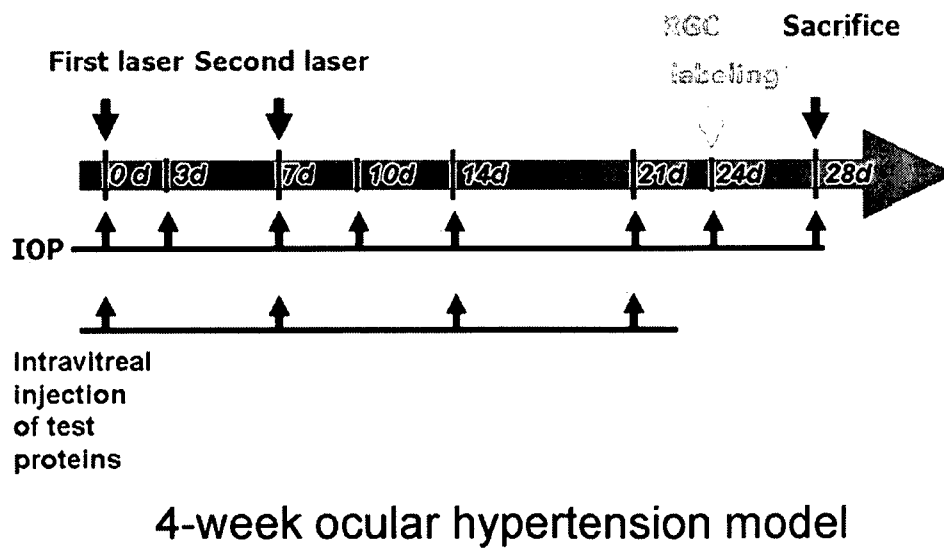


Fig. 1

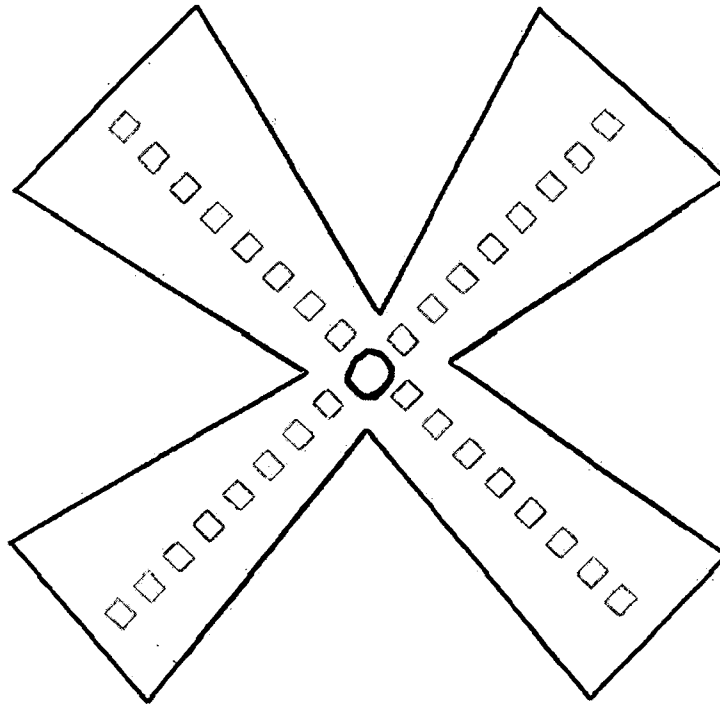
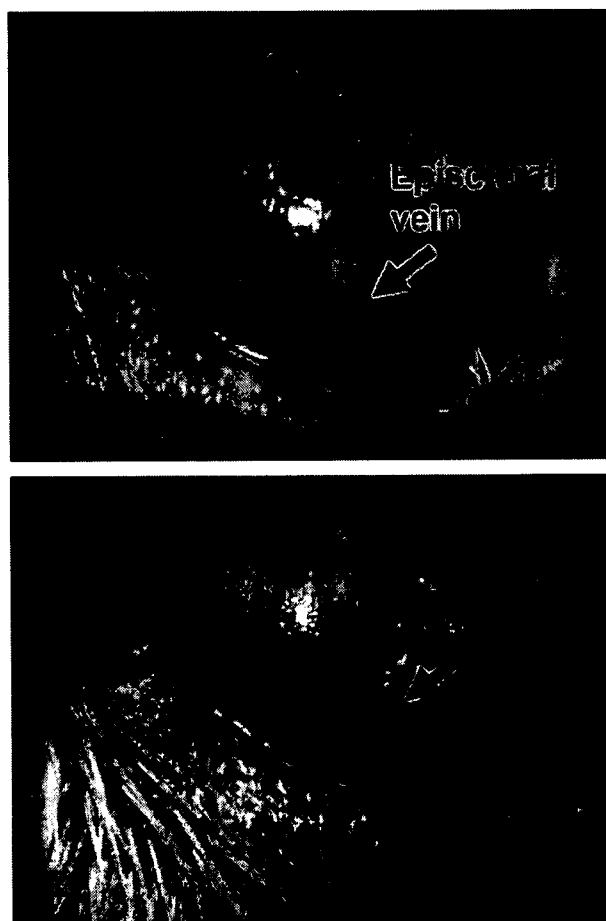


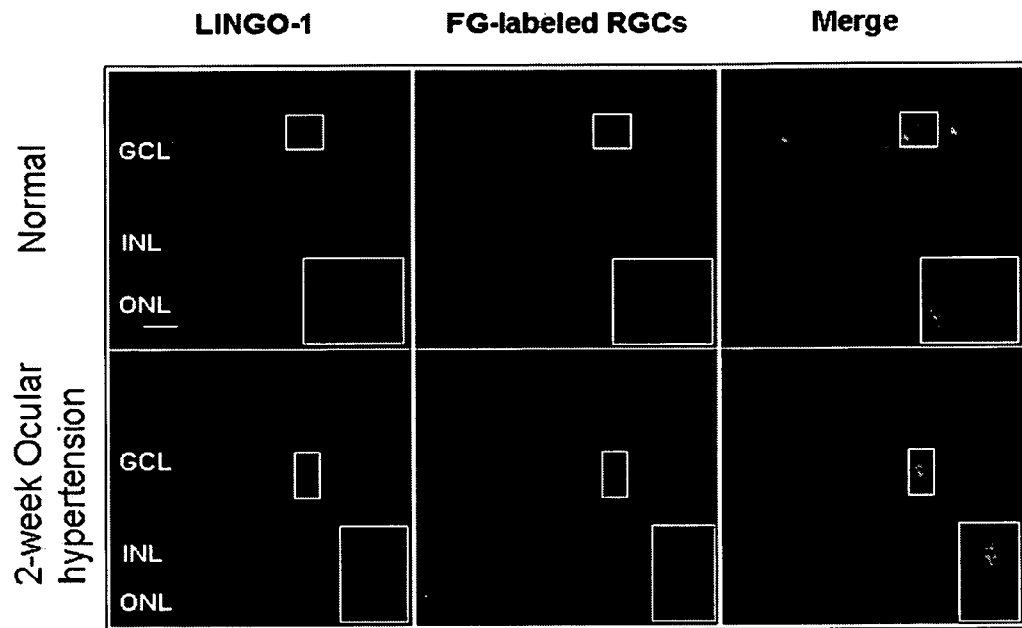
Fig. 2



A: Prelaser treatment
B: Postlaser treatment

Fig. 3

A



B

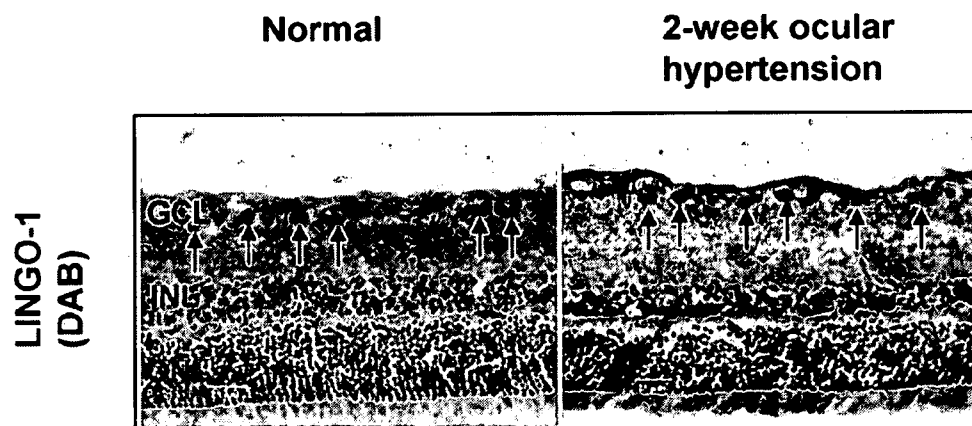


Fig. 4

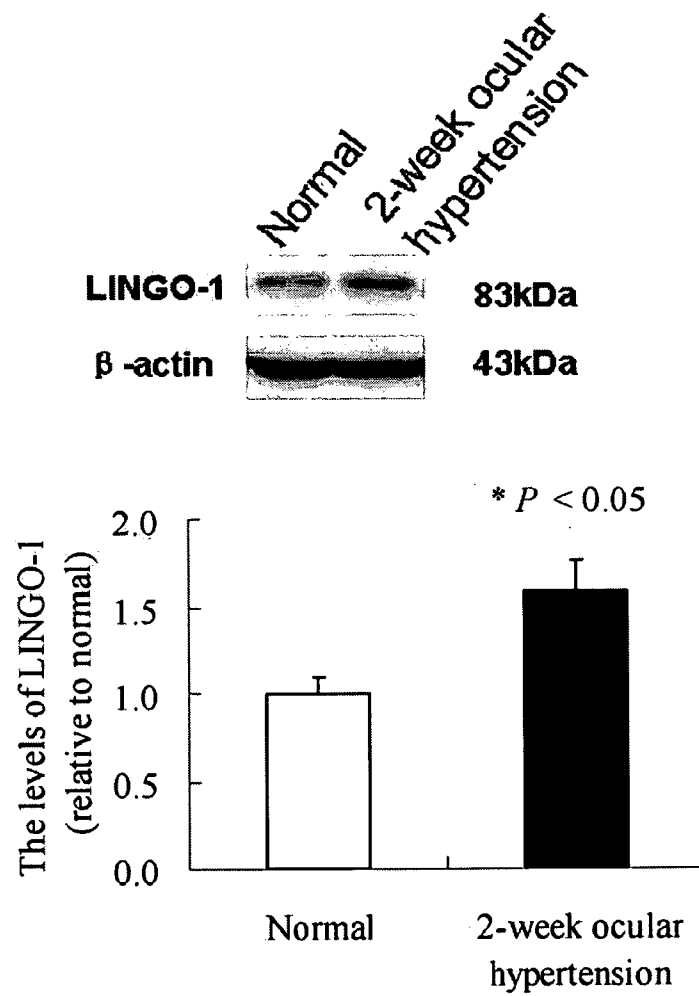


Fig. 5

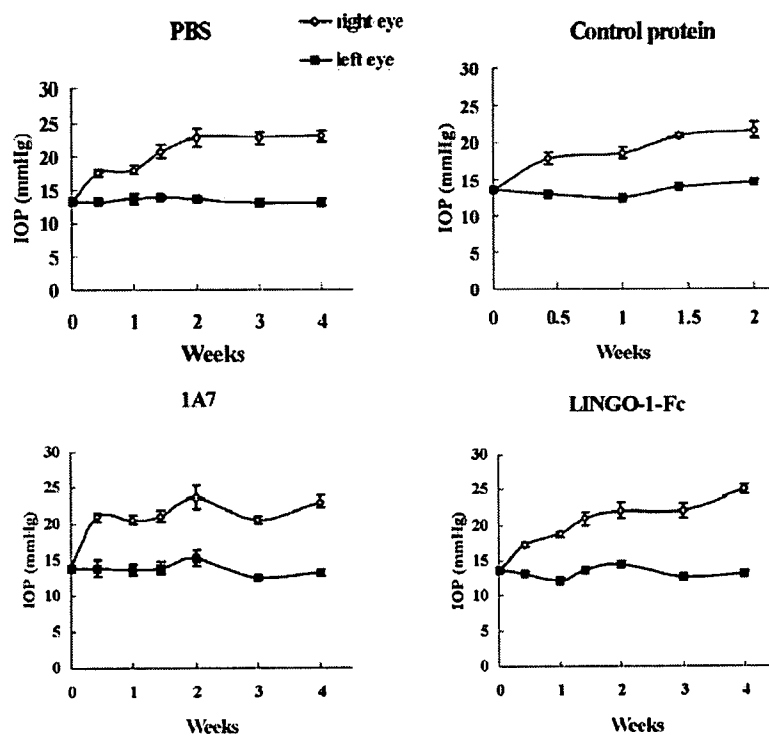
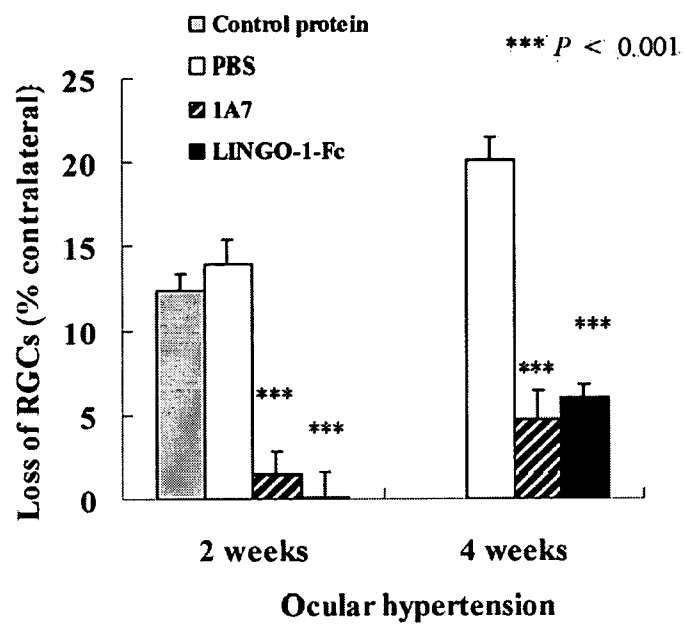


Fig. 6

A



B

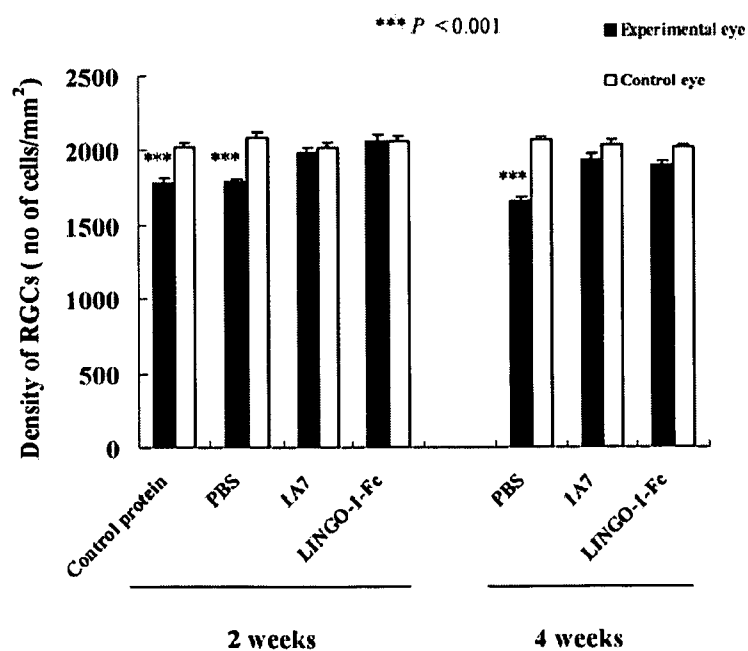


Fig. 7

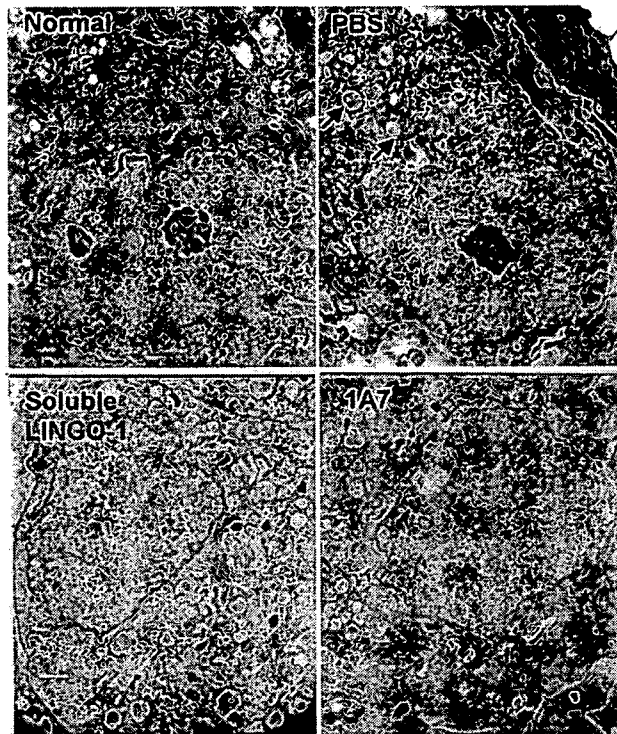
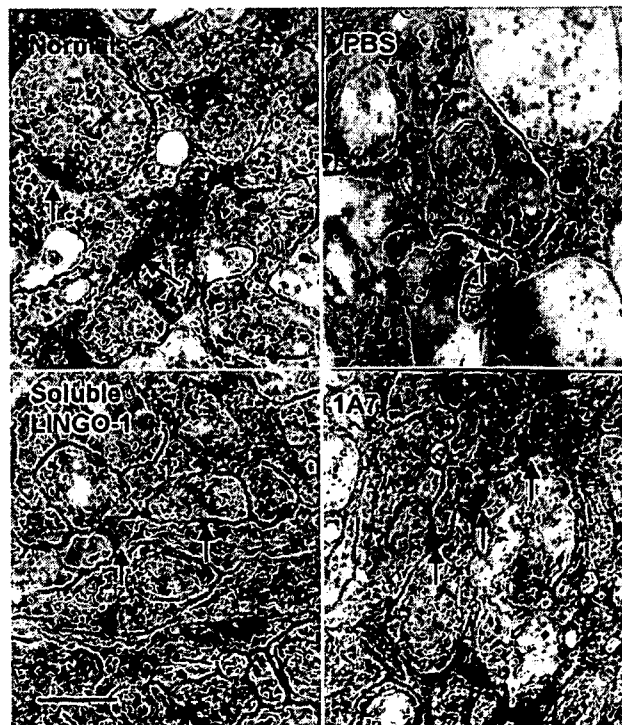
A Retinal ganglion cells**B** Inner plexiform layer

Fig. 8

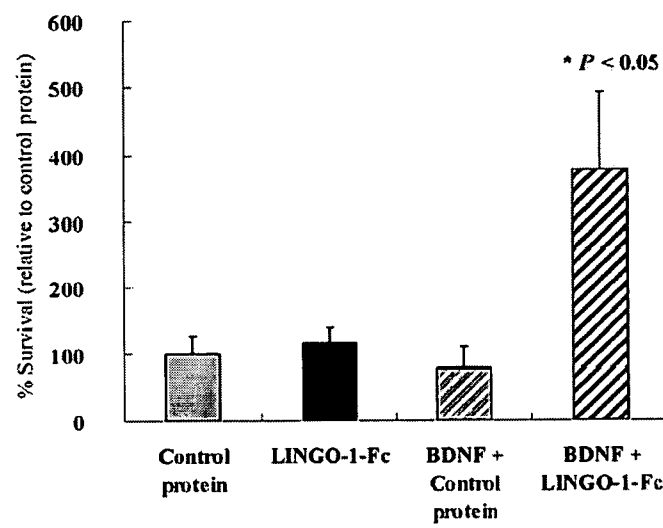


Fig. 9

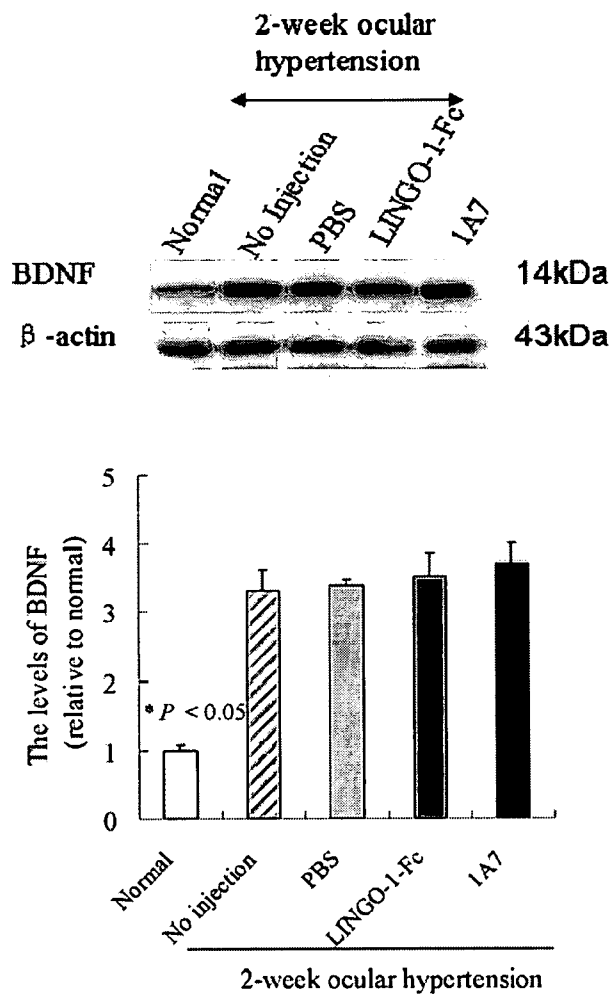
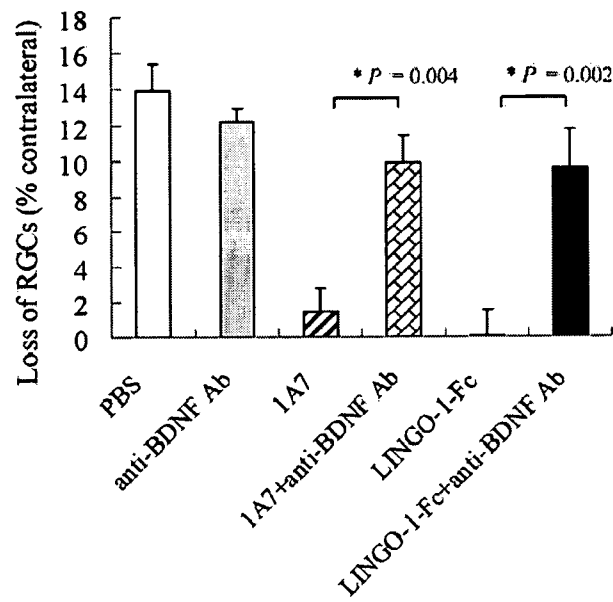


Fig. 10

A



B

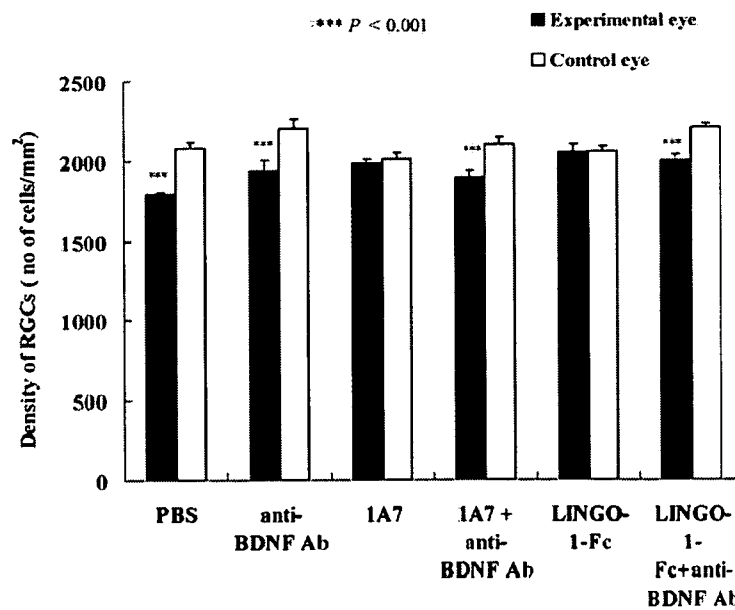
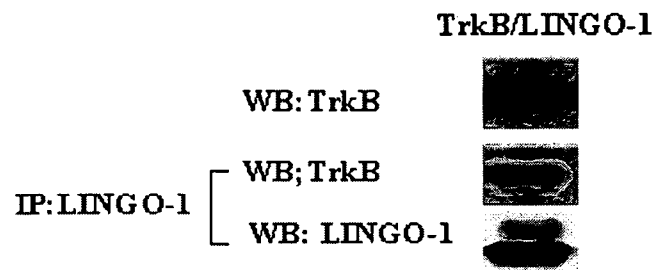


Fig. 11

A



B

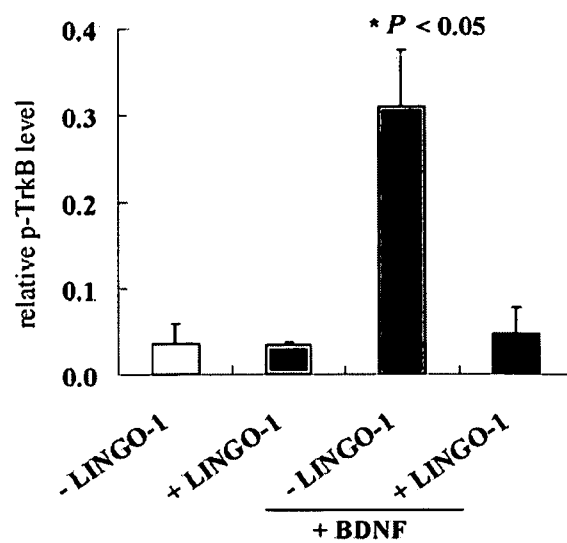
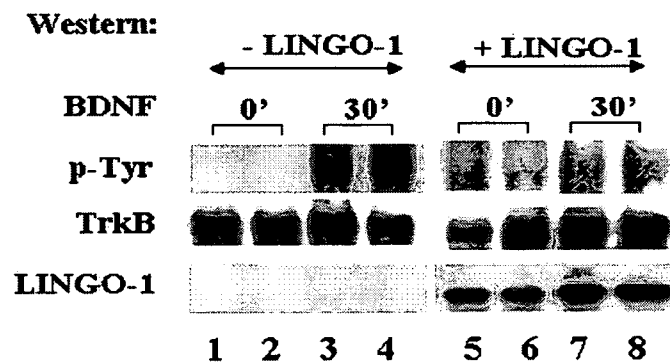
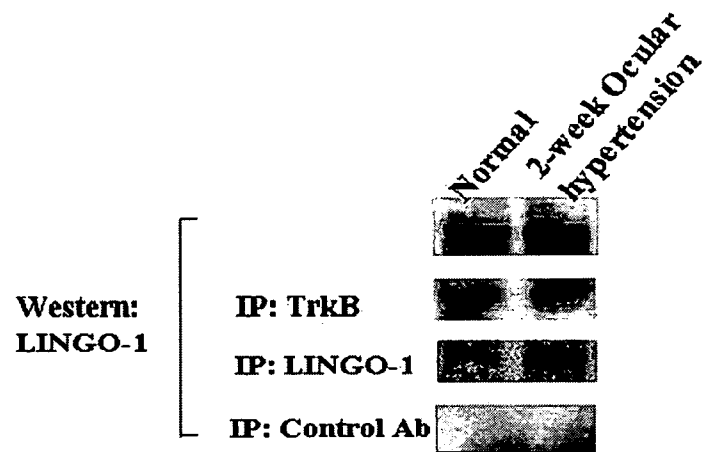


Fig. 12

A



B

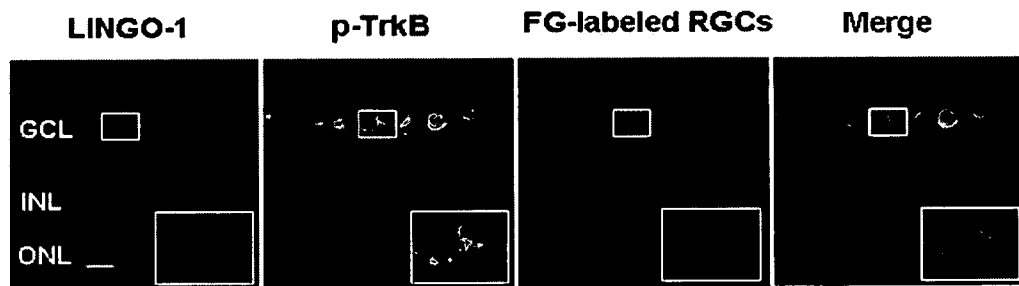


Fig. 13

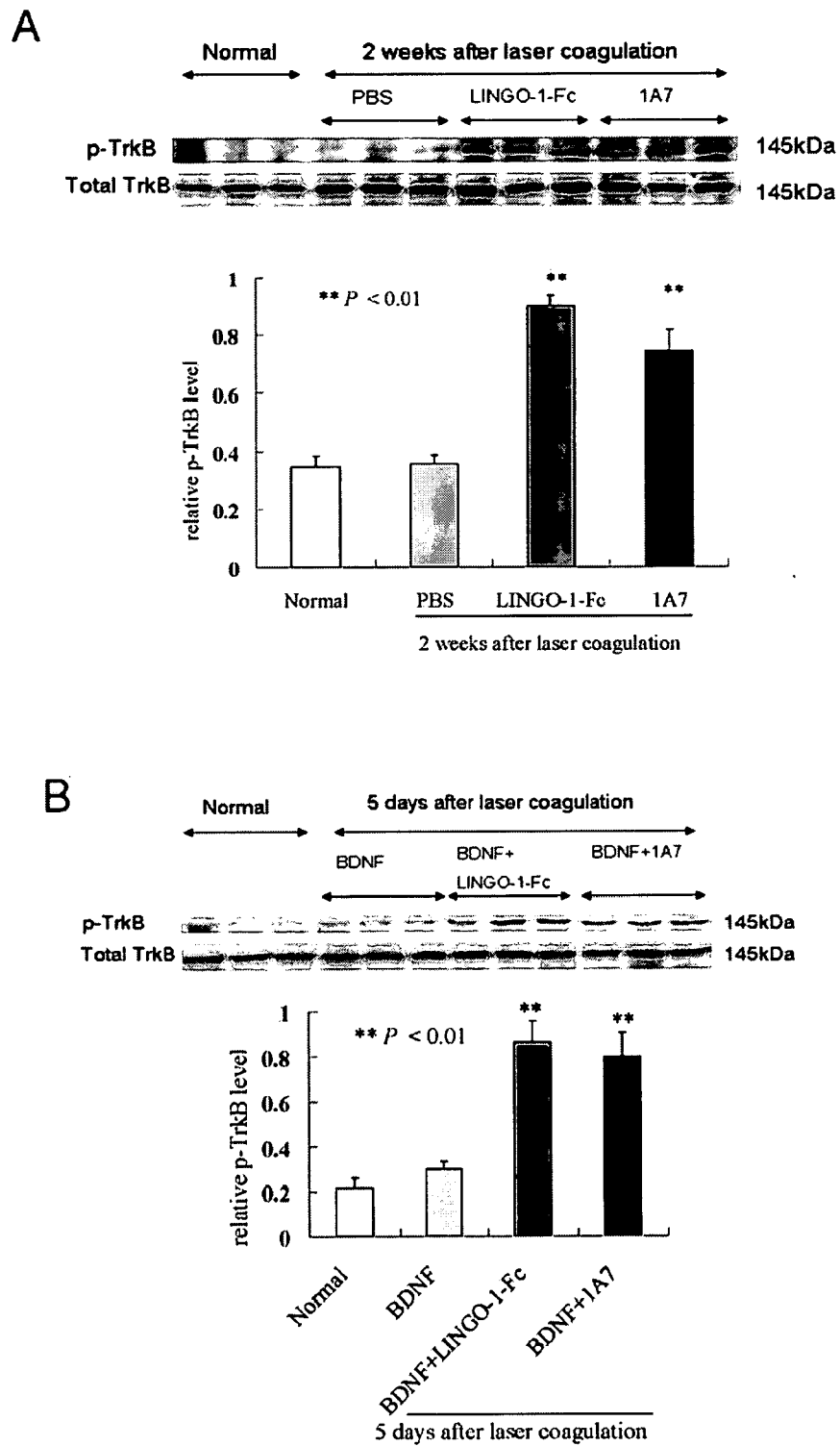


Fig. 14

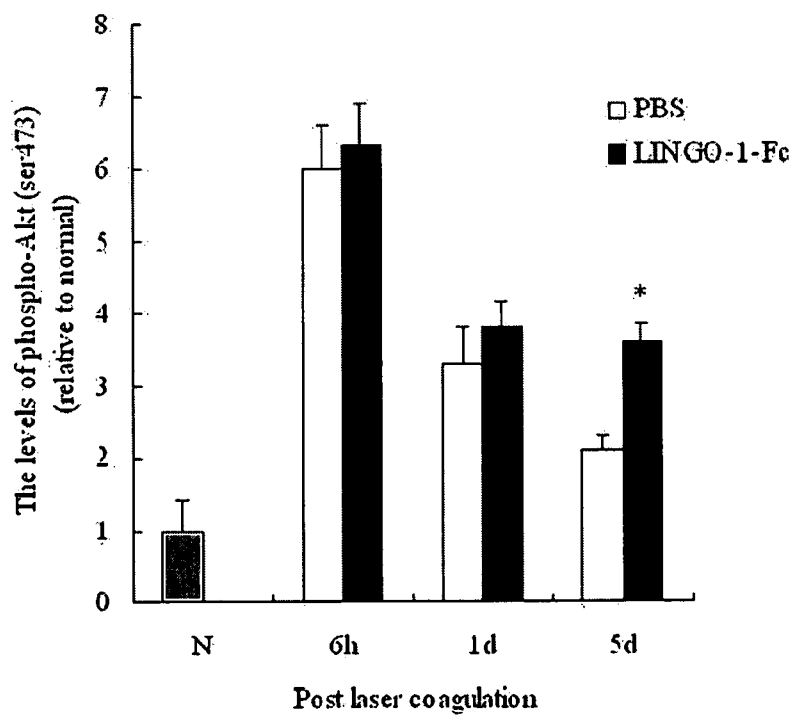
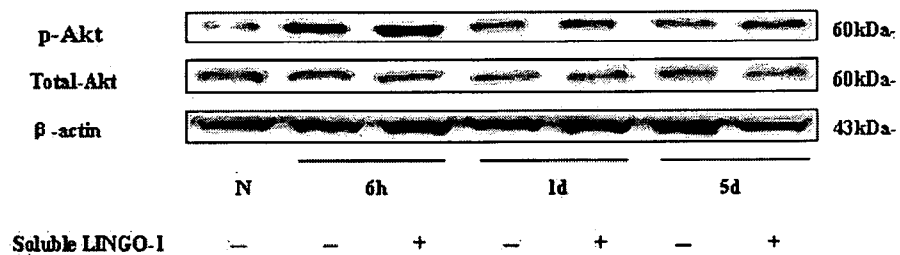


Fig. 15

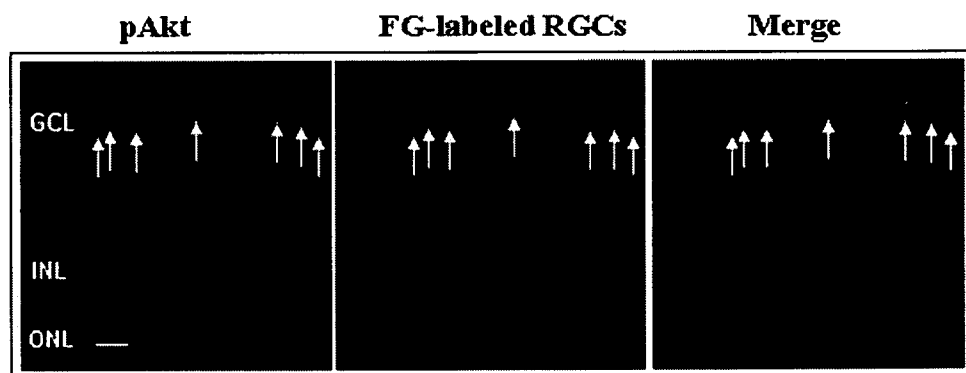
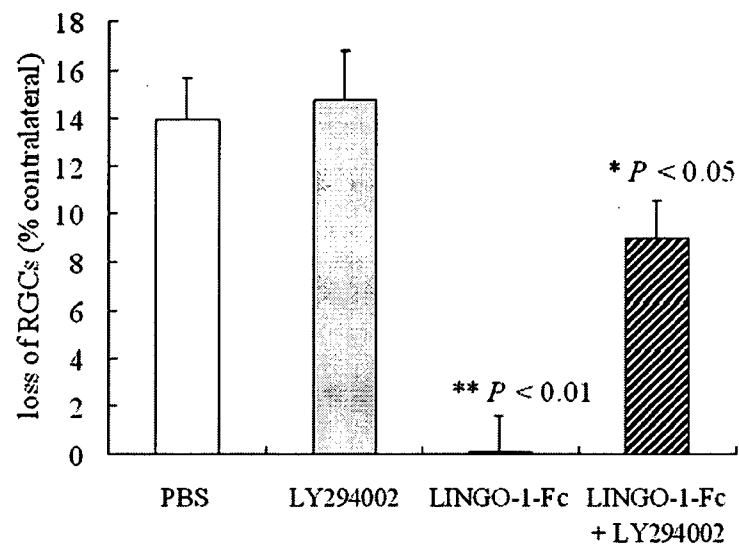


Fig. 16

A



B

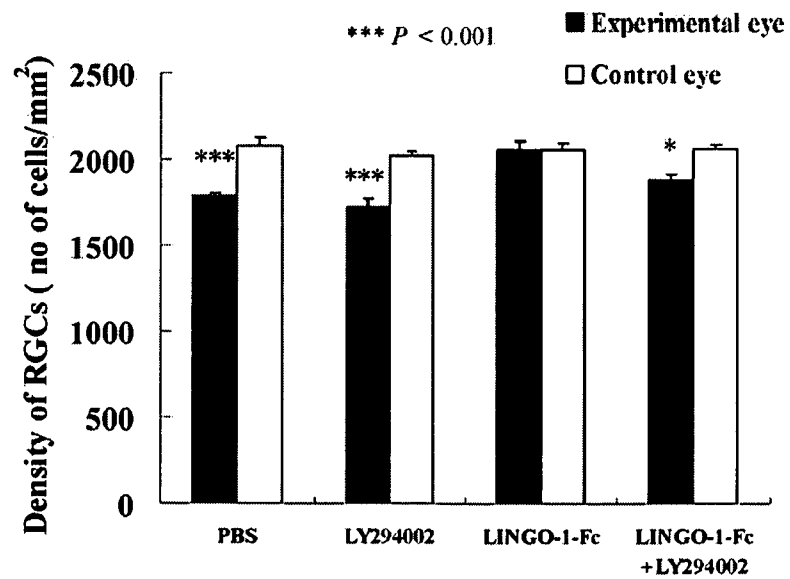


Fig. 17

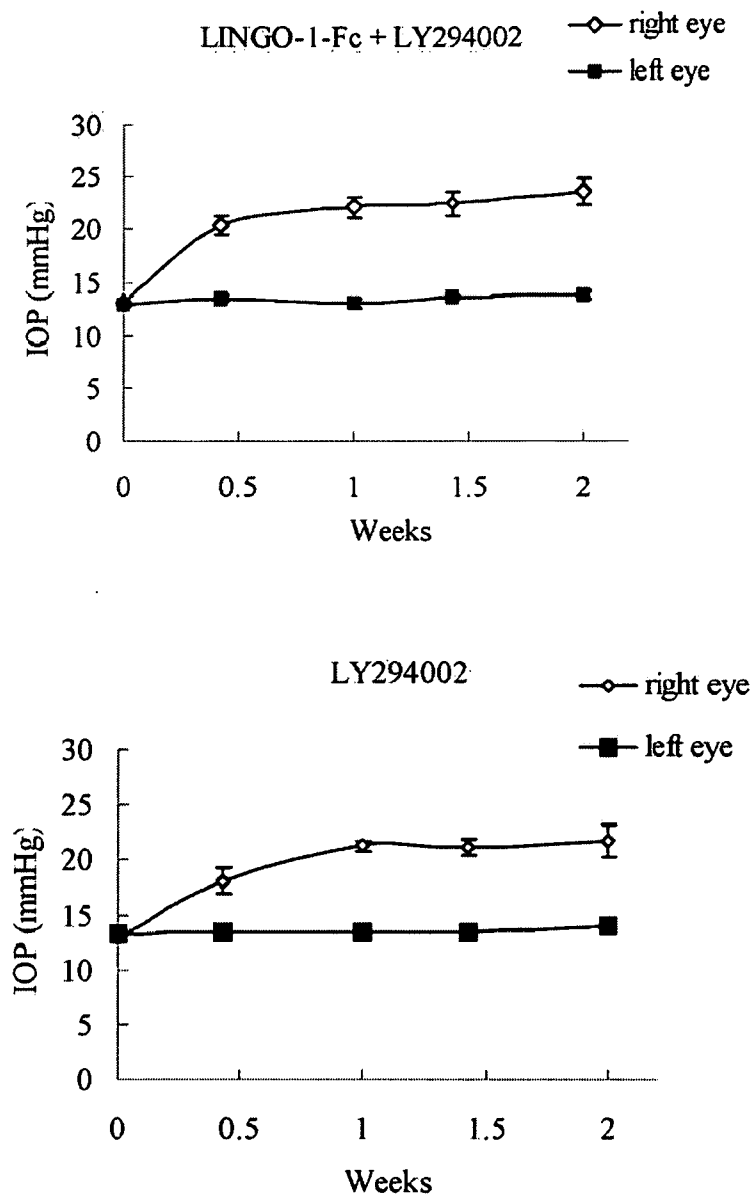


Fig. 18

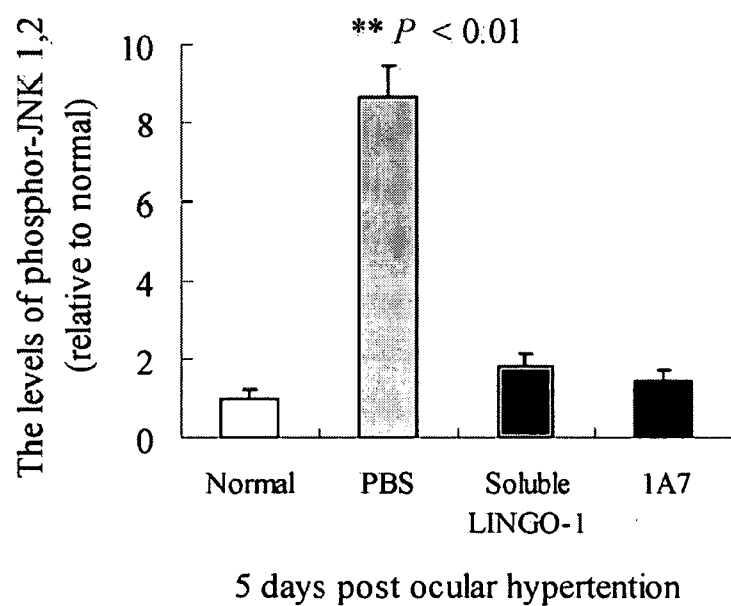
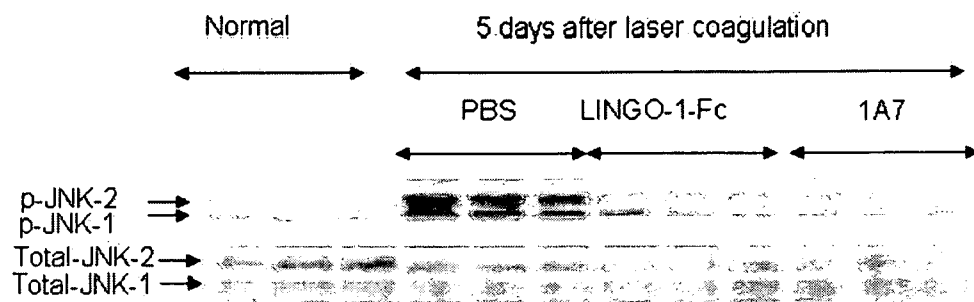


Fig. 19

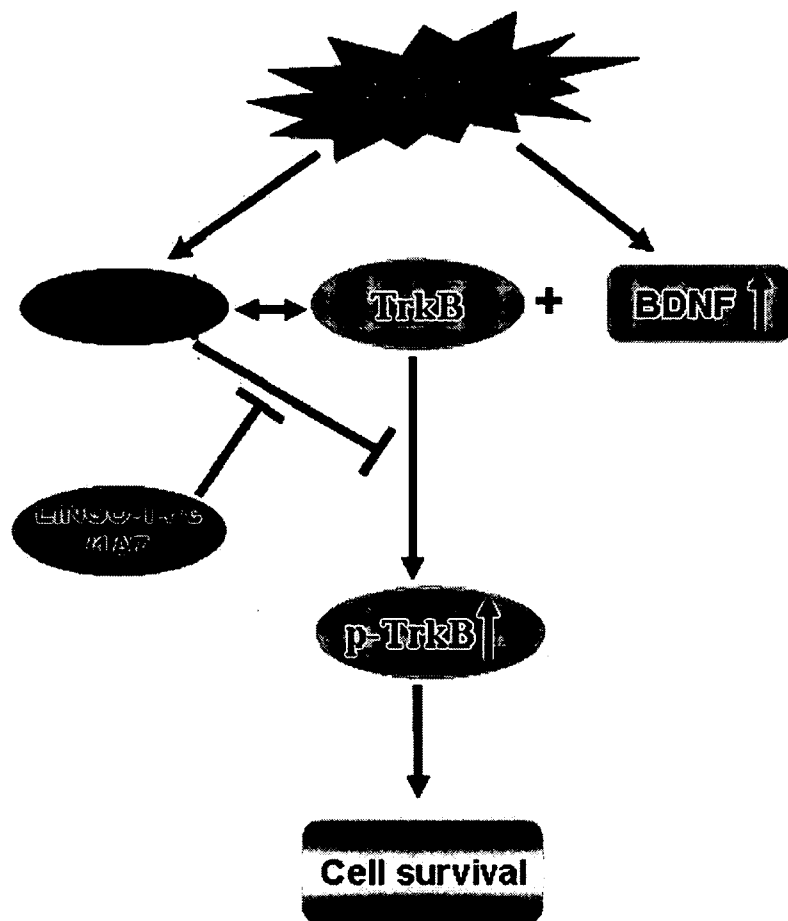


Fig. 20

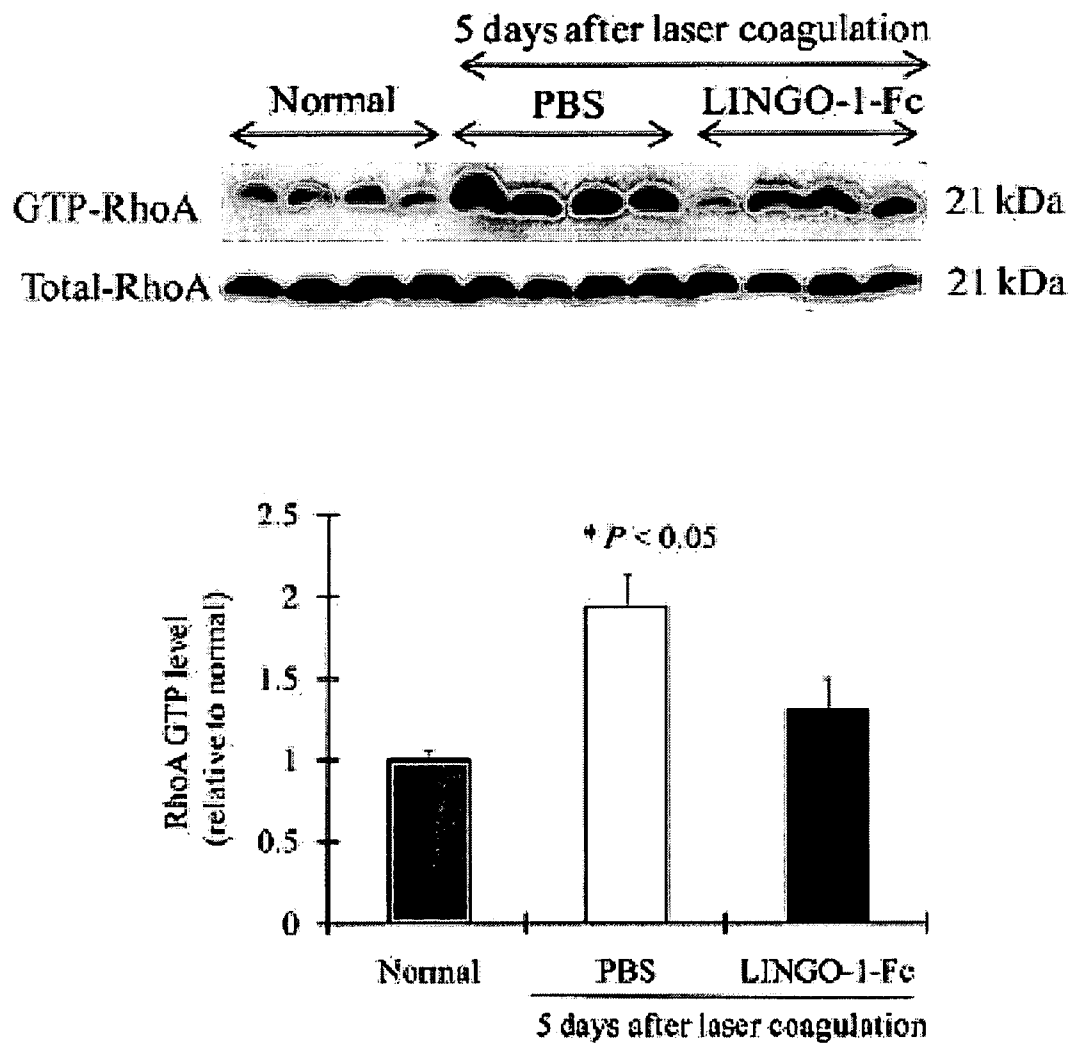


Fig. 21

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 08/11633

A. CLASSIFICATION OF SUBJECT MATTER

IPC(8) - A01N 37/18; A61K 31/00 (2009.01)

USPC - 514/2, 514/44

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC(8) - A01N 37/18; A61K 31/00 (2009.01)

USPC - 514/2, 514/44

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
424/130.1, 78.08

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

WEST - DB=PGPB,USPT,USOC,EPAB,JPAB; PLUR=YES; OP=ADJ; google; google scholar

Search terms: LINGO, LINGO1, LINGO-1, leucine-rich repeat, leucine rich repeat, TrkB, BDNF/NT-3, NTRK2, NTRK 2, NTRK-2, neurotrophin tyrosine kinase, inhibit, inhibitor, inhibition, inhibited, inhibiting, repressor, repression, repressed, repressed,

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 2007/0186296 A1 (GAO et al.) 09 August 2007 (09.08.2007) para [0004]; para [0006]; para [0007]; para [0016]; para [0028]; para [0027]; para [0054]; para [0069]; para [0070]; para [0071];	1, 45, 46
Y	para [0072]; para [0074]; para [0076]; para [0120]; para [0128]; para [0129]; para [0130]; para [0132]; para [0133]; para [0136].	2-10, 55
Y	US 2007/0060526 A1 (LONGO et al.) 15 March 2007 (15.03.2007) para [0013]; para [0018]; para [0196]; para [0209]; para [0345].	2, 3, 6, 8-10
Y	US 2003/0162734 A1 (MILLER et al.) 28 August 2003 (28.08.2003) para [0022]; para [0035]; para [0036]; para [0042].	4, 5
Y	(CHENG et al.) TrkB Gene Transfer Protects Retinal Ganglion Cells from Axotomy-Induced Death In Vivo. Journal of Neuroscience, 22(10):3977-3986 (15.05.2002) abstract; p. 3959.	7, 8, 55
Y	(HARTMANN et al.) Truncated TrkB receptor-induced outgrowth of dendritic filopodia involves the p75 neurotrophin receptor. Journal of Cell Science 117, 5803-5814 (2004) abstract; p. 5813.	2, 3

☐ Further documents are listed in the continuation of Box C.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&" document member of the same patent family

Date of the actual completion of the international search

26 January 2009 (26.01.2009)

Date of mailing of the international search report

18 FEB 2009

Name and mailing address of the ISA/US

Mail Stop PCT, Attn: ISA/US, Commissioner for Patents
P.O. Box 1450, Alexandria, Virginia 22313-1450

Facsimile No. 571-273-3201

Authorized officer:

Lee W. Young

PCT Helpdesk: 571-272-4300

PCT OSP: 571-272-7774

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 08/11633

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☒ Claims Nos.: 11-44, 47-54, and 56-107
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.
- ☐ The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
- ☐ No protest accompanied the payment of additional search fees.