TIE-2 LIGANDS, METHODS OF MAKING AND USES THEREOF

This International Application claims priority of copending United States Applications U.S. Serial No. 418,595, filed April 6, 1995, and U.S. Serial No. 373,579, filed January 17, 1995, and U.S. Serial No. 353,503, filed December 9, 1994, and U.S. Serial No. 348,492, filed December 2, 1994, and U.S. Serial No. 330,261 filed October 27, 1994, and U.S. Serial No. 319,932, filed October 7, 1994, the contents of each of which are hereby incorporated by reference. Throughout this application various publications are referenced. The disclosures of these publications in their entireties are hereby incorporated by reference into this application.

INTRODUCTION

The present invention relates generally to the field of genetic engineering and more particularly to genes for receptor tyrosine kinases and their cognate ligands, their insertion into recombinant DNA vectors, and the production of the encoded proteins in recipient strains of micro-organisms and recipient eukaryotic cells. More specifically, the present invention is directed to novel ligands, known as the TIE-2 ligands, that bind the TIE-2 receptor, as well as to methods of making and using the TIE-2 ligands. The invention further provides nucleic acid sequences encoding TIE-2 ligands, and methods for the generation of nucleic acids encoding TIE-2 ligands and their gene products. The TIE-2 ligands, as well as nucleic acids encoding them, may be useful in the diagnosis and treatment of certain diseases involving endothelial cells and associated TIE receptors, such as neoplastic diseases involving tumor angiogenesis, wound healing, thromboembolic diseases,

atherosclerosis and inflammatory diseases. More generally, biologically active TIE-2 ligands may be used to promote the growth, survival and/or differentiation of cells expressing the TIE-2 receptor. Biologically active TIE-2 ligand may be used for the in vitro maintenance of TIE-2 receptor expressing cells in culture. Cells and tissues expressing TIE-2 receptor include, for example, cardiac and vascular endothelial cells, lens epithelium and heart epicardium. Alternatively, such ligand may be used to support cells which are engineered to express TIE-2 receptor. Further, TIE-2 ligands and their cognate receptor may be used in assay systems to identify agonists or antagonists of the TIE-2 receptor.

BACKGROUND OF THE INVENTION

The cellular behavior responsible for the development, maintenance, and repair of differentiated cells and tissues is regulated, in large part, by intercellular signals conveyed via growth factors and similar ligands and their receptors. The receptors are located on the cell surface of responding cells and they bind peptides or polypeptides known as growth factors as well as other hormone-like ligands. The results of this interaction are rapid biochemical changes in the responding cells, as well as a rapid and a long-term readjustment of cellular gene expression. Several receptors associated with various cell surfaces may bind specific growth factors.

The phosphorylation of tyrosines on proteins by tyrosine kinases is one of the key modes by which signals are transduced across the plasma membrane. Several currently known protein tyrosine kinase genes encode transmembrane receptors for polypeptide growth factors and hormones such as epidermal growth factor (EGF), insulin, insulin-

like growth factor-I (IGF-I), platelet derived growth factors (PDGF-A and -B), and fibroblast growth factors (FGFs). (Heldin et al., Cell Regulation, 1: 555-566 (1990); Ullrich, et al., Cell, 61: 243-54 (1990)). In each instance, these growth factors exert their action by binding to the extracellular portion of their cognate receptors, which leads to activation of the intrinsic tyrosine kinase present on the cytoplasmic portion of the receptor. Growth factor receptors of endothelial cells are of particular interest due to the possible involvement of growth factors in several important physiological and pathological processes, such as vasculogenesis, angiogenesis, atherosclerosis, and inflammatory diseases. (Folkman, et al. Science, 235: 442-447 (1987)). Also, the receptors of several hematopoietic growth factors are tyrosine kinases; these include c-fms, which is the colony stimulating factor 1 receptor, Sherr, et al., Cell, 41: 665-676 (1985), and c-kit, a primitive hematopoietic growth factor receptor reported in Huang, et al., Cell, 63: 225-33 (1990).

The receptor tyrosine kinases have been divided into evolutionary subfamilies based on the characteristic structure of their ectodomains. (Ullrich, et al. Cell, 61: 243-54 (1990)). Such subfamilies include, EGF receptor-like kinase (subclass I) and insulin receptor-like kinase (subclass II), each of which contains repeated homologous cysteine-rich sequences in their extracellular domains. A single cysteine-rich region is also found in the extracellular domains of the eph-like kinases. Hirai, et al., Science, 238: 1717-1720 (1987); Lindberg, et al. Mol. Cell. Biol., 10: 6316-24 (1990); Lhotak, et al., Mol. Cell. Biol. 11: 2496-2502 (1991). PDGF receptors as well as c-fms and c-kit receptor tyrosine kinases may be grouped into subclass III; while the FGF receptors form subclass IV. Typical for the members of both of these subclasses are extracellular folding units stabilized by intrachain

disulfide bonds. These so-called immunoglobulin (Ig)-like folds are found in the proteins of the immunoglobulin superfamily which contains a wide variety of other cell surface receptors having either cell-bound or soluble ligands. Williams, et al., Ann. Rev. Immunol., 6: 381-405 (1988).

Receptor tyrosine kinases differ in their specificity and affinity. In general, receptor tyrosine kinases are glycoproteins, which consist of (1) an extracellular domain capable of binding the specific growth factor(s); (2) a transmembrane domain which usually is an alpha-helical portion of the protein; (3) a juxtamembrane domain where the receptor may be regulated by, e.g., protein phosphorylation; (4) a tyrosine kinase domain which is the enzymatic component of the receptor; and (5) a carboxyterminal tail which in many receptors is involved in recognition and binding of the substrates for the tyrosine kinase.

Processes such as alternative exon splicing and alternative choice of gene promoter or polyadenylation sites have been reported to be capable of producing several distinct polypeptides from the same gene. These polypeptides may or may not contain the various domains listed above. As a consequence, some extracellular domains may be expressed as separate, secreted proteins and some forms of the receptors may lack the tyrosine kinase domain and contain only the extracellular domain inserted in the plasma membrane via the transmembrane domain plus a short carboxyl terminal tail.

A gene encoding an endothelial cell transmembrane tyrosine kinase, originally identified by RT-PCR as an unknown tyrosine kinase-homologous cDNA fragment from human leukemia cells, was described by Partanen, et al., Proc. Natl. Acad. Sci. USA, 87: 8913-8917 (1990). This gene and its encoded protein are called "tie" which is an

abbreviation for "tyrosine kinase with Ig and EGF homology domains." Partanen, et al. Mol. Cell. Biol. 12: 1698-1707 (1992).

It has been reported that tie mRNA is present in all human fetal and mouse embryonic tissues. Upon inspection, tie message has been localized to the cardiac and vascular endothelial cells. tie mRNA has been localized to the endothelia of blood vessels and endocardium of 9.5 to 18.5 day old mouse embryos. Enhanced tie expression was shown during neovascularization associated with developing ovarian follicles and granulation tissue in skin wounds. Korhonen, et al. Blood 80: 2548-2555 (1992). Thus tie has been suggested to play a role in angiogenesis, which is important for developing treatments for solid tumors and several other angiogenesis-dependent diseases such as diabetic retinopathy, psoriasis, atherosclerosis and arthritis.

Two structurally related rat TIE receptor proteins have been reported to be encoded by distinct genes with related profiles of expression. One gene, termed tie-1, is the rat homolog of human tie. Maisonpierre, et al., Oncogene 8: 1631-1637 (1993). The other gene, tie-2, may be the rat homolog of the murine tek gene, which, like tie, has been reported to be expressed in the mouse exclusively in endothelial cells and their presumptive progenitors. Dumont, et al. Oncogene 8: 1293-1301 (1993).

Both genes were found to be widely expressed in endothelial cells of embryonic and postnatal tissues. Significant levels of tie-2 transcripts were also present in other embryonic cell populations, including lens epithelium, heart epicardium and regions of mesenchyme. Maisonpierre, et al., Oncogene 8: 1631-1637 (1993).

The predominant expression of the TIE receptor in vascular endothelia suggests that TIE plays a role in the development and maintenance of the vascular system. This could include roles in

endothelial cell determination, proliferation, differentiation and cell migration and patterning into vascular elements. It has been reported that analyses of mouse embryos deficient in TIE-2 showed that it is important in angiogenesis, particular for vascular network formation in endothelial cells. Sato, T.N., et al., Nature 376:70-74 (1994). In the mature vascular system, TIE could function in endothelial cell survival, maintenance and response to pathogenic influences.

SUMMARY OF THE INVENTION

The present invention provides for a composition comprising a TIE-2 ligand substantially free of other proteins. The invention also provides for an isolated nucleic acid molecule encoding a TIE-2 ligand. The isolated nucleic acid may be DNA, cDNA or RNA. The invention also provides for a vector comprising an isolated nucleic acid molecule encoding a TIE-2 ligand. The invention further provides for a host-vector system for the production in a suitable host cell of a polypeptide having the biological activity of a TIE-2 ligand. The suitable host cell may be bacterial, yeast, insect or mammalian. The invention also provides for a method of producing a polypeptide having the biological activity of a TIE-2 ligand which comprises growing cells of the host-vector system under conditions permitting production of the polypeptide and recovering the polypeptide so produced.

The invention herein described of an isolated nucleic acid molecule encoding a TIE-2 ligand further provides for the development of the ligand, a fragment or derivative thereof, or another molecule which is a receptor agonist or antagonist, as a therapeutic for the treatment of patients suffering from disorders involving cells, tissues or organs which express the TIE receptor. The present invention also provides for an antibody which specifically binds such a therapeutic molecule. The antibody may be monoclonal or polyclonal. The invention also provides for a method of using such a monoclonal or polyclonal antibody to measure the amount of the therapeutic molecule in a sample taken from a patient for purposes of monitoring the course of therapy.

The present invention also provides for an antibody which specifically binds a TIE-2 ligand. The antibody may be monoclonal or

polyclonal. Thus the invention further provides for therapeutic compositions comprising an antibody which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle. The invention also provides for a method of blocking blood vessel growth in a mammal by administering an effective amount of a therapeutic composition comprising an antibody which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle.

The invention further provides for therapeutic compositions comprising a TIE-2 ligand in a pharmaceutically acceptable vehicle. The invention also provides for a method of promoting neovascularization in a patient by administering an effective amount of a therapeutic composition comprising a TIE-2 ligand in a pharmaceutically acceptable vehicle. In one embodiment, the method may be used to promote wound healing. In another embodiment, the method may be used to treat ischemia.

Alternatively, the invention provides that a TIE-2 ligand may be conjugated to a cytotoxic agent and a therapeutic composition prepared therefrom. The invention further provides for a receptor body which specifically binds a TIE-2 ligand. The invention further provides for therapeutic compositions comprising a receptor body which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle. The invention also provides for a method of blocking blood vessel growth in a mammal by administering an effective amount of a therapeutic composition comprising a receptor body which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle.

The invention also provides for a TIE-2 receptor antagonist as well as a method of inhibiting TIE-2 ligand biological activity in a mammal comprising administering to the mammal an effective amount of a TIE-2 antagonist. According to the invention, the antagonist may

be an antibody or other molecule capable of specifically binding either TIE-2 ligand or TIE-2 receptor. For example, the antagonist may be a TIE-2 receptorbody.

BRIEF DESCRIPTION OF THE FIGURES

FIGURES 1A and 1B - TIE-2 receptorbody (TIE-2 RB) inhibits the development of blood vessels in the embryonic chicken chorioallantoic membrane (CAM). A single piece of resorbable gelatin foam (Gelfoam) soaked with 6 μg of RB was inserted immediately under the CAM of 1-day chick embryos. After 3 further days of incubation, 4 day old embryos and surrounding CAM were removed and examined. FIGURE 1A: embryos treated with EHK-1 RB (rEHK-1 ecto / h IgG1 Fc) were viable and possessed normally developed blood vessels in their surrounding CAM. FIGURE 1B : all embryos treated with TIE-2 RB (r TIE-2 ecto / h IgG1 Fc) were dead, diminished in size and were almost completely devoid of surrounding blood vessels.

FIGURE 2 - Vector pJFE14.

FIGURE 3 - Restriction map of λgt10.

FIGURE 4 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand from clone λgt10 encoding htie-2 ligand 1.

FIGURE 5 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand from T98G clone.

FIGURE 6 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand from clone pBluescript KS encoding human TIE 2 ligand 2.

FIGURE 7 - Western blot showing activation of TIE-2 receptor by TIE-2 ligand 1 (Lane L1) but not by TIE-2 ligand 2 (Lane L2) or control (Mock).

FIGURE 8 - Western blot showing that prior treatment of HAEC cells with excess TIE-2 ligand 2 (Lane 2) antagonizes the subsequent ability of dilute TIE-2 ligand 1 to activate the TIE-2 receptor (TIE2-R) as compared with prior treatment of HAEC cells with MOCK medium (Lane 1).

FIGURE 9 - Histogram representation of binding to rat TIE2 IgG immobilized surface by TIE2 ligand in C2C12 ras, Rat2 ras, SHEP, and T98G concentrated (10x) conditioned medium. Rat TIE2 (rTIE2) specific binding is demonstrated by the significant reduction in the binding activity in the presence of 25 µg/ml soluble rat TIE2 RB as compared to a minor reduction in the presence of soluble trkB RB.

FIGURE 10 - Binding of recombinant human TIE-2 ligand 1 (hTL1) and human TIE-2 ligand 2 (hTL2), in cos cell supernatants, to a human TIE-2 RB immobilized surface. Human TIE-2-specific binding was determined by incubating the samples with 25 µg/ml of either soluble human TIE2 RB or trkB RB; significant reduction in the binding activity is observed only for the samples incubated with human TIE2 RB.

FIGURE 11 - Western blot showing that TIE-2 receptorbody (denoted TIE-2 RB or, as here, TIE2-Fc) blocks the activation of TIE-2 receptors by TIE-2 ligand 1 (TL1) in HUVEC cells, whereas an unrelated receptorbody (TRKB-Fc) does not block this activation.

DETAILED DESCRIPTION OF THE INVENTION

As described in greater detail below, applicants have isolated, by expression cloning, a novel ligand that binds the TIE-2 receptor. The present invention comprises a TIE-2 ligand as well as its amino acid sequence and also functionally equivalent molecules in which amino acid residues are substituted for residues within the sequence resulting in a silent change. For example, one or more amino acid residues within the sequence can be substituted by another amino acid(s) of a similar polarity which acts as a functional equivalent, resulting in a silent alteration. Substitutes for an amino acid within the sequence may be selected from other members of the class to which the amino acid belongs. For example, the class of nonpolar (hydrophobic) amino acids include alanine, leucine, isoleucine, valine, proline, phenylalanine, tryptophan and methionine. The polar neutral amino acids include glycine, serine, threonine, cysteine, tyrosine, asparagine, and glutamine. The positively charged (basic) amino acids include arginine, lysine and histidine. The negatively charged (acidic) amino acids include aspartic acid and glutamic acid. Also included within the scope of the invention are proteins or fragments or derivatives thereof which exhibit the same or similar biological activity and derivatives which are differentially modified during or after translation, e.g., by glycosylation, proteolytic cleavage, linkage to an antibody molecule or other cellular ligand, etc.

The present invention also encompasses the nucleotide sequence that encodes the protein described herein as TIE-2 ligand 1, as well as cells which are genetically engineered to produce the protein, by e.g. transfection, transduction, infection, electroporation, or microinjection of nucleic acid encoding the TIE-2 ligand 1 described herein in a suitable expression vector.

The present invention further encompasses the nucleotide sequence that encodes the protein described herein as TIE-2 ligand 2, as well as cells which are genetically engineered to produce the protein, by e.g. transfection, transduction, infection, electroporation, or microinjection of nucleic acid encoding the TIE-2 ligand 2 described herein in a suitable expression vector.

One skilled in the art will also recognize that the present invention encompasses DNA and RNA sequences that hybridize to a deduced TIE-2 ligand encoding sequence, under conditions of moderate stringency, as defined in, for example, Sambrook, et al. Molecular Cloning: A Laboratory Manual, 2 ed. Vol. 1, pp. 101-104, Cold Spring Harbor Laboratory Press (1989). Thus, a nucleic acid molecule contemplated by the invention includes one having a sequence deduced from an amino acid sequence of a TIE-2 ligand prepared as described herein, as well as a molecule having a sequence of nucleic acids that hybridizes to such a nucleic acid sequence, and also a nucleic acid sequence which is degenerate of the above sequences as a result of the genetic code, but which encodes a ligand that binds the TIE-2 receptor.

Any of the methods known to one skilled in the art for the insertion of DNA fragments into a vector may be used to construct expression vectors encoding TIE-2 ligand using appropriate transcriptional/translational control signals and the protein coding sequences. These methods may include in vitro recombinant DNA and

synthetic techniques and in vivo recombinations (genetic recombination). Expression of a nucleic acid sequence encoding a TIE-2 ligand or peptide fragments thereof may be regulated by a second nucleic acid sequence so that the protein or peptide is expressed in a host transformed with the recombinant DNA molecule. For example, expression of a TIE-2 ligand described herein may be controlled by any promoter/enhancer element known in the art. Promoters which may be used to control expression of the ligand include, but are not limited to the long terminal repeat as described in Squinto et al., (Cell 65:1-20 (1991)); the SV40 early promoter region (Bernoist and Chambon, Nature 290:304-310), the CMV promoter, the M-MuLV 5' terminal repeat, the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto, et al., Cell 22:787-797 (1980)), the herpes thymidine kinase promoter (Wagner et al., Proc. Natl. Acad. Sci. U.S.A. 78:144-1445 (1981)), the adenovirus promoter, the regulatory sequences of the metallothioein gene (Brinster et al., Nature 296:39-42 (1982)); prokaryotic expression vectors such as the β-lactamase promoter (Villa-Kamaroff, et al., Proc. Natl. Acad. Sci. U.S.A. 75:3727-3731 (1978)), or the tac promoter (DeBoer, et al., Proc. Natl. Acad. Sci. U.S.A. 80:21-25 (1983)), see also "Useful proteins from recombinant bacteria" in Scientific American, 242:74-94 (1980); promoter elements from yeast or other fungi such as the Gal 4 promoter, the ADH (alcohol dehydrogenase) promoter, PGK (phosphoglycerol kinase) promoter, alkaline phosphatase promoter, and the following animal transcriptional control regions, which exhibit tissue specificity and have been utilized in transgenic animals: elastase I gene control region which is active in pancreatic acinar cells (Swift et al., Cell 38:639-646 (1984); Ornitz et al., Cold Spring Harbor Symp. Quant. Biol. 50:399-409 (1986); MacDonald, Hepatology 7:425-515 (1987); insulin gene

control region which is active in pancreatic beta cells (Hanahan, Nature 315:115-122 (1985)), immunoglobulin gene control region which is active in lymphoid cells (Grosschedl et al., 1984, Cell 38:647-658; Adames et al., 1985, Nature 318:533-538; Alexander et al., 1987, Mol. Cell. Biol. 7:1436-1444), mouse mammary tumor virus control region which is active in testicular, breast, lymphoid and mast cells (Leder et al., 1986, Cell 45:485-495), albumin gene control region which is active in liver (Pinkert et al., 1987, Genes and Devel. 1:268-276), alpha-fetoprotein gene control region which is active in liver (Krumlauf et al., 1985, Mol. Cell. Biol. 5:1639-1648; Hammer et al., 1987, Science 235:53-58); alpha 1-antitrypsin gene control region which is active in the liver (Kelsey et al, 1987, Genes and Devel. 1:161-171), beta-globin gene control region which is active in myeloid cells (Mogram et al., 1985, Nature 315:338-340; Kollias et al., 1986, Cell 46:89-94); myelin basic protein gene control region which is active in oligodendrocyte cells in the brain (Readhead et al., 1987, Cell 48:703-712); myosin light chain-2 gene control region which is active in skeletal muscle (Shani, 1985, Nature 314:283-286), and gonadotropic releasing hormone gene control region which is active in the hypothalamus (Mason et al., 1986, Science 234:1372-1378). The invention further encompasses the production of antisense compounds which are capable of specifically hybridizing with a sequence of RNA encoding a TIE-2 ligand to modulate its expression. (Ecker, U.S. Patent No. 5,166,195, issued November 24, 1992).

Thus, according to the invention, expression vectors capable of being replicated in a bacterial or eukaryotic host comprising a nucleic acid encoding a TIE-2 ligand as described herein, are used to transfect a host and thereby direct expression of such nucleic acid to produce the TIE-2 ligand, which may then be recovered in a biologically active form.

As used herein, a biologically active form includes a form capable of binding to the TIE-2 receptor and causing a biological response such as a differentiated function or influencing the phenotype of the cell expressing the receptor. Such biologically active forms would, for example, induce phosphorylation of the tyrosine kinase domain of the TIE-2 receptor.

Expression vectors containing the gene inserts can be identified by four general approaches: (a) DNA-DNA hybridization, (b) presence or absence of "marker" gene functions, (c) expression of inserted sequences and (d) PCR detection. In the first approach, the presence of a foreign gene inserted in an expression vector can be detected by DNA-DNA hybridization using probes comprising sequences that are homologous to an inserted TIE-2 ligand encoding gene. In the second approach, the recombinant vector/host system can be identified and selected based upon the presence or absence of certain "marker" gene functions (e.g., thymidine kinase activity, resistance to antibiotics, transformation phenotype, occlusion body formation in baculovirus, etc.) caused by the insertion of foreign genes in the vector. For example, if a nucleic acid encoding a TIE-2 ligand is inserted within the marker gene sequence of the vector, recombinants containing the insert can be identified by the absence of the marker gene function. In the third approach, recombinant expression vectors can be identified by assaying the foreign gene product expressed by the recombinant. Such assays can be based, for example, on the physical or functional properties of a TIE-2 ligand gene product, for example, by binding of the ligand to the TIE-2 receptor or portion thereof which may be tagged with, for example, a detectable antibody or portion thereof or by binding to antibodies produced against the TIE-2 ligand protein or a portion thereof. Cells of the present invention may transiently or,

preferably, constitutively and permanently express TIE-2 ligands as described herein. In the fourth approach, DNA nucleotide primers can be prepared corresponding to a tie-2 specific DNA sequence. These primers could then be used to PCR a tie-2 gene fragment. (PCR Protocols: A Guide To Methods and Applications, Edited by Michael A. Innis et al., Academic Press (1990)).

The recombinant ligand may be purified by any technique which allows for the subsequent formation of a stable, biologically active protein. For example, and not by way of limitation, the ligand may be recovered from cells either as soluble proteins or as inclusion bodies, from which they may be extracted quantitatively by 8M guanidinium hydrochloride and dialysis. In order to further purify the ligand, conventional ion exchange chromatography, hydrophobic interaction chromatography, reverse phase chromatography or gel filtration may be used.

In additional embodiments of the invention, a recombinant TIE-2 ligand encoding gene may be used to inactivate or "knock out" the endogenous gene by homologous recombination, and thereby create a TIE-2 ligand deficient cell, tissue, or animal. For example, and not by way of limitation, the recombinant TIE-2 ligand encoding gene may be engineered to contain an insertional mutation, for example the neo gene, which would inactivate the native TIE-2 ligand encoding gene. Such a construct, under the control of a suitable promoter, may be introduced into a cell, such as an embryonic stem cell, by a technique such as transfection, transduction, or injection. Cells containing the construct may then be selected by G418 resistance. Cells which lack an intact TIE-2 ligand encoding gene may then be identified, e.g. by Southern blotting, PCR detection, Northern blotting or assay of expression. Cells lacking an intact TIE-2 ligand encoding gene may then

be fused to early embryo cells to generate transgenic animals deficient in such ligand. Such an animal may be used to define specific in vivo processes, normally dependent upon the ligand.

The present invention also provides for antibodies to the TIE-2 ligands described herein which are useful for detection of the ligands in, for example, diagnostic applications. For preparation of monoclonal antibodies directed toward TIE-2 ligand, any technique which provides for the production of antibody molecules by continuous cell lines in culture may be used. For example, the hybridoma technique originally developed by Kohler and Milstein (1975, Nature 256:495-497), as well as the trioma technique, the human B-cell hybridoma technique (Kozbor et al., 1983, Immunology Today 4:72), and the EBV-hybridoma technique to produce human monoclonal antibodies (Cole et al., 1985, in "Monoclonal Antibodies and Cancer Therapy," Alan R. Liss, Inc. pp. 77-96) and the like are within the scope of the present invention.

The monoclonal antibodies may be human monoclonal antibodies or chimeric human-mouse (or other species) monoclonal antibodies. Human monoclonal antibodies may be made by any of numerous techniques known in the art (e.g., Teng et al., 1983, Proc. Natl. Acad. Sci. U.S.A. 80:7308-7312; Kozbor et al., 1983, Immunology Today 4:72-79; Olsson et al., 1982, Meth. Enzymol. 92:3-16). Chimeric antibody molecules may be prepared containing a mouse antigen-binding domain with human constant regions (Morrison et al., 1984, Proc. Natl. Acad. Sci. U.S.A. 81:6851, Takeda et al., 1985, Nature 314:452).

Various procedures known in the art may be used for the production of polyclonal antibodies to epitopes of the TIE-2 ligands described herein. For the production of antibody, various host animals can be immunized by injection with a TIE-2 ligand, or a fragment or derivative thereof, including but not limited to rabbits, mice and rats.

Various adjuvants may be used to increase the immunological response, depending on the host species, and including but 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.

A molecular clone of an antibody to a selected TIE-2 ligand epitope can be prepared by known techniques. Recombinant DNA methodology (see e.g., Maniatis et al., 1982, Molecular Cloning, A Laboratory Manual, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York) may be used to construct nucleic acid sequences which encode a monoclonal antibody molecule, or antigen binding region thereof.

The present invention provides for antibody molecules as well as fragments of such antibody molecules. Antibody fragments which contain the idiotype of the molecule can be generated by known techniques. For example, such fragments include but are not limited to: the $F(ab')_2$ fragment which can be produced by pepsin digestion of the antibody molecule; the Fab' fragments which can be generated by reducing the disulfide bridges of the $F(ab')_2$ fragment, and the Fab fragments which can be generated by treating the antibody molecule with papain and a reducing agent. Antibody molecules may be purified by known techniques, e.g., immunoabsorption or immunoaffinity chromatography, chromatographic methods such as HPLC (high performance liquid chromatography), or a combination thereof.

The present invention further encompasses an immunoassay for measuring the amount of a TIE-2 ligand in a biological sample by

a) contacting the biological sample with at least one antibody which specifically binds the TIE-2 ligand so that the antibody forms a complex with any TIE-2 ligand present in the sample; and

b) measuring the amount of the complex and thereby measuring the amount of the TIE-2 ligand in the biological sample.

The invention further encompasses an assay for measuring the amount of TIE-2 receptor in a biological sample by

a) contacting the biological sample with at least one ligand of the invention so that the ligand forms a complex with the TIE-2 receptor; and

b) measuring the amount of the complex and thereby measuring the amount of the TIE-2 receptor in the biological sample.

The present invention also provides for the utilization of a TIE-2 ligand to support the survival and/or growth and/or differentiation of TIE-2 receptor expressing cells. Thus, the ligand may be used as a supplement to support, for example, endothelial cells in culture. Further, the discovery by applicants of a cognate ligand for the TIE-2 receptor enables the utilization of assay systems useful for the identification of agonists or antagonists of the TIE-2 receptor. Such assay systems would be useful in identifying molecules capable of promoting or inhibiting angiogenesis. For example, in one embodiment, antagonists of the TIE-2 receptor may be identified as test molecules that are capable of interfering with the interaction of the TIE-2 receptor with a biologically active TIE-2 ligand. Such antagonists are identified by their ability to 1) block the binding of a biologically

active TIE-2 ligand to the receptor as measured, for example, using BIAcore biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ); or 2) block the ability of a biologically active TIE-2 ligand to cause a biological response. Such biological responses include, but are not limited to, phosphorylation of the TIE-2 receptor or downstream components of the TIE-2 signal transduction pathway, or survival, growth or differentiation of TIE-2 receptor bearing cells.

In one embodiment, cells engineered to express the TIE-2 receptor may be dependent for growth on the addition of TIE-2 ligand. Such cells provide useful assay systems for identifying additional agonists of the TIE-2 receptor, or antagonists capable of interfering with the activity of TIE-2 ligand on such cells. Alternatively, autocrine cells, engineered to be capable of co-expressing both TIE-2 ligand and receptor, may provide useful systems for assaying potential agonists or antagonists.

Therefore, the present invention provides for introduction of the TIE-2 receptor into cells that do not normally express this receptor, thus allowing these cells to exhibit profound and easily distinguishable responses to a ligand which binds this receptor. The type of response elicited depends on the cell utilized, and not the specific receptor introduced into the cell. Appropriate cell lines can be chosen to yield a response of the greatest utility for assaying, as well as discovering, molecules that can act on tyrosine kinase receptors. The molecules may be any type of molecule, including but not limited to peptide and non-peptide molecules, that will act in systems to be described in a receptor specific manner.

One of the more useful systems to be exploited involves the introduction of the TIE-2 receptor into a fibroblast cell line (e.g., NIH3T3 cells) thus such a receptor which does not normally mediate

proliferative responses can, following introduction into fibroblasts, nonetheless be assayed by a variety of well established methods to quantitate effects of fibroblast growth factors (e.g. thymidine incorporation or other types of proliferation assays; see van Zoelen, 1990, "The Use of Biological Assays For Detection Of Polypeptide Growth Factors" in Progress Factor Research, Vol. 2, pp. 131-152; Zhan and M. Goldfarb, 1986, Mol. Cell. Biol., Vol. 6, pp. 3541-3544). These assays have the added advantage that any preparation can be assayed both on the cell line having the introduced receptor as well as the parental cell line lacking the receptor; only specific effects on the cell line with the receptor would be judged as being mediated through the introduced receptor. Such cells may be further engineered to express the TIE-2 ligand, thus creating an autocrine system useful for assaying for molecules that act as antagonists/agonists of this interaction. Thus, the present invention provides for host cells comprising nucleic acid encoding TIE-2 ligand and nucleic acid encoding TIE-2 receptor.

The TIE-2 receptor/TIE-2 ligand interaction also provides a useful system for identifying small molecule agonists or antagonists of the TIE-2 receptor. For example, fragments, mutants or derivatives of a TIE-2 ligand may be identified that bind the TIE-2 receptor but do not induce biological activity. Alternatively, the characterization of a TIE-2 ligand enables the determination of active portions of the molecule. Further, the identification of a ligand enables the determination of the X-ray crystal structure of the receptor/ligand complex, thus enabling identification of the binding site on the receptor. Knowledge of the binding site will provide useful insight into the rational design of novel agonists and antagonists.

The specific binding of a test molecule to the TIE-2 receptor may be measured in a number of ways. For example, the actual binding of

test molecule to cells expressing tie-2 may be detected or measured, by detecting or measuring (i) test molecule bound to the surface of intact cells; (ii) test molecule cross-linked to TIE-2 protein in cell lysates; or (iii) test molecule bound to TIE-2 in vitro. The specific interaction between test molecule and TIE-2 may be evaluated by using reagents that demonstrate the unique properties of that interaction.

As a specific, nonlimiting example, the methods of the invention may be used as follows. Consider a case in which the TIE-2 ligand in a sample is to be measured. Varying dilutions of the sample (the test molecule), in parallel with a negative control (NC) containing no TIE-2 ligand activity, and a positive control (PC) containing a known amount of a TIE-2 ligand, may be exposed to cells that express tie-2 in the presence of a detectably labeled TIE-2 ligand (in this example, radioiodinated ligand). The amount of TIE-2 ligand in the test sample may be evaluated by determining the amount of ^{125}I-labeled TIE-2 ligand that binds to the controls and in each of the dilutions, and then comparing the sample values to a standard curve. The more TIE-2 ligand in the sample, the less ^{125}I-ligand that will bind to TIE-2.

The amount of ^{125}I-ligand bound may be determined by measuring the amount of radioactivity per cell, or by cross-linking the TIE-2 ligand to cell surface proteins using DSS, as described in Meakin and Shooter, 1991, Neuron 6:153-163, and detecting the amount of labeled protein in cell extracts using, for example, SDS polyacrylamide gel electrophoresis, which may reveal a labeled protein having a size corresponding to TIE-2 ligand/TIE-2 receptor. The specific test molecule/TIE-2 interaction may further be tested by adding to the assays various dilutions of an unlabeled control ligand that does not bind the TIE-2 receptor and therefore should have no substantial affect on the competition between labeled TIE-2 ligand and test molecule for

TIE-2 binding. Alternatively, a molecule known to be able to disrupt TIE-2 ligand/TIE-2 binding, such as, but not limited to, anti-TIE-2 antibody, or TIE-2 receptorbody as described herein, may be expected to interfere with the competition between ^{125}I-TIE-2 ligand and test molecule for TIE-2 receptor binding.

Detectably labeled TIE-2 ligand includes, but is not limited to, TIE-2 ligand linked covalently or noncovalently to a radioactive substance, a fluorescent substance, a substance that has enzymatic activity, a substance that may serve as a substrate for an enzyme (enzymes and substrates associated with colorimetrically detectable reactions are preferred) or to a substance that can be recognized by an antibody molecule that is preferably a detectably labeled antibody molecule.

Alternatively, the specific binding of test molecule to TIE-2 may be measured by evaluating the secondary biological effects of TIE-2 ligand/TIE-2 receptor binding, including, but not limited to, cell growth and/or differentiation or immediate early gene expression or phosphorylation of TIE-2. For example, the ability of the test molecule to induce differentiation can be tested in cells that lack tie-2 and in comparable cells that express tie-2; differentiation in tie-2-expressing cells but not in comparable cells that lack tie-2 would be indicative of a specific test molecule/TIE-2 interaction. A similar analysis could be performed by detecting immediate early gene (e.g. fos and jun) induction in tie-2-minus and tie-2-plus cells, or by detecting phosphorylation of TIE-2 using standard phosphorylation assays known in the art. Such analysis might be useful in identifying agonists or antagonists that do not competitively bind to TIE-2.

Similarly, the present invention provides for a method of identifying an molecule that has the biological activity of a TIE-2

ligand comprising (i) exposing a cell that expresses tie-2 to a test molecule and (ii) detecting the specific binding of the test molecule to TIE-2 receptor, in which specific binding to TIE-2 positively correlates with TIE-2 like activity. Specific binding may be detected by either assaying for direct binding or the secondary biological effects of binding, as discussed supra. Such a method may be particularly useful in identifying new members of the TIE ligand family or, in the pharmaceutical industry, in screening a large array of peptide and non-peptide molecules (e.g., peptidomimetics) for TIE associated biological activity. In a preferred, specific, nonlimiting embodiment of the invention, a large grid of culture wells may be prepared that contain, in alternate rows, PC12 (or fibroblasts, see infra) cells that are either tie-2-minus or engineered to be tie-2-plus. A variety of test molecules may then be added such that each column of the grid, or a portion thereof, contains a different test molecule. Each well could then be scored for the presence or absence of growth and/or differentiation. An extremely large number of test molecules could be screened for such activity in this manner.

In additional embodiments, the invention provides for methods of detecting or measuring TIE-like activity or identifying a molecule as having such activity comprising (i) exposing a test molecule to a TIE-2 receptor protein in vitro under conditions that permit binding to occur and (ii) detecting binding of the test molecule to the TIE-2 protein, in which binding of test molecule to TIE-2 correlates with TIE-like activity. According to such methods, the TIE-2 may or may not be substantially purified, may be affixed to a solid support (e.g. as an affinity column or as an ELISA assay), or may be incorporated into an artificial membrane. Binding of test molecule to TIE-2 may be evaluated by any method known in the art. In preferred embodiments,

the binding of test molecule may be detected or measured by evaluating its ability to compete with detectably labeled known TIE-2 ligands for TIE-2 receptor binding.

The present invention also provides for a method of detecting the ability of a test molecule to function as an antagonist of TIE-like activity comprising detecting the ability of the molecule to inhibit an effect of TIE ligand binding to TIE-2 on a cell that expresses tie-2. Such an antagonist may or may not interfere with TIE-2 ligand/TIE-2 receptor binding. Effects of TIE-2 ligand binding to TIE-2 receptor are preferably biological or biochemical effects, including, but not limited to, cell survival or proliferation, cell transformation, immediate early gene induction, or TIE-2 phosphorylation.

The invention further provides for both a method of identifying antibodies or other molecules capable of neutralizing the ligand or blocking binding to the receptor, as well as the molecules identified by the method. By way of nonlimiting example, the method may be performed via an assay which is conceptually similar to an ELISA assay. For example, TIE receptorbody may be bound to a solid support, such as a plastic multiwell plate. As a control, a known amount of TIE ligand which has been Myc-tagged may then be introduced to the well and any tagged TIE ligand which binds the receptorbody may then be identified by means of a reporter antibody directed against the Myc-tag. This assay system may then be used to screen test samples for molecules which are capable of i) binding to the tagged ligand or ii) binding to the receptorbody and thereby blocking binding to the receptorbody by the tagged ligand. For example, a test sample containing a putative molecule of interest together with a known amount of tagged ligand may be introduced to the well and the amount of tagged ligand which binds to the receptorbody may be measured. By

comparing the amount of bound tagged ligand in the test sample to the amount in the control, samples containing molecules which are capable of blocking ligand binding to the receptor may be identified. The molecules of interest thus identified may be isolated using methods well known to one of skill in the art.

Once a blocker of ligand binding is found, one of skill in the art would know to perform secondary assays to determine whether the blocker is binding to the receptor or to the ligand, as well as assays to determine if the blocker molecule can neutralize the biological activity of the ligand. For example, by using a binding assay which employs BIAcore biosensor technology (or the equivalent), in which either TIE receptorbody or TIE ligand is covalently attached to a solid support (e.g. carboxymethyl dextran on a gold surface), one of skill in the art would be able to determine if the blocker molecule is binding specifically to the ligand or to the receptorbody. To determine if the blocker molecule can neutralize the biological activity of the ligand, one of skill in the art could perform a phosphorylation assay (see Example 5) or alternatively, a functional bioassay, such as a survival assay, by using primary cultures of, for example, endothelial cells. Alternatively, a blocker molecule which binds to the receptorbody could be an agonist and one of skill in the art would know to how to determine this by performing an appropriate assay for identifying additional agonists of the TIE-2 receptor.

Because TIE-2 receptor has been identified in association with endothelial cells and, as demonstrated herein, blocking of the ligand appears to prevent vascularization, applicants have demonstrated that the TIE-2 ligand will be useful for the induction of vascularization in diseases or disorders where such vascularization is indicated. Such diseases or disorders would include wound healing, ischaemia and

diabetes. On the other hand, antagonists of the TIE-2 receptor, such as receptorbodies as described herein in Examples 2 and 3, and TIE-2 ligand 2 as described in Example 9, would be useful to prevent or attenuate vascularization, thus preventing or attenuating, for example, tumor growth.

Because TIE-2 receptor bearing cells appear to be upregulated in vascularization, TIE-2 ligandbodies may also be useful in preventing or attenuating, for example, tumor growth. TIE-2 ligandbodies may be useful for the delivery of toxins to a receptor bearing cell. Alternatively, other molecules, such as growth factors, cytokines or nutrients, may be delivered to a TIE-2 receptor bearing cell via TIE-2 ligandbodies. TIE-2 ligandbodies could also be used as a diagnostic reagent for the TIE-2 receptor, to detect the receptor in vivo or in vitro. Where the TIE-2 receptor is associated with a disease state, TIE-2 ligandbodies may be useful as diagnostic reagents for detecting the disease by, for example, tissue staining or whole body imaging.

The present invention also provides for pharmaceutical compositions comprising the TIE-2 ligands or ligandbodies described herein, peptide fragments thereof, or derivatives in a pharmacologically acceptable vehicle. The TIE-2 ligand proteins, peptide fragments, or derivatives may be administered systemically or locally. Any appropriate mode of administration known in the art may be used, including, but not limited to, intravenous, intrathecal, intraarterial, intranasal, oral, subcutaneous, intraperitoneal, or by local injection or surgical implant. Sustained release formulations are also provided for.

The present invention further provides for an isolated and purified nucleic acid molecule comprising a nucleic acid sequence encoding a human TIE-2 ligand, wherein the nucleic acid sequence is selected from the group consisting of:

(a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4, Figure 5 or Figure 6;

(b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; and

(c) a nucleic acid sequence that is degenerate as a result of the genetic code to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

In another embodiment the invention provides for an isolated nucleic acid molecule encoding a TIE-2 ligand wherein the nucleic acid sequence is:

(a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4, Figure 5 or Figure 6;

(b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or

(c) a nucleic acid sequence which, but for the degeneracy of the genetic code would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

The present invention further provides for an isolated and purified human TIE-2 ligand encoded by an isolated nucleic acid molecule of the invention. The invention also provides a vector which comprises an

isolated nucleic acid molecule comprising a nucleic acid sequence encoding a human TIE-2 ligand. In one embodiment, the vector is designated as pBluescript KS encoding human TIE 2 ligand 2.

The invention further provides for an expression vector comprising a DNA molecule encoding a human TIE-2 ligand, wherein the DNA molecule is operatively linked to an expression control sequence. The invention also provides a host-vector system for the production of a polypeptide having the biological activity of a human TIE-2 ligand which comprises the expression vector of the invention in a suitable host cell. In one embodiment, the suitable host cell may be a bacterial cell, yeast cell, insect cell, or mammalian cell. The invention further provides for a method of producing a polypeptide having the activity of a biologically active TIE-2 ligand which comprises growing cells of the host-vector system of the invention, under conditions permitting production of the polypeptide and recovering the polypeptide so produced.

The invention herein described of an isolated nucleic acid molecule encoding a TIE-2 ligand further provides for the development of the ligand, a fragment or derivative thereof, or another molecule which is a receptor agonist or antagonist, as a therapeutic for the treatment of patients suffering from disorders involving cells, tissues or organs which express the TIE receptor. By way of nonlimiting example, the present invention provides for a ligandbody which specifically binds the TIE-2 receptor or the TIE-2. In an alternative embodiment, the present invention provides for a ligandbody which comprises a TIE-2 ligand fused to an immunoglobulin constant region. In one embodiment, the ligandbody comprises TIE-2 ligand 1 or TIE-2 ligand 2 and the Fc portion of human IgG1. The ligandbody may be used

in a method of treatment of the human or animal body, or in a method of diagnosis.

The present invention also provides for an antibody which specifically binds such a therapeutic molecule. The antibody may be monoclonal or polyclonal. The invention also provides for a method of using such a monoclonal or polyclonal antibody to measure the amount of the therapeutic molecule in a sample taken from a patient for purposes of monitoring the course of therapy.

The invention further provides for a therapeutic composition comprising a human TIE-2 ligand or ligandbody and a cytotoxic agent conjugated thereto. In one embodiment, the cytotoxic agent may be a radioisotope or toxin.

The invention also provides for an antibody which specifically binds a human TIE-2 ligand. The antibody may be monoclonal or polyclonal.

The invention further provides for a method of purifying a human TIE-2 ligand comprising:

a) coupling at least one TIE-2 binding substrate to a solid matrix;

b) incubating the substrate of a) with a cell lysate so that the substrate forms a complex with any human TIE-2 ligand in the cell lysate;

c) washing the solid matrix; and

d) eluting the human TIE-2 ligand from the coupled substrate.

The substrate may be any substance that specifically binds the human TIE-2 ligand. In one embodiment, the substrate is selected from the group consisting of anti-TIE-2 ligand antibody, TIE-2 receptor and TIE-2 receptorbody. The invention further provides for a receptorbody which specifically binds a human TIE-2 ligand, as well as a therapeutic

composition comprising the receptorbody in a pharmaceutically acceptable vehicle, and a method of blocking blood vessel growth in a human comprising administering an effective amount of the therapeutic composition.

The invention also provides for a therapeutic composition comprising a human TIE-2 ligand or ligandbody in a pharmaceutically acceptable vehicle, as well as a method of promoting neovascularization in a patient comprising administering to the patient an effective amount of the therapeutic composition.

In addition, the present invention provides for a method for identifying a cell which expresses TIE-2 receptor which comprises contacting a cell with a detectably labeled TIE-2 ligand or ligandbody, under conditions permitting binding of the detectably labeled ligand to the TIE-2 receptor and determining whether the detectably labeled ligand is bound to the TIE-2 receptor, thereby identifying the cell as one which expresses TIE-2 receptor. The present invention also provides for a therapeutic composition comprising a TIE-2 ligand or ligandbody and a cytotoxic agent conjugated thereto. The cytotoxic agent may be a radioisotope or toxin.

The invention also provides a method of detecting expression of TIE-2 ligand by a cell which comprises obtaining mRNA from the cell, contacting the mRNA so obtained with a labelled nucleic acid molecule encoding a TIE-2 ligand, under hybridizing conditions, determining the presence of mRNA hybridized to the labelled molecule, and thereby detecting the expression of the TIE-2 ligand in the cell.

The invention further provides a method of detecting expression of a TIE-2 ligand in tissue sections which comprises contacting the tissue sections with a labelled nucleic acid molecule encoding a TIE-2 ligand, under hybridizing conditions, determining the presence of mRNA

hybridized to the labelled molecule, and thereby detecting the expression of the TIE-2 ligand in tissue sections.

EXAMPLE 1 - IDENTIFICATION OF THE ABAE CELL LINE AS REPORTER CELLS FOR THE TIE-2 RECEPTOR

Adult BAE cells are registered in the European Cell Culture Repository, under ECACC#92010601. (See PNAS 75:2621 (1978)). Northern (RNA) analyses revealed moderate levels of tie-2 transcripts in the ABAE (Adult Bovine Arterial Endothelial) cell line, consistent with in situ hybridization results that demonstrated almost exclusive localization of tie-2 RNAs to vascular endothelial cells. We therefore examined ABAE cell lysates for the presence of TIE-2 protein, as well as the extent to which this TIE-2 protein is tyrosine-phosphorylated under normal versus serum-deprived growth conditions. ABAE cell lysates were harvested and subjected to immunoprecipitation, followed by Western blot analyses of immunoprecipitated proteins with TIE-2 specific and phosphotyrosine-specific antisera. Omission or inclusion of TIE-2 peptides as specific blocking molecules during TIE-2 immunoprecipitation allowed unambiguous identification of TIE-2 as a moderately detectable protein of ~150 kD whose steady-state phosphotyrosine levels diminish to near undetectable levels by prior serum-starvation of the cells.

Culture of ABAE cells and harvest of cell lysates was done as follows. Low-passage-number ABAE cells were plated as a monolayer at a density of 2 x 10^6 cells/150mm plastic petri plate (Falcon) and cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% bovine calf serum (10 % BCS), 2 mM L-glutamine (Q) and 1% each of

penicillin and streptomycin (P-S) in an atmosphere of 5% CO_2. Prior to harvest of cell lysates, cells were serum-starved for 24 hours in DMEM/Q/P-S, followed by aspiration of the medium and rinsing of the plates with ice-cold phosphate buffered saline (PBS) supplemented with sodium orthovanadate, sodium fluoride and sodium benzamidine. Cells were lysed in a small volume of this rinse buffer that had been supplemented with 1% NP40 detergent and the protease inhibitors, PMSF and aprotinin. Insoluble debris was removed from the cell lysates by centrifugation at 14,000 xG for 10 minutes, at 4°C and the supernatants were subjected to immune-precipitation with antisera specific for TIE-2 receptor, with or without the presence of blocking peptides added to ~20 µg/ml lysate. Immunoprecipitated proteins were resolved by PAGE (7.5% Laemmli gel), and then electro-transferred to PVDF membrane and incubated either with various TIE-2- or phosphotyrosine-specific antisera. TIE-2-protein was visualized by incubation of the membrane with HRP-linked secondary antisera followed by treatment with ECL reagent (Amersham).

EXAMPLE 2 - CLONING AND EXPRESSION OF TIE-2 RECEPTORBODY FOR AFFINITY-BASED STUDY OF TIE-2 LIGAND INTERACTIONS

An expression construct was created that would yield a secreted protein consisting of the entire extracellular portion of the rat TIE-2 receptor fused to the human immunoglobulin gamma-1 constant region (IgG1 Fc). This fusion protein is called a TIE-2 "receptorbody" (RB), and would be normally expected to exist as a dimer in solution based on

formation of disulfide linkages between individual IgG1 Fc tails. The Fc portion of the TIE-2 RB was prepared as follows. A DNA fragment encoding the Fc portion of human IgG1 that spans from the hinge region to the carboxy-terminus of the protein, was amplified from human placental cDNA by PCR with oligonucleotides corresponding to the published sequence of human IgG1; the resulting DNA fragment was cloned in a plasmid vector. Appropriate DNA restriction fragments from a plasmid encoding the full-length TIE-2 receptor and from the human IgG1 Fc plasmid were ligated on either side of a short PCR-derived fragment that was designed so as to fuse, in-frame, the TIE-2 and human IgG1 Fc protein-coding sequences. Thus, the resulting TIE-2 ectodomain-Fc fusion protein precisely substituted the IgG1 Fc in place of the region spanning the TIE-2 transmembrane and cytoplasmic domains. An alternative method of preparing RBs is described in Goodwin, et. al. Cell 73:447-456 (1993).

Milligram quantities of TIE-2 RB were obtained by cloning the TIE-2 RB DNA fragment into the pVL1393 baculovirus vector and subsequently infecting the Spodoptera frugiperda SF-21AE insect cell line. Alternatively, the cell line SF-9 (ATCC Accession No. CRL-1711) or the cell line BTI-TN-5b1-4 may be used. DNA encoding the TIE-2 RB was cloned as an Eco RI-NotI fragment into the baculovirus transfer plasmid pVL1393. Plasmid DNA purified by cesium chloride density gradient centrifugation was recombined into viral DNA by mixing 3 μg of plasmid DNA with 0.5 μg of Baculo-Gold DNA (Pharminigen), followed by introduction into liposomes using 30μg Lipofectin (GIBCO-BRL). DNA-liposome mixtures were added to SF-21AE cells (2x 10^6 cells/60mm dish) in TMN-FH medium (Modified Grace's Insect Cell Medium (GIBCO-BRL) for 5 hours at 27°C, followed by incubation at 27°C for 5 days in

TMN-FH medium supplemented with 5% fetal calf serum. Tissue culture medium was harvested for plaque purification of recombinant viruses, which was carried out using methods previously described (O'Reilly, D.R., L.K. Miller, and V.A. Luckow, Baculovirus Expression Vectors - A Laboratory Manual. 1992, New York: W.H. Freeman) except that the agarose overlay contained 125 µg/mL X-gal (5-bromo-4-chloro-3-indolyl-β-D-galactopyranoside; GIBCO-BRL). After 5 days of incubation at 27°C, non-recombinant plaques were scored by positive chromogenic reaction to the X-gal substrate, and their positions marked. Recombinant plaques were then visualized by addition of a second overlay containing 100 µg/mL MTT (3-[4,5-dimethylthiazol-2-yl]2,5,diphenyltetrazolium bromide; Sigma). Putative recombinant virus plaques were picked by plug aspiration, and purified by multiple rounds of plaque isolation to assure homogeneity. Virus stocks were generated by serial, low-multiplicity passage of plaque-purified virus. Low passage stocks of one virus clone (vTIE-2 receptor body) were produced.

SF-21AE cells were cultured in serum free medium (SF-900 II, Gibco BRL) containing 1X antibiotic/antimycotic solution (Gibco BRL) and 25 mg/L Gentamycin (Gibco BRL). Pluronic F-68 was added as a surfactant to a final concentration of 1g/L. Cultures (4L) were raised in a bioreactor (Artisan Cell Station System) for at least three days prior to infection. Cells were grown at 27°C, with gassing to 50 % dissolved oxygen, at a gas flow rate of 80 mL/min (aeration at a sparge ring). Agitation was by means of a marine impeller at a rate of 100 rpm. Cells were harvested in mid-logarithmic growth phase (~2 X10 ^{6} cells per mL), concentrated by centrifugation, and infected with 5 plaque forming units of vTIE-2 Receptor Body per cell. Cells and inoculum

were brought to 400mL with fresh medium, and virus was adsorbed for 2 hours at 27°C in a spinner flask. The culture was then resuspended in a final volume of 8L with fresh serum-free medium, and the cells incubated in the bioreactor using the previously described conditions.

Culture medium from vTIE-2 Receptor Body-infected SF21AE cells were collected by centrifugation (500x g, 10 minutes) at 72 hours post-infection. Cell supernatants were brought to pH 8 with NaOH. EDTA was added to a final concentration of 10 mM and the supernatant pH was readjusted to 8. Supernatants were filtered (0.45 µm, Millipore) and loaded on a protein A column (protein A sepharose 4 fast flow or HiTrap protein A, both from Pharmacia). The column was washed with PBS containing 0.5 M NaCl until the absorbance at 280 nm decreased to baseline. The column was washed in PBS and eluted with 0.5 M acetic acid. Column fractions were immediately neutralized by eluting into tubes containing 1 M Tris pH 9. The peak fractions containing the TIE-2 RB were pooled and dialyzed versus PBS.

EXAMPLE 3 - DEMONSTRATION THAT TIE-2 HAS A CRITICAL ROLE IN DEVELOPMENT OF THE VASCULATURE

Given the absence of a known ligand for TIE-2 receptor, it was reasoned that it might be possible to gain insight into the function of TIE-2 by introduction of "excess" soluble TIE-2 receptor body (TIE-2 RB) into a developing system. The potential ability of TIE-2 RB to bind, and thereby neutralize, available TIE-2 ligand could result in an observable disruption of normal vascular development and characterization of the ligand. To examine whether TIE-2 RB could be used to disrupt vascular development in early chick embryos, small pieces of a biologically

resorbable foam were soaked with TIE-2 RB and inserted immediately beneath the chorioallantoic membrane at positions just lateral to the primitive embryo.

Early chicken embryos develop atop the yolk from a small disk of cells that is covered by the chorioallantoic membrane (CAM). The endothelial cells that will come to line the vasculature in the embryo arise from both extra- and intra-embryonic cell sources. Extraembryonically-derived endothelial cells, which provide the major source for endothelial cells in the embryo, originate from accretions of mesenchyme that are situated laterally around the embryo-proper, just underneath the CAM. As these mesenchyme cells mature, they give rise to a common progenitor of both the endothelial and hematopoietic cell lineages, termed the hemangioblast. In turn, the hemangioblast gives rise to a mixed population of angioblasts (the endothelial cell progenitor) and hematoblasts (the pluripotential hematopoietic precursor). Formation of rudiments of the circulatory system begins when endothelial cell progeny segregate to form a one-cell-thick vesicle that surrounds the primitive blood cells. Proliferation and migration of these cellular components eventually produces a vast network of blood-filled microvessels under the CAM that will ultimately invade the embryo to join with limited, intraembryonically-derived vascular elements.

Newly fertilized chicken eggs obtained from Spafas, Inc. (Boston, MA) were incubated at 99.5°F, 55 % RH. At about 24 hrs. of development, the egg shell was wiped down with 70% ethanol and a dentist's drill was used to make a 1.5 cm. hole in the blunt apex of each egg. The shell membrane was removed to reveal an air space directly above the

embryo. Small rectangular pieces of sterile Gelfoam (Upjohn) were cut with a scalpel and soaked in equal concentrations of either TIE-2- or EHK-1 receptorbody. EHK-1 receptorbody was made as set forth in Example 2 using the EHK-1 extracellular domain instead of the TIE-2 extracellular domain (Maisonpierre et al., Oncogene 8:3277-3288 (1993). Each Gelfoam piece absorbed approximately 6 μg of protein in 30 μl. Sterile watchmakers forceps were used to make a small tear in the CAM at a position several millimeters lateral to the primitive embryo. The majority of the piece of RB-soaked Gelfoam was inserted under the CAM and the egg shell was sealed over with a piece of adhesive tape. Other similarly-staged eggs were treated in parallel with RB of the unrelated, neuronally expressed receptor tyrosine kinase, EHK-1 (Maisonpierre et al., Oncogene 8:3277-3288 (1993). Development was allowed to proceed for 4 days and then the embryos were examined by visual inspection. Embryos were removed by carefully breaking the shells in dishes of warmed PBS and carefully cutting away the embryo with surrounding CAM. Of 12 eggs treated with each RB, 6 TIE-2 RB and 5 EHK-1 RB treated embryos had developed beyond the stage observed at the start of the experiment. A dramatic difference was seen between these developed embryos, as shown in Figures 1A and 1B. Those treated with EHK-1 RB appeared to have developed relatively normally. Four out of five EHK-1 embryos were viable as judged by the presence of a beating heart. Furthermore, the extra-embryonic vasculature, which is visually obvious due to the presence of red blood cells, was profuse and extended several centimeters laterally under the CAM. By contrast, those treated with TIE-2 RB were severely stunted, ranging from 2-5 mm. in diameter, as compared with more than 10 mm. in diameter for the EHK-1 RB embryos. All of the TIE-2 RB treated embryos were dead and their

CAMs were devoid of blood vessels. The ability of TIE-2 RB to block vascular development in the chicken demonstrates that TIE-2 ligand is necessary for development of the vasculature.

EXAMPLE 4 - IDENTIFICATION OF A TIE-2-SPECIFIC BINDING ACTIVITY IN CONDITIONED MEDIUM FROM THE ras ONCOGENE-TRANSFORMED C2C12 MOUSE MYOBLAST CELL LINE

Screening of ten-fold-concentrated cell-conditioned media (10X CCM) from various cell lines for the presence of soluble, TIE-2-specific binding activity (BIAcore; Pharmacia Biosensor, Piscataway, NJ) revealed binding activity in serum-free medium from oncogenic-ras-transformed C2C12 cells (C2C12-ras), RAT 2-ras (which is a ras transformed fibroblast cell line), human glioblastoma T98G and the human neuroblastoma cell line known as SHEP-1.

The C2C12-ras 10X CCM originated from a stably-transfected line of C2C12 myoblasts that was oncogenically transformed by transfection with the T-24 mutant of H-ras by standard calcium phosphate-based methods. An SV40 based neomycin-resistance expression plasmid was physically linked with the ras expression plasmid in order to permit selection of transfected clones. Resulting G418-resistant ras-C2C12 cells were routinely maintained as a monolayer on plastic dishes in DMEM/glutamine/penicillin-streptomycin supplemented with 10 % fetal calf serum (FCS). Serum-free C2C12-ras 10X CCM was made by plating the cells at 60% confluence in a serum free defined media for 12 hours. (Zhan and Goldfarb, Mol. Cell. Biol. 6: 3541-3544 (1986)); Zhan, et al. Oncogene 1: 369-376 (1987)). The medium was discarded and replaced

with fresh DMEM/Q/P-S for 24 hours. This medium was harvested and cells were refed fresh DMEM/Q/P-S, which was also harvested after a further 24 hours. These CCM were supplemented with the protease inhibitors PMSF (1mM) and aprotinin (10µg/ml), and ten-fold concentrated on sterile size-exclusion membranes (Amicon). TIE-2-binding activity could be neutralized by incubation of the medium with an excess of TIE-2 RB, but not by incubation with EHK-1 RB, prior to BIAcore analysis.

Binding activity of the 10x CCM was measured using biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ) which monitors biomolecular interactions in real-time via surface plasmon resonance. Purified TIE-2 RB was covalently coupled through primary amines to the carboxymethyl dextran layer of a CM5 research grade sensor chip (Pharmacia Biosensor; Piscataway, NJ). The sensor chip surface was activated using a mixture of N-hydroxysuccinimide (NHS) and N-ethyl-N'-(3-dimethylaminopropyl)carbodiimide (EDC), followed by immobilization of TIE-2 RB (25 µg/mL, pH 4.5) and deactivation of unreacted sites with 1.0 M ethanolamine (pH 8.5). A negative control surface of the EHK-1 receptorbody was prepared in a similar manner.

The running buffer used in the system was HBS (10 mM Hepes, 3.4 mM edta, 150 mM NaCl, 0.005% P20 surfactant, pH 7.4). The 10x CCM samples were centrifuged for 15 min at 4 C and further clarified using a sterile, low protein-binding 0.45 µm filter (Millipore; Bedford, MA). Dextran (2mg/ml) and P20 surfactant (0.005%) were added to each CCM sample. Aliquots of 40 µL were injected across the immobilized surface (either TIE-2 or EHK-1) at a flow rate of 5 µL/min and the receptor binding was monitored for 8 min. The binding activity

(resonance units, RU) was measured as the difference between a baseline value determined 30 s prior to the sample injection and a measurement taken at 30 s post-injection. Regeneration of the surface was accomplished with one 12-μL pulse of 3 M $MgCl_2$.

The instrument noise level is 20 RU; therefore, any binding activity with a signal above 20 RU may be interpreted as a real interaction with the receptor. For C2C12-ras conditioned media, the binding activities were in the range 60-90 RU for the TIE-2 RB immobilized surface. For the same samples assayed on a EHK-1 RB immobilized surface, the measured activities were less than 35 RU. Specific binding to the TIE-2 receptorbody was evaluated by incubating the samples with an excess of either soluble TIE-2 or EHK-1 RB prior to assaying the binding activity. The addition of soluble EHK-1 RB had no effect on the TIE-2 binding activity of any of the samples, while in the presence of soluble TIE-2 binding to the surface is two-thirds less than that measured in the absence of TIE-2. A repeat assay using >50x concentrated C2C12-ras CCM resulted in a four-fold enhancement over background of the TIE-2 specific binding signal.

EXAMPLE 5 - C2C12-ras CCM CONTAINS AN ACTIVITY THAT INDUCES TYROSINE PHOSPHORYLATION OF TIE-2 RECEPTOR

C2C12-ras 10X CCM was examined for its ability to induce tyrosine phosphorylation of TIE-2 in ABAE cells. Serum-starved ABAE cells were briefly incubated with C2C12-ras CCM, lysed and subjected to immunoprecipitation and Western analyses as described above. Stimulation of serum-starved ABAE cells with serum-free C2C12-ras

10X CCM was done as follows. The medium of ABAE cells starved as described above was removed and replaced with either defined medium or 10X CCM that had been pre-warmed to 37°C. After 10 minutes, the media were removed and the cells were twice rinsed on ice with an excess of chilled PBS supplemented with orthovanadate/NaF/benzamidine. Cell lysis and TIE-2-specific immunoprecipitation was done as described above.

ABAE cells incubated for 10 minutes with defined medium showed no induction of TIE-2 tyrosine phosphorylation, whereas incubation with C2C12-ras CCM stimulated at least a 100 X increase in TIE-2 phosphorylation. This activity was almost totally depleted by pre-incubation of the C2C12-ras 10X CCM for 90 minutes at room temperature with 13 ug of TIE-2 RB coupled to protein G-Sepharose beads. Medium incubated with protein G Sepharose alone was not depleted of this phosphorylating activity.

EXAMPLE 6 - EXPRESSION CLONING OF TIE-2 LIGAND

COS-7 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% fetal bovine serum (FBS), 1% each of penicillin and streptomycin (P/S) and 2 mM glutamine in an atmosphere of 5% CO_2. The mouse myoblast C2C12 ras cell line was cultured in Eagle's minimal essential medium (EMEM) with 10% FBS, (P/S) and 2 mM glutamine. Full length mouse TIE-2 ligand cDNA clones were obtained by screening a C2C12 ras cDNA library in the pJFE14 vector expressed in COS cells. This vector, as shown in Figure 2, is a modified version of the vector pSR_α (Takebe, et al. 1988, Mol. Cell. Biol. 8:466-472). The

library was created using the two BSTX1 restriction sites in the pJFE14 vector.

COS-7 cells were transiently transfected with either the pJFE14 library or control vector by the DEAE-dextran transfection protocol. Briefly, COS-7 cells were plated at a density of 1.0 x 10^6 cells per 100 mm plate 24 hours prior to transfection. For transfection, the cells were cultured in serum-free DMEM containing 400 µg/ml of DEAE-dextran, 1 µM chloroquine, and 2 mM glutamine, and 1 µg of the appropriate DNA for 3-4 hours at 37°C in an atmosphere of 5% CO_2. The transfection media was aspirated and replaced with phosphate-buffered saline with 10% DMSO for 2-3 min. Following this DMSO "shock", the COS-7 cells were placed into DMEM with 10% FBS, 1% each of penicillin and streptomycin, and 2 mM glutamine for 48 hours.

Because the TIE-2 ligand is secreted it was necessary to permeabilize the cells to detect binding of the receptorbody probe to the ligand. Two days after transfection the cells were rinsed with PBS and then incubated with PBS containing 1.8% formaldehyde for 15-30 min. at room temperature. Cells were then washed with PBS and incubated for 15 min. with PBS containing 0.1% Triton X-100 and 10% Bovine Calf Serum to permeabilize the cells and block non-specific binding sites. The screening was conducted by direct localization of staining using a TIE-2 receptorbody, which consisted of the extracellular domain of TIE-2 fused to the IgG1 constant region. This receptorbody was prepared as set forth in Example 2. A 100 mm dish of transfected, fixed and permeabilized COS cells was probed by incubating them for 30 min with TIE-2-RB. The cells were then washed twice with PBS and

incubated for an additional 30 min with PBS/10% Bovine Calf Serum/anti-human IgG-alkaline phosphatase conjugate. After three PBS washes, cells were incubated in alkaline-phosphatase substrate for 30-60 min. The dish was then inspected microscopically for the presence of stained cells. For each stained cell, a small area of cells including the stained cell was scraped from the dish using a plastic pipette tip and plasmid DNA was then rescued and used to electroporate bacterial cells. Single bacterial colonies resulting from the electroporation were picked and plasmid DNA prepared from these colonies was used to transfect COS-7 cells which were probed for TIE-2 ligand expression as evidenced by binding to TIE-2 receptorbodies. This allowed identification of single clones coding for TIE-2 ligand. Confirmation of TIE-2 ligand expression was obtained by phosphorylation of the TIE-2 receptor using the method set forth in Example 5. A plasmid clone encoding the TIE-2 ligand was deposited with the ATCC on October 7, 1994 and designated as "pJFE14 encoding TIE-2 ligand" under ATCC Accession No. 75910.

EXAMPLE 7 - ISOLATION AND SEQUENCING OF FULL LENGTH cDNA CLONE ENCODING HUMAN TIE-2 LIGAND

A human fetal lung cDNA library in lambda gt-10 (see Figure 3) was obtained from Clontech Laboratories, Inc. (Palo Alto, CA). Plaques were plated at a density of 1.25 x 10^6/20x20 cm plate, and replica filters taken following standard procedures (Sambrook, et al., Molecular Cloning: A Laboratory Manual, 2nd Ed., page 8.46, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York).

Isolation of human tie-2 ligand clones was carried out as follows. A 2.2 kb XhoI fragment from the deposited tie-2 ligand clone (ATCC NO. 75910 - see Example 6 above) was labeled by random priming to a specific activity of approximately 5x10^{8}cpm/ng. Hybridization was carried out at 65°C in hybridization solution containing 0.5 mg/ml salmon sperm DNA. The filters were washed at 65°C in 2 x SSC, 0.1 % SDS and exposed to Kodak XAR-5 film overnight at -70°C. Positive phage were plaque purified. High titre phage lysates of pure phage were used for isolation of DNA via a Qiagen column using standard techniques (Qiagen, Inc., Chatsworth, CA, 1995 catalog, page 36). Phage DNA was digested with EcoRI to release the cloned cDNA fragment for subsequent subcloning. A lambda phage vector containing human tie-2 ligand DNA was deposited with the ATCC on October 26, 1994 under the designation λgt10 encoding htie-2 ligand 1 (ATCC Accession No. 75928). Phage DNA may be subjected directly to DNA sequence analysis by the dideoxy chain termination method (Sanger, et al., 1977, Proc. Natl. Acad. Sci. U.S.A. 74: 5463-5467).

Subcloning human TIE-2 ligand into a mammalian expression vector

The clone λgt10 encoding htie-2 ligand 1 contains an EcoRI site located 490 base pairs downstream from the start of the coding sequence for the human tie-2 ligand. The coding region may be excised using unique restriction sites upstream and downstream of the initiator and stop codons respectively. For example, an SpeI site, located 70 bp 5' to the initiator codon, and a Bpu1102i (also known as BlpI) site, located 265 bp 3' to the stop codon, may be used to excise the complete coding region. This may then be subcloned into the pJFE14 cloning vector,

using the XbaI (compatible to the SpeI overhang) and the PstI sites (the PstI and Bpu1102i sites are both made blunt ended).

Sequencing of human TIE-2 ligand

The coding region from the clone λgt10 encoding htie-2 ligand 1 was sequenced using the ABI 373A DNA sequencer and Taq Dyedeoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). The nucleotide and deduced amino acid sequence of human TIE-2 ligand from the clone λgt10 encoding htie-2 ligand 1 is shown in Figure 4.

In addition, full length human TIE-2 ligand cDNA clones were obtained by screening a human glioblastoma T98G cDNA library in the pJFE14 vector. Clones encoding human tie-2 ligand were identified by DNA hybridization using a 2.2 kb XhoI fragment from the deposited tie-2 ligand clone (ATCC NO. 75910) as a probe (see Example 6 above). The coding region was sequenced using the ABI 373A DNA sequencer and Taq Dyedeoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). This sequence was nearly identical to that of clone λgt10 encoding htie-2 ligand 1. As shown in Figure 4, the clone λgt10 encoding htie-2 ligand 1 contains an additional glycine residue which is encoded by nucleotides 1114-1116. The coding sequence of the T98G clone does not contain this glycine residue but otherwise is identical to the coding sequence of the clone λgt10 encoding htie-2 ligand 1. Figure 5 sets forth the nucleotide and deduced amino acid sequence of human TIE-2 ligand from the T98G clone.

EXAMPLE 8 - ISOLATION AND SEQUENCING OF SECOND FULL LENGTH cDNA CLONE A ENCODING HUMAN TIE-2 LIGAND

A human fetal lung cDNA library in lambda gt-10 (see Figure 3) was obtained from Clontech Laboratories, Inc. (Palo Alto, CA). Plaques were plated at a density of 1.25 x 10^6/20x20 cm plate, and replica filters taken following standard procedures (Sambrook, et al., Molecular Cloning: A Laboratory Manual, 2nd Ed., page 8.46, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York). Duplicate filters were screened at low stringency (2 x SSC, 55° C) with probes made to the human tie 2 L-1 sequence. One of the duplicate filters was probed with a 5' probe, encoding amino acids 25 - 265 of human tie 2L-1 as set forth in Figure 4. The second duplicate filter was probed with a 3' probe, encoding amino acids 282 - 498 of human tie 2 L-1 sequence (see Figure 4). Both probes were hybridized at 55 C in hybridization solution containing 0.5 mg/ml salmon sperm DNA. Filters were washed in 2 x SSC at 55° C and exposed overnight to X-ray film. In addition, duplicate filters were also hybridized at normal stringency (2 x SSC, 65° C) to the full length coding probe of mouse tie2L (F3-15, XhoI insert). Three positive clones were picked that fulfilled the following criteria: i. hybridization had not been seen to the full length (mouse) probe at normal stringency, and ii. hybridization was seen at low stringency to both 5' and 3' probes. EcoRI digestion of phage DNA obtained from these clones indicated two independent clones with insert sizes of approximately 2.2kb and approximately 1.8 kb. The 2.2kb EcoRI insert was subcloned into the EcoRI sites of both pBluescript KS (Stratagene) and a mammalian expression vector suitable for use in COS cells. Two orientations were identified for the mammalian expression vector. The 2.2kb insert in pBluescript KS was

deposited with the ATCC on December 9, 1994 and designated as pBluescript KS encoding human TIE 2 ligand 2. The start site of the TIE-2 ligand 2 coding sequence is approximately 355 base pairs downstream of the pBluescript EcoRI site.

COS-7 cells were transiently transfected with either the expression vector or control vector by the DEAE-dextran transfection protocol. Briefly, COS-7 cells were plated at a density of 1.0 x 10^6 cells per 100 mm plate 24 hours prior to transfection. For transfection, the cells were cultured in serum-free DMEM containing 400 µg/ml of DEAE-dextran, 1 µM chloroquine, and 2 mM glutamine, and 1 µg of the appropriate DNA for 3-4 hours at 37°C in an atmosphere of 5% CO_2. The transfection media was aspirated and replaced with phosphate-buffered saline with 10% DMSO for 2-3 min. Following this DMSO "shock", the COS-7 cells were placed into DMEM with 10% FBS, 1% each of penicillin and streptomycin, and 2 mM glutamine for 48 hours.

Because the TIE-2 ligand is secreted it was necessary to permeabilize the cells to detect binding of the receptorbody probe to the ligand. Transfected COS-7 cells were plated at a density of 1.0 x 10^6 cells per 100 mm plate. The cells were rinsed with PBS and then incubated with PBS containing 1.8% formaldehyde for 15-30 min. at room temperature. Cells were then washed with PBS and incubated for 15 min. with PBS containing 0.1% Triton X-100 and 10% Bovine Calf Serum to permeabilize the cells and block non-specific binding sites.
The screening was conducted by direct localization of staining using a TIE-2 receptorbody, which consisted of the extracellular domain of TIE-2 fused to the IgG1 constant region. This receptorbody was

prepared as set forth in Example 2. Transfected COS cells were probed by incubating them for 30 min with TIE-2-RB. The cells were then washed twice with PBS, fixed with methanol, and then incubated for an additional 30 min with PBS/10% Bovine Calf Serum/anti-human IgG-alkaline phosphatase conjugate. After three PBS washes, cells were incubated in alkaline-phosphatase substrate for 30-60 min. The dish was then inspected microscopically for the presence of stained cells. Cells expressing one orientation of the clone, but not the other orientation, were seen to bind the TIE-2 receptor body.

One of skill in the art will readily see that the described methods may be used to further identify other related members of the TIE ligand family.

Sequencing of second human TIE-2 ligand

The coding region from the clone pBluescript KS encoding human TIE-2 ligand 2 was sequenced using the ABI 373A DNA sequencer and Taq Dyedeoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). The nucleotide and deduced amino acid sequence of human TIE-2 ligand from the clone pBluescript KS encoding human TIE-2 ligand 2 is shown in Figure 6.

EXAMPLE 9 - TIE-2 LIGAND 2 IS A RECEPTOR ANTAGONIST

Conditioned media from COS cells expressing either TIE-2 ligand 2 (TL2) or TIE-2 ligand 1 (TL1) were compared for their ability to activate TIE-2 receptors naturally present in a human endothelial cell line.

Lipofectamine reagent (GIBCO-BRL, Inc.) and recommended protocols were used to transfect COS-7 cells with either the pJFE14 expression vector alone, pJFE14 vector containing the human TIE-2 ligand 1 cDNA, or with a pMT21 expression vector (Kaufman, R.J., 1985, Proc. Natl. Acad. Sci. USA 82: 689-693) containing the human TIE-2 ligand 2 cDNA. COS media containing secreted ligands were harvested after three days and concentrated 20-fold by diafiltration (DIAFLO ultrafiltration membranes, Amicon, Inc.). The quantity of active TIE-2 ligand 1 and TIE-2 ligand 2 present in these media was determined and expressed as the amount (in resonance units, R.U.) of TIE-2 receptor specific binding activity measured by a BIAcore binding assay.

Northern (RNA) analyses revealed significant levels of TIE-2 transcripts in HAEC (Human Aortic Endothelial Cell) human primary endothelial cells (Clonetics, Inc.). Therefore, these cells were used to examine whether TIE-2 receptor is tyrosine-phosphorylated when exposed to COS media containing the TIE-2 ligands. HAEC cells were maintained in a complete endothelial cell growth medium (Clonetics, Inc.) that contained 5% fetal bovine serum, soluble bovine brain extract, 10 ng/ml human EGF, 1 mg/ml hydrocortisone, 50 mg/ml gentamicin and 50 ng/ml amphotericin-B. Assessment of whether TL1 and TL2 could activate TIE-2 receptor in the HAEC cells was done as follows. Semi-confluent HAEC cells were serum-starved for two hours in high-glucose Dulbecco's MEM with added L-glutamine and penicillin-streptomycin at 37°C followed by replacement of the starvation medium with ligand-containing conditioned COS media for 7 minutes at 37°C in a 5% CO2 incubator. The cells were subsequently lysed and TIE-2 receptor protein was recovered by immunoprecipitation of the

lysates with TIE-2 peptide antiserum, followed by Western blotting with antiphosphotyrosine antiserum, exactly as described in example 1. The results are shown in Figure 7. Phosphotyrosine levels on the TIE-2 receptor (TIE2-R) were induced by treatment of HEAC cells with TIE-2 ligand 1 (Lane L1) but not by TIE-2 ligand 2 (Lane L2) conditioned COS media. MOCK is conditioned media from COS transfected with JFE14 empty vector.

Evidence that both TL1 and TL2 specifically bind to the TIE-2 receptor was demonstrated by using a BIAcore to assay the TIE-2 receptor specific binding activities in transfected COS media and by immunostaining of TL1- and TL2-expressing COS cells with TIE-2 receptorbodies.

Because TL2 did not activate the TIE-2 receptor, applicants set out to determine whether TL2 might be capable of serving as an antagonist of TL1 activity. HAEC phosphorylation assays were performed in which cells were first incubated with an "excess" of TL2, followed by addition of dilute TL1. It was reasoned that prior occupancy of TIE-2 receptor due to high levels of TL2 might prevent subsequent stimulation of the receptor following exposure to TL1 present at a limiting concentration.

Semi-confluent HAEC cells were serum-starved as described above and then incubated for 3 min., at 37°C with 1-2 ml. of 20X COS/JFE14-TL2 conditioned medium. Control plates were treated with 20X COS/JFE14-only medium (MOCK). The plates were removed from the incubator and various dilutions of COS/JFE14-TL1 medium were then added, followed by further incubation of the plates for 5-7 min. at 37°C. Cells were

subsequently rinsed, lysed and TIE-2-specific tyrosine phosphorylation in the lysates was examined by receptor immunoprecipitation and Western blotting, as described above. TL1 dilutions were made using 20X COS/JFE14-TL1 medium diluted to 2X, 0.5X, 0.1X, or 0.02X by addition of 20X COS/JFE14-alone medium. An assay of the initial 20X TL1 and 20X TL2 COS media using BIAcore biosensor technology indicated that they contained similar amounts of TIE-2-specific binding activities, i.e., 445 R.U. and 511 R.U. for TL1 and TL2, respectively. The results of the antiphosphotyrosine Western blot, shown in Figure 8, indicate that when compared to prior treatment of HAEC cells with MOCK medium (lane 1), prior treatment of HAEC cells with excess TIE-2 ligand 2 (lane 2) antagonizes the subsequent ability of dilute TIE-2 ligand 1 to activate the TIE-2 receptor (TIE2-R).

These data indicate that, unlike TL1, TL2 was not able to stimulate TIE-2 receptor kinase activity in HAEC cells. Furthermore, pre-incubation of the endothelial cells with high concentrations of TL2 followed by addition of TL1 blocked the ability of TL1 to stimulate the TIE-2 receptor, indicating that TL2 is a TIE-2 receptor antagonist.

EXAMPLE 10 - IDENTIFICATION OF TIE-2-SPECIFIC BINDING ACTIVITY IN CONDITIONED MEDIUM AND COS CELL SUPERNATANTS

Binding activity of 10x CCM from the cell lines C2C12 ras, Rat2 ras, SHEP, and T98G, or cos cell supernatants after transfection with either human TIE-2 ligand 1 (hTL1) or human TIE-2 ligand 2 (hTL2) was measured using biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ) which monitors biomolecular interactions in real-time

via surface plasmon resonance (SPR). Purified rat or human TIE-2 RB was covalently coupled through primary amines to the carboxymethyl dextran layer of a CM5 research grade sensor chip (Pharmacia Biosensor; Piscataway, NJ). The sensor chip surface was activated using a mixture of N-hydroxysuccinimide (NHS) and N-ethyl-N'-(3-dimethylaminopropyl)carbodiimide (EDC), followed by immobilization of TIE-2 RB (25 μg/mL, pH 4.5) and deactivation of unreacted sites with 1.0 M ethanolamine (pH 8.5). In general, 9000-10000 RU of each receptorbody was coupled to the sensor chip.

The running buffer used in the system was HBS (10 mM Hepes, 150 mM NaCl, 0.005% P20 surfactant, pH 7.4). The samples were centrifuged for 15 min at 4°C and further clarified using a sterile, low protein-binding 0.45 μm filter (Millipore; Bedford, MA). Dextran (2mg/ml) and P20 surfactant (0.005%) were added to each sample. Aliquots of 40 μL were injected across the immobilized surface (either rat or human TIE2) at a flow rate of 5 μL/min and the receptor binding was monitored for 8 min. The binding activity (resonance units, RU) was measured as the difference between a baseline value determined 30 s prior to the sample injection and a measurement taken at 30 s post-injection. Regeneration of the surface was accomplished with one 15-μL pulse of 3 M $MgCl_2$.

The CCM samples (C2C12 ras, Rat2 ras, SHEP, T98G) were tested on the rat TIE2 RB immobilized surface, while the recombinant hTL1 and hTL2 were tested on the human TIE2 RB immobilized surface. In each case, specific binding to the TIE-2 receptorbody was evaluated by incubating the samples with 25 μg/ml of either soluble TIE-2 (rat or human) RB or

trkB RB prior to assaying the binding activity. As shown in Figures 9 and 10, the addition of soluble trkB RB causes a slight decrease in the TIE-2 binding activity, while the addition of soluble TIE-2 RB significantly reduces the binding activity as compared to that measured in the absence of TIE-2 RB.

EXAMPLE 11 - TIE-2 RB SPECIFICALLY BLOCKS ACTIVATION OF THE TIE-2 RECEPTOR BY TIE-2 LIGAND 1

The applicants sought to determine whether soluble TIE-2 RB can serve as a competitive inhibitor to block activation of TIE-2 receptor by TIE-2 ligand 1 (TL1). To do this, TL1-containing COS media were preincubated with either TIE2- or TrkB-RB and then compared for their ability to activate TIE-2 receptors naturally present in a human endothelial cell line.

Conditioned COS media were generated from COS-7 cells transfected with either the pJFE14 expression vector alone (MOCK), or pJFE14 vector containing the human TIE-2 ligand 1 cDNA (TL1) and harvested as described in Example 9 hereinabove, with the exception that the media were sterile filtered but not concentrated. The quantity of TL1 was determined and expressed as the amount (in resonance units, R.U.) of TIE-2 receptor- specific binding activity measured by BIAcore binding assay.

Northern (RNA) analyses revealed significant levels of TIE-2 transcripts in HUVEC (Human Umbilical Vein Endothelial Cell) human primary endothelial cells (Clonetics, Inc.). Therefore, these cells were

used to examine whether TIE-2 receptor can be tyrosine-phosphorylated when exposed in the presence of TIE2- or TrkB-RBs to COS media containing TL1. HUVEC cells were maintained at 37°C, 5% CO2 in a complete endothelial cell growth medium (Clonetics, Inc.) that contained 5% fetal bovine serum, soluble bovine brain extract with 10 ug/ml heparin, 10 ng/ml human EGF, 1 ug/ml hydrocortisone, 50 ug/ml gentamicin and 50 ng/ml amphotericin-B. Assessment of whether TL1 could activate TIE-2 receptor in the HUVEC cells was done as follows. Confluent dishes of HUVEC cells were serum-starved for two-to-four hours in low-glucose Dulbecco's MEM at 37°C, 5% CO2, followed by 10 minute incubation in starvation medium that included 0.1 mM sodium orthovanadate, a potent inhibitor of phosphotyrosine phosphatases. Meanwhile, conditioned COS media were preincubated 30 min. at room temperature with either TIE2- or TrkB-RB added to 50 ug/ml. The starvation medium was then removed from the HUVEC dishes and incubated with the RB-containing COS media for 7 minutes at 37°C. HUVEC cells were subsequently lysed and TIE-2 receptor protein was recovered by immunoprecipitation with TIE-2 peptide antiserum, followed by Western blotting with an anti-phosphotyrosine antibody, as described in Example 1. The results are shown in Figure 11. Phosphotyrosine levels on the TIE-2 receptor were induced by treatment of HUVEC cells with TIE-2 ligand 1 (TL1) relative to that seen with control medium (MOCK) and this induction is specifically blocked by prior incubation with TIE2-RB (TIE2-Fc) but not by incubation with TrkB-RB (TrkB-Fc). These data indicate that soluble TIE-2 RB can serve as a selective inhibitor to block activation of TIE-2 receptor by TIE-2 ligand 1.

EXAMPLE 12 - CONSTRUCTION OF TIE-2 LIGANDBODIES

An expression construct was created that would yield a secreted protein consisting of the entire coding sequence of human TIE-2 ligand 1 (TL1) or TIE-2 ligand 2 (TL2) fused to the human immunoglobulin gamma-1 constant region (IgG1 Fc). These fusion proteins are called TIE-2 "ligandbodies" (TL1-Fc or TL2-Fc). The Fc portion of TL1-Fc and TL2-Fc was prepared as follows. A DNA fragment encoding the Fc portion of human IgG1 that spans from the hinge region to the carboxy-terminus of the protein, was amplified from human placental cDNA by PCR with oligonucleotides corresponding to the published sequence of human IgG1; the resulting DNA fragment was cloned in a plasmid vector. Appropriate DNA restriction fragments from a plasmid encoding full-length TL1 or TL2 and from the human IgG1 Fc plasmid were ligated on either side of a short PCR-derived fragment that was designed so as to fuse, in-frame, TL1 or TL2 with human IgG1 Fc protein-coding sequences.

Milligram quantities of TL2-Fc were obtained by cloning the TL2-Fc DNA fragment into the pVL1393 baculovirus vector and subsequently infecting the Spodoptera frugiperda SF-21AE insect cell line. Alternatively, the cell line SF-9 (ATCC Accession No. CRL-1711) or the cell line BTI-TN-5b1-4 may be used. DNA encoding the TL2-Fc was cloned as an Eco RI-NotI fragment into the baculovirus transfer plasmid pVL1393. Plasmid DNA was recombined into viral DNA by mixing 3 mg of plasmid DNA with 0.5 mg of Baculo-Gold DNA (Pharminigen), followed by introduction into liposomes using 30mg Lipofectin (GIBCO-BRL). DNA-liposome mixtures were added to SF-21AE cells (2x 106 cells/60mm dish) in TMN-FH medium (Modified Grace's Insect Cell

Medium (GIBCO-BRL) for 5 hours at 27°C, followed by incubation at 27°C for 5 days in TMN-FH medium supplemented with 5% fetal calf serum. Tissue culture medium was harvested for plaque purification of recombinant viruses, which was carried out using methods previously described (O'Reilly, D.R., L.K. Miller, and V.A. Luckow, Baculovirus Expression Vectors - A Laboratory Manual. 1992, New York: W.H. Freeman) except that the agarose overlay contained 125 mg/mL X-gal (5-bromo-4-chloro-3-indolyl-b- D-galactopyranoside; GIBCO-BRL). After 5 days of incubation at 27°C, non-recombinant plaques were scored by positive chromogenic reaction to the X-gal substrate, and their positions marked. Recombinant plaques were then visualized by addition of a second overlay containing 100 mg/mL MTT (3-[4,5-dimethylthiazol-2-yl]2,5,diphenyltetrazolium bromide; Sigma). Putative recombinant virus plaques were picked by plug aspiration, and purified by multiple rounds of plaque isolation to assure homogeneity. Virus stocks were generated by serial, low-multiplicity passage of plaque-purified virus. Low passage stocks of one virus clone (vTL2-Fc Clone #7) were produced.

SF-21AE cells were cultured in serum free medium (SF-900 II, Gibco BRL) containing 1X antibiotic/antimycotic solution (Gibco BRL) and 25 mg/L Gentamycin (Gibco BRL). Pluronic F-68 was added as a surfactant to a final concentration of 1g/L. Cultures (4L) were raised in a bioreactor (Artisan Cell Station System) for at least three days prior to infection. Cells were grown at 27°C, with gassing to 50 % dissolved oxygen, at a gas flow rate of 80 mL/min (aeration at a sparge ring). Agitation was by means of a marine impeller at a rate of 100 rpm. Cells were harvested in mid-logarithmic growth phase (~2 X10 6 cells per mL), concentrated by centrifugation, and infected with 5 plaque

forming units of vTL2-Fc per cell. Cells and inoculum were brought to 400mL with fresh medium, and virus was adsorbed for 2 hours at 27°C in a spinner flask. The culture was then resuspended in a final volume of 8L with fresh serum-free medium, and the cells incubated in the bioreactor using the previously described conditions.

Culture medium from vTL2-Fc-infected SF21AE cells were collected by centrifugation (500x g, 10 minutes) at 72 hours post-infection. Cell supernatants were brought to pH 8 with NaOH. EDTA was added to a final concentration of 10 mM and the supernatant pH was readjusted to 8. Supernatants were filtered (0.45 mm, Millipore) and loaded on a protein A column (protein A sepharose 4 fast flow or HiTrap protein A, both from Pharmacia). The column was washed with PBS containing 0.5 M NaCl until the absorbance at 280 nm decreased to baseline. The column was washed in PBS and eluted with 0.5 M acetic acid. Column fractions were immediately neutralized by eluting into tubes containing 1 M Tris pH 9. The peak fractions containing the TL2-Fc were pooled and dialyzed versus PBS.

DEPOSITS

The following have been deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852 in accordance with the Budapest Treaty. A plasmid clone encoding a TIE-2 ligand was deposited with the ATCC on October 7, 1994 and designated as "pJFE14 encoding TIE-2 ligand" under ATCC Accession No. 75910. Recombinant Autographa californica baculovirus encoding TIE-2 receptor body was deposited with the ATCC on October 7, 1994 and designated as "vTIE-2 receptor body" under ATCC Accession No. VR2484. A lambda phage vector containing human tie-2 ligand DNA was deposited with the ATCC on October 26, 1994 and designated as λgt10 encoding htie-2 ligand 1 under ATCC Accession No. 75928. A plasmid clone encoding a second TIE-2 ligand was deposited with the ATCC on December 9, 1994 and designated as "pBluescript KS encoding human TIE 2 ligand 2" under ATCC Accession No. 75963.

The present invention is not to be limited in scope by the specific embodiments described herein. Indeed, various modifications of the invention in addition to those described herein will become apparent to those skilled in the art from the foregoing description and accompanying figures. Such modifications are intended to fall within the scope of the appended claims.

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WHAT IS CLAIMED IS:

1. An isolated nucleic acid molecule encoding a TIE-2 ligand.

2. An isolated nucleic acid molecule according to claim 1 wherein the nucleic acid sequence is
   (a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4, Figure 5 or Figure 6;
   (b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or
   (c) a nucleic acid sequence which, but for the degeneracy of the genetic code would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

3. A vector which comprises a nucleic acid molecule of claim 1 or 2.

4. A vector according to claim 3 wherein the nucleic acid molecule of claim 1 or 2 is operatively linked to an expression control sequence capable of directing its expression in a host cell.

5. A vector according to claim 3 or 4 which is a plasmid.

6. A plasmid according to claim 5 designated pJFE14 encoding TIE-2 ligand (ATCC Accession No. 75910).

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7. A plasmid according to claim 5 designated pBluescript KS encoding human TIE-2 ligand 2 (ATCC Accession No. 75963).

8. A vector according to claim 3 or 4 designated as λgtl0 encoding hTIE-2 ligand 1 (ATCC Accession No. 75928.)

9. An isolated TIE-2 ligand substantially free of other proteins.

10. Isolated TIE-2 ligand according to claim 9 encoded by a nucleic acid molecule according to claim 1 or 2.

11. A host-vector system for the production of a ligand according to claim 9 or 10 which comprises a vector according to any one of claims 3 to 8 in a host cell.

12. A host-vector system according to claim 11 wherein the host cell is a bacterial, yeast, insect or mammalian cell.

13. A host vector system comprising the host vector system of claim 11 or 12 and a nucleic acid encoding the TIE-2 receptor.

14. A method of producing a ligand as defined in claim 9 or 10 which comprises growing cells of a host-vector system according to any one of claims 10 to 12 under conditions permitting production of the ligand, and recovering the ligand so produced.

15. An antibody which specifically binds the ligand of claim 9 or 10.

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16. An antibody according to claim 15 which is a monoclonal antibody.

17. A receptorbody which specifically binds the ligand of claim 9 or 10.

18. An isolated nucleic acid molecule encoding a receptorbody according to claim 17.

19. A vector comprising a nucleic acid molecule according to claim 18.

20. A vector according to claim 19 which is a plasmid.

21. A plasmid according to claim 20 designated vTIE-2 receptorbody (ATCC Deposit VR2484).

22. A conjugate comprising a ligand according to claim 9 or 10 and, conjugated thereto, a cytotoxic agent.

23. A conjugate according to claim 22 wherein the cytotoxic agent is a radioisotope or toxin.

24. A pharmaceutical composition comprising a TIE-2 ligand according to claim 9 or 10 and a pharmaceutically acceptable carrier.

25. A pharmaceutical composition comprising an antibody according to claim 15 or 16 and a pharmaceutically acceptable carrier.

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26. A pharmaceutical composition comprising a receptorbody according to claim 17 and a pharmaceutically acceptable carrier.

27. A pharmaceutical composition comprising a conjugate according to claim 22 or 23 and a pharmaceutically acceptable carrier.

28. A ligand according to claim 9 or 10, an antibody according to claim 15, a receptorbody according to claim 17, a conjugate according to claim 22 or 23, or a composition according to any one of claims 24 to 25 for use in a method of treatment of the human or animal body, or in a method of diagnosis.

29. An antibody or receptorbody according to claim 28 for use in a method of blocking blood vessel growth in a mammal.

30. An antibody or receptorbody according to claim 29 for use in a method wherein the mammal is a human.

31. A ligand according to claim 28 for use in method of promoting neovascularization in a mammal.

32. A ligand according to claim 31 for use in the promotion of wound healing.

33. A ligand according to claim 31 for use in the treatment of ischemia.

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34. A TIE-2 antagonist for use in a method of inhibiting TIE-2 ligand activity in a mammal.

35. An antagonist according to claim 34 which is an antibody capable of specifically binding TIE-2 receptor.

36. An antibody according to claim 35 which is an antibody according to claim 15 or 16.

37. An antagonist according to claim 34 which is a receptorbody according to claim 17.

38. An antagonist according to claim 34 which is a ligand according to claim 9 or 10.

39. An antagonist according to any one of claims 34 to 38 for use in a method wherein the mammal is a human.

40. An antagonist according to any one of claims 34 to 39 for use in a method of attenuating or preventing tumour growth in a human.

41. A method of maintaining a TIE-2 receptor expressing cell in culture, which method comprises administering to the TIE-2 receptor expressing cell an effective amount of the ligand of claim 9 or 10.

42. A method according to claim 41 wherein the TIE-2 receptor expressing cell is an endothelial cell.

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43. A method of identifying a TIE-2 receptor antagonist comprising contacting cells expressing the TIE-2 receptor with:

a) a test compound; and

b) a ligand according to claim 9 or 10; under conditions permitting binding of the ligand to the receptor and determining whether the test compound is capable of interfering with the binding of the ligand to the receptor.

44. A polypeptide produced by the method of claim 14.

45. A nucleic acid according to claim 1 or 18, substantially as hereinbefore described with reference to any one of the foregoing Examples.

46. A vector according to claim 3, substantially as hereinbefore described with reference to any one of the foregoing Examples.

47. A ligand according to claim 9, 10, or 28, substantially as hereinbefore described with reference to any one of the foregoing Examples.

48. A host vector system according to claim 11, substantially as hereinbefore described with reference to any one of the foregoing Examples.

49. A method according to claim 14, 41, or 43, substantially as hereinbefore described with reference to any one of the foregoing Examples.

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* C L M S *

50. An antibody according to claim 15 or 28, substantially as hereinbefore described.

51. A receptorbody according to claim 17 or 28, substantially as hereinbefore described with reference to any one of the foregoing Examples.

52. A conjugate according to claim 22 or 28, substantially as hereinbefore described.

53. A composition according to any one of claims 24 to 27 or 28, substantially as hereinbefore described with reference to any one of the foregoing Examples.

54. A TIE-2 antagonist according to claim 34, substantially as hereinbefore described with reference to any one of the foregoing Examples.

55. A ligandbody which specifically binds the TIE-2 receptor or the receptorbody of claim 51.

56. A ligandbody which comprises a TIE-2 ligand fused to an immunoglobulin constant region.

57. The ligandbody of claim 56 wherein the TIE-2 ligand is TIE-2 ligand 1 or TIE-2 ligand 2 and the immunoglobulin constant region is the Fc portion of human IgG1.

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58. A ligandbody according to claims 55 to 57 for use in a method of treatment of the human or animal body, or in a method of diagnosis.

59. A method of treating a human or animal subject comprising administering to the subject an effective amount of a ligand according to claim 9 or 10, an antibody according to claim 15, a receptorbody according to claim 17, a conjugate according to claim 22 or 23, or a composition according to any one of claims 24 to 27.

60. A method according to claim 59, the method being as defined in any one of claims 29 to 40.

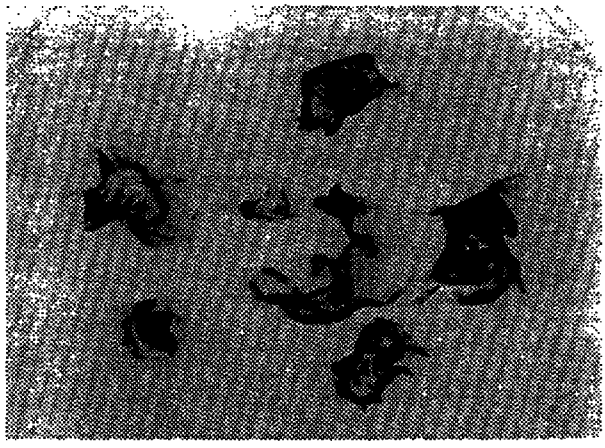
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Fig.1A.



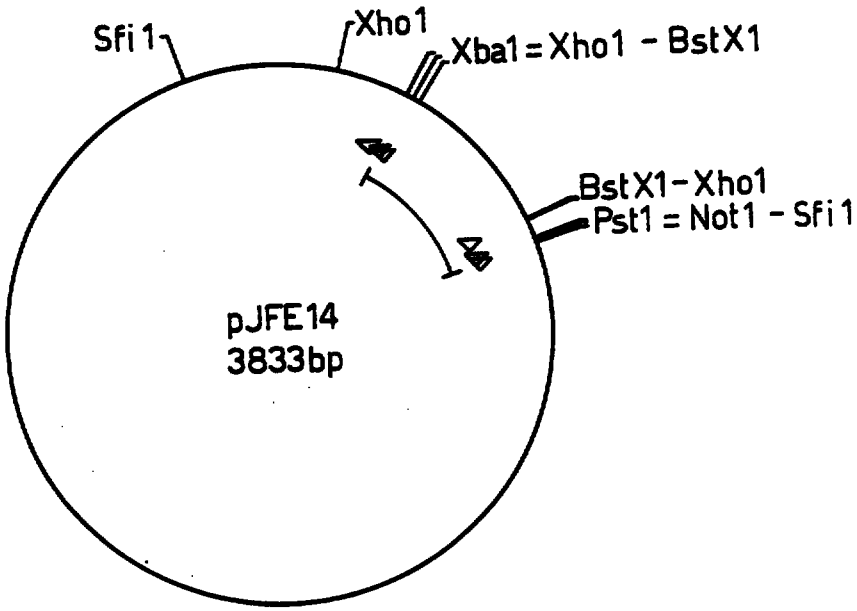
r EHK-1 ecto/h IgG1 Fc
Gelfoam (6ug)

Fig.1B.



r TIE-2 ecto/h IgG1 Fc
Gelfoam (6ug)

Fig.2.


Sfi 1
Xho1
Xba1 = Xho1 - BstX1
BstX1 - Xho1
Pst1 = Not1 - Sfi 1
pJFE14
3833bp

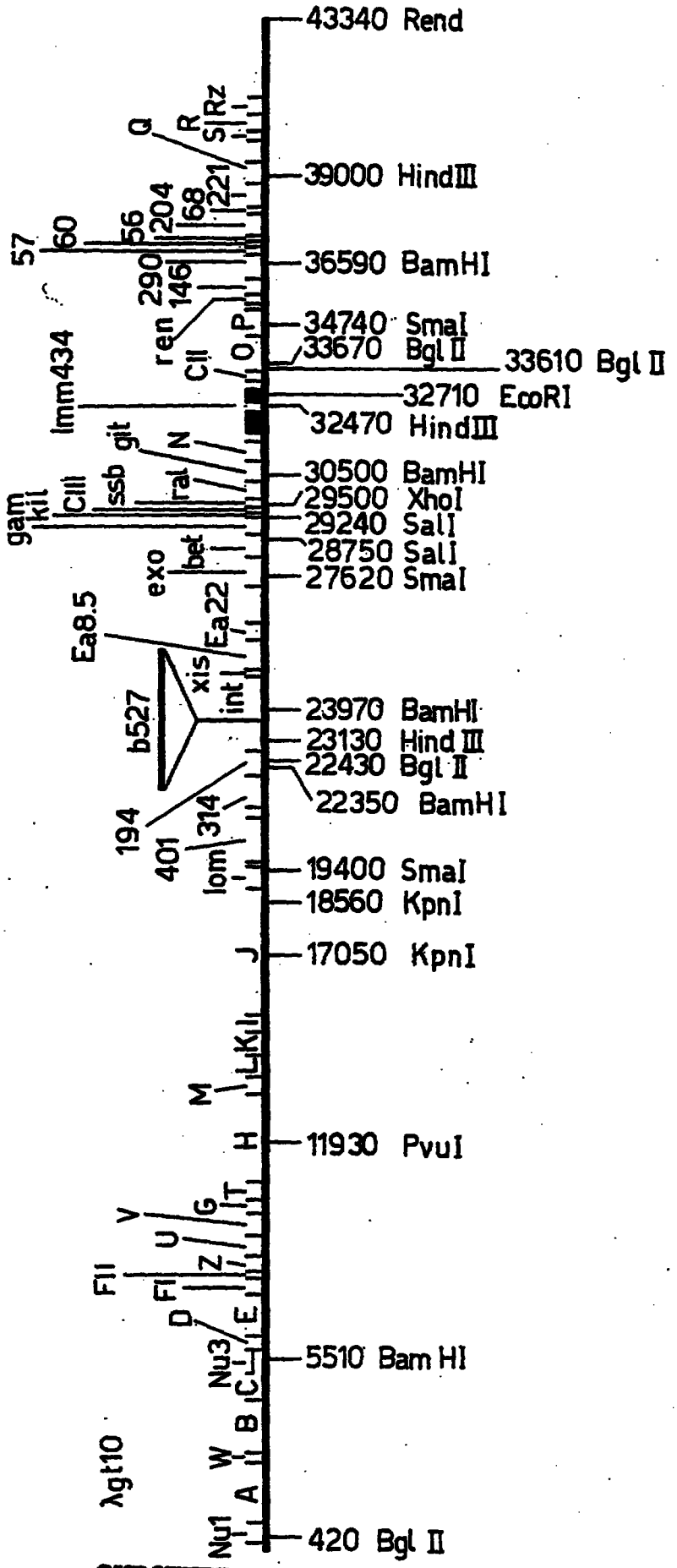

43340 Rend
39000 HindIII
36590 BamHI
34740 SmaI
33670 BglII
33610 BglII
32710 EcoRI
32470 HindIII
30500 BamHI
29500 XhoI
29240 SalI
28750 SalI
27620 SmaI
23970 BamHI
23130 HindIII
22430 BglII
22350 BamHI
19400 SmaI
18560 KpnI
17050 KpnI
11930 PvuI
5510 BamHI
420 BglII
λgt10
Nu1
A
W
B
C
Nu3
D
E
FI
FII
Z
U
V
G
T
H
J
K
L
M
lom
314
401
194
b527
xis
int
Ea22
Ea8.5
exo
bet
gam
kil
CIII
ssb
ral
git
N
imm434
CII
O
P
ren
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146
57
60
56
204
68
221
S
Rz
R
Q

Fig.3.

Fig.4.

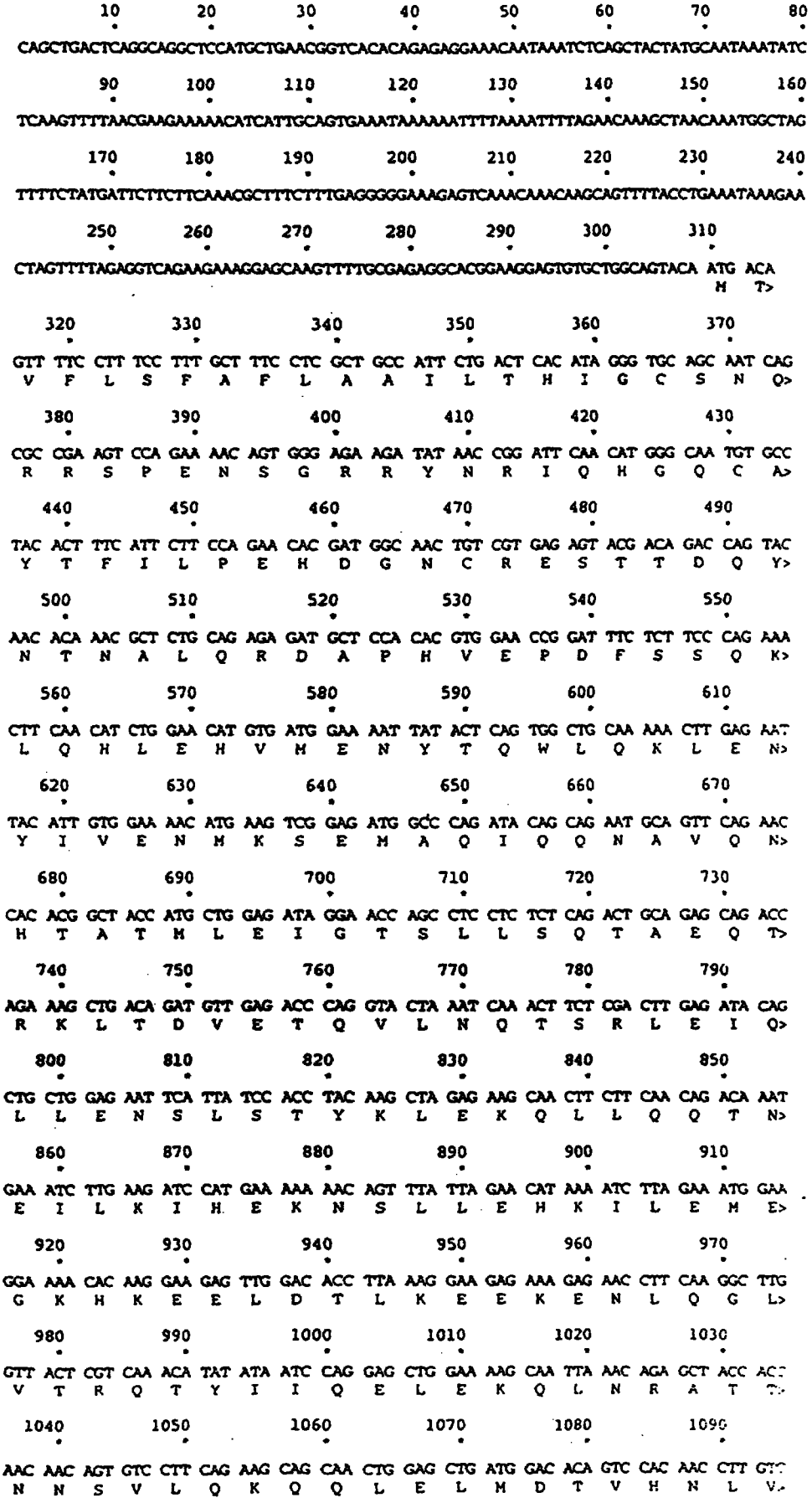

10 20 30 40 50 60 70 80
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90 100 110 120 130 140 150 160
TCAAGTTTTAACGAAGAAAAACATCATTGCAGTGAAATAAAAAATTTTAAAATTTTAGAACAAAGCTAACAAATGGCTAG
170 180 190 200 210 220 230 240
TTTTCTATGATTCTTCTTCAAACGCTTTCTTTGAGGGGGAAAGAGTCAAACAAACAAGCAGTTTTACCTGAAATAAAGAA
250 260 270 280 290 300 310
CTAGTTTTAGAGGTCAGAAGAAAGGAGCAAGTTTTGCGAGAGGCACGGAAGGAGTGTGCTGGCAGTACA ATG ACA
M T>
320 330 340 350 360 370
GTT TTC CTT TCC TTT GCT TTC CTC GCT GCC ATT CTG ACT CAC ATA GGG TGC AGC AAT CAG
V F L S F A F L A A I L T H I G C S N Q>
380 390 400 410 420 430
CGC CGA AGT CCA GAA AAC AGT GGG AGA AGA TAT AAC CGG ATT CAA CAT GGG CAA TGT GCC
R R S P E N S G R R Y N R I Q H G Q C A>
440 450 460 470 480 490
TAC ACT TTC ATT CTT CCA GAA CAC GAT GGC AAC TGT CGT GAG AGT ACG ACA GAC CAG TAC
Y T F I L P E H D G N C R E S T T D Q Y>
500 510 520 530 540 550
AAC ACA AAC GCT CTG CAG AGA GAT GCT CCA CAC GTG GAA CCG GAT TTC TCT TCC CAG AAA
N T N A L Q R D A P H V E P D F S S Q K>
560 570 580 590 600 610
CTT CAA CAT CTG GAA CAT GTG ATG GAA AAT TAT ACT CAG TGG CTG CAA AAA CTT GAG AAT
L Q H L E H V M E N Y T Q W L Q K L E N>
620 630 640 650 660 670
TAC ATT GTG GAA AAC ATG AAG TCG GAG ATG GCC CAG ATA CAG CAG AAT GCA GTT CAG AAC
Y I V E N M K S E M A Q I Q Q N A V Q N>
680 690 700 710 720 730
CAC ACG GCT ACC ATG CTG GAG ATA GGA ACC AGC CTC CTC TCT CAG ACT GCA GAG CAG ACC
H T A T M L E I G T S L L S Q T A E Q T>
740 750 760 770 780 790
AGA AAG CTG ACA GAT GTT GAG ACC CAG GTA CTA AAT CAA ACT TCT CGA CTT GAG ATA CAG
R K L T D V E T Q V L N Q T S R L E I Q>
800 810 820 830 840 850
CTG CTG GAG AAT TCA TTA TCC ACC TAC AAG CTA GAG AAG CAA CTT CTT CAA CAG ACA AAT
L L E N S L S T Y K L E K Q L L Q Q T N>
860 870 880 890 900 910
GAA ATC TTG AAG ATC CAT GAA AAA AAC AGT TTA TTA GAA CAT AAA ATC TTA GAA ATG GAA
E I L K I H E K N S L L E H K I L E M E>
920 930 940 950 960 970
GGA AAA CAC AAG GAA GAG TTG GAC ACC TTA AAG GAA GAG AAA GAG AAC CTT CAA GGC TTG
G K H K E E L D T L K E E K E N L Q G L>
980 990 1000 1010 1020 1030
GTT ACT CGT CAA ACA TAT ATA ATC CAG GAG CTG GAA AAG CAA TTA AAC AGA GCT ACC ACC
V T R Q T Y I I Q E L E K Q L N R A T T>
1040 1050 1060 1070 1080 1090
AAC AAC AGT GTC CTT CAG AAG CAG CAA CTG GAG CTG ATG GAC ACA GTC CAC AAC CTT GTC
N N S V L Q K Q Q L E L M D T V H N L V>
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Fig.4. (Cont.)

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   1100         1110         1120         1130         1140         1150
    .            .            .            .            .            .
AAT CTT TGC ACT AAA GAA GGT GTT TTA CTA AAG GGA GGA AAA AGA GAG GAA GAG AAA CCA
 N   L   C   T   K   E   G   V   L   L   K   G   G   K   R   E   E   E   K   P>

   1160         1170         1180         1190         1200         1210
    .            .            .            .            .            .
TTT AGA GAC TGT GCA GAT GTA TAT CAA GCT GGT TTT AAT AAA AGT GGA ATC TAC ACT ATT
 F   R   D   C   A   D   V   Y   Q   A   G   F   N   K   S   G   I   Y   T   I>

   1220         1230         1240         1250         1260         1270
    .            .            .            .            .            .
TAT ATT AAT AAT ATG CCA GAA CCC AAA AAG GTG TTT TGC AAT ATG GAT GTC AAT GGG GGA
 Y   I   N   N   M   P   E   P   K   K   V   F   C   N   M   D   V   N   G   G>

   1280         1290         1300         1310         1320         1330
    .            .            .            .            .            .
GGT TGG ACT GTA ATA CAA CAT CGT GAA GAT GGA AGT CTA GAT TTC CAA AGA GGC TGG AAG
 G   W   T   V   I   Q   H   R   E   D   G   S   L   D   F   Q   R   G   W   K>

   1340         1350         1360         1370         1380         1390
    .            .            .            .            .            .
GAA TAT AAA ATG GGT TTT GGA AAT CCC TCC GGT GAA TAT TGG CTG GGG AAT GAG TTT ATT
 E   Y   K   M   G   F   G   N   P   S   G   E   Y   W   L   G   N   E   F   I>

   1400         1410         1420         1430         1440         1450
    .            .            .            .            .            .
TTT GCC ATT ACC AGT CAG AGG CAG TAC ATG CTA AGA ATT GAG TTA ATG GAC TGG GAA GGG
 F   A   I   T   S   Q   R   Q   Y   M   L   R   I   E   L   M   D   W   E   G>

   1460         1470         1480         1490         1500         1510
    .            .            .            .            .            .
AAC CGA GCC TAT TCA CAG TAT GAC AGA TTC CAC ATA GGA AAT GAA AAG CAA AAC TAT AGG
 N   R   A   Y   S   Q   Y   D   R   F   H   I   G   N   E   K   Q   N   Y   R>

   1520         1530         1540         1550         1560         1570
    .            .            .            .            .            .
TTG TAT TTA AAA GGT CAC ACT GGG ACA GCA GGA AAA CAG AGC AGC CTG ATC TTA CAC GGT
 L   Y   L   K   G   H   T   G   T   A   G   K   Q   S   S   L   I   L   H   G>

   1580         1590         1600         1610         1620         1630
    .            .            .            .            .            .
GCT GAT TTC AGC ACT AAA GAT GCT GAT AAT GAC AAC TGT ATG TGC AAA TGT GCC CTC ATG
 A   D   F   S   T   X   D   A   D   N   D   N   C   M   C   K   C   A   L   M>

   1640         1650         1660         1670         1680         1690
    .            .            .            .            .            .
TTA ACA GGA GGA TGG TGG TTT GAT GCT TGT GGC CCC TCC AAT CTA AAT GGA ATG TTC TAT
 L   T   G   G   W   W   F   D   A   C   G   P   S   N   L   N   G   M   F   Y>

   1700         1710         1720         1730         1740         1750
    .            .            .            .            .            .
ACT GCG GGA CAA AAC CAT GGA AAA CTG AAT GGG ATA AAG TGG CAC TAC TTC AAA GGG CCC
 T   A   G   Q   N   H   G   K   L   N   G   I   K   W   H   Y   F   K   G   P>

   1760         1770         1780         1790         1800         1810
    .            .            .            .            .            .
AGT TAC TCC TTA CGT TCC ACA ACT ATG ATG ATT CGA CCT TTA GAT TTT TGA AAG CGCAATGT
 S   Y   S   L   R   S   T   T   M   M   I   R   P   L   D   F   *

1820         1830         1840         1850         1860         1870         1880         1890
 .            .            .            .            .            .            .            .
CAGAAGCGATTATGAAAGCAACAAAGAAATCCGGAGAAGCTGCCAGGTGAGAAACTGTTTGAAAACTTCAGAAGCAAACA

1900         1910         1920         1930         1940         1950         1960         1970
 .            .            .            .            .            .            .            .
ATATTGTCTCCCTTCCAGCAATAAGTGGTAGTTATGTGAAGTCACCAAGGTTCTTGACCGTGAATCTGGAGCCGTTTGAG

1980         1990         2000         2010         2020         2030         2040         2050
 .            .            .            .            .            .            .            .
TTCACAAGAGTCTCTACTTGGGGTGACAGTGCTCACGTGGCTCGACTATAGAAAACTCCACTGACTGTCGGGCTTTAAAA

2060         2070         2080         2090         2100         2110         2120         2130
 .            .            .            .            .            .            .            .
AGGGAAGAAACTGCTGAGCTTGCTGTGCTTCAAACTACTACTGGACCTTATTTTGGAACTATGGTAGCCAGATGATAAAT

2140
 .
ATGGTTAATTTC
```

Fig.5.

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        10         20         30         40         50         60         70         80
CAGCTGACTCAGGCAGGCTCCATGCTGAACGGTCACACAGAGAGGAAACAATAAATCTCAGCTACTATGCAATAAATATC

        90        100        110        120        130        140        150        160
TCAAGTTTTAACGAAGAAAAACATCATTGCAGTGAAATAAAAAATTTTAAAATTTTAGAACAAAGCTAACAAATGGCTAG

       170        180        190        200        210        220        230        240
TTTTCTATGATTCTTCTTCAAACGCTTTCTTTGAGGGGGAAAGAGTCAAACAAACAAGCAGTTTTACCTGAAATAAAGAA

       250        260        270        280        290        300        310
CTAGTTTTAGAGGTCAGAAGAAAGGAGCAAGTTTTGCGAGAGGCACGGAAGGAGTGTGCTGGCAGTACA ATG ACA
                                                                        M   T>

       320        330        340        350        360        370
GTT TTC CTT TCC TTT GCT TTC CTC GCT GCC ATT CTG ACT CAC ATA GGG TGC AGC AAT CAG
 V   F   L   S   F   A   F   L   A   A   I   L   T   H   I   G   C   S   N   Q>

       380        390        400        410        420        430
CGC CGA AGT CCA GAA AAC AGT GGG AGA AGA TAT AAC CGG ATT CAA CAT GGG CAA TGT GCC
 R   R   S   P   E   N   S   G   R   R   Y   N   R   I   Q   H   G   Q   C   A>

       440        450        460        470        480        490
TAC ACT TTC ATT CTT CCA GAA CAC GAT GGC AAC TGT CGT GAG AGT ACG ACA GAC CAG TAC
 Y   T   F   I   L   P   E   H   D   G   N   C   R   E   S   T   T   D   Q   Y>

       500        510        520        530        540        550
AAC ACA AAC GCT CTG CAG AGA GAT GCT CCA CAC GTG GAA CCG GAT TTC TCT TCC CAG AAA
 N   T   N   A   L   Q   R   D   A   P   H   V   E   P   D   F   S   S   Q   K>

       560        570        580        590        600        610
CTT CAA CAT CTG GAA CAT GTG ATG GAA AAT TAT ACT CAG TGG CTG CAA AAA CTT GAG AAT
 L   Q   H   L   E   H   V   M   E   N   Y   T   Q   W   L   Q   K   L   E   N>

       620        630        640        650        660        670
TAC ATT GTG GAA AAC ATG AAG TCG GAG ATG GCC CAG ATA CAG CAG AAT GCA GTT CAG AAC
 Y   I   V   E   N   M   K   S   E   M   A   Q   I   Q   Q   N   A   V   Q   N>

       680        690        700        710        720        730
CAC ACG GCT ACC ATG CTG GAG ATA GGA ACC AGC CTC CTC TCT CAG ACT GCA GAG CAG ACC
 H   T   A   T   M   L   E   I   G   T   S   L   L   S   Q   T   A   E   Q   T>

       740        750        760        770        780        790
AGA AAG CTG ACA GAT GTT GAG ACC CAG GTA CTA AAT CAA ACT TCT CGA CTT GAG ATA CAG
 R   K   L   T   D   V   E   T   Q   V   L   N   Q   T   S   R   L   E   I   Q>

       800        810        820        830        840        850
CTG CTG GAG AAT TCA TTA TCC ACC TAC AAG CTA GAG AAG CAA CTT CTT CAA CAG ACA AAT
 L   L   E   N   S   L   S   T   Y   K   L   E   K   Q   L   L   Q   Q   T   N>

       860        870        880        890        900        910
GAA ATC TTG AAG ATC CAT GAA AAA AAC AGT TTA TTA GAA CAT AAA ATC TTA GAA ATG GAA
 E   I   L   K   I   H   E   K   N   S   L   L   E   H   K   I   L   E   M   E>

       920        930        940        950        960        970
GGA AAA CAC AAG GAA GAG TTG GAC ACC TTA AAG GAA GAG AAA GAG AAC CTT CAA GGC TTG
 G   K   H   K   E   E   L   D   T   L   K   E   E   K   E   N   L   Q   G   L>

       980        990       1000       1010       1020       1030
GTT ACT CGT CAA ACA TAT ATA ATC CAG GAG CTG GAA AAG CAA TTA AAC AGA GCT ACC ACC
 V   T   R   Q   T   Y   I   I   Q   E   L   E   K   Q   L   N   R   A   T   T>

      1040       1050       1060       1070       1080       1090
AAC AAC AGT GTC CTT CAG AAG CAG CAA CTG GAG CTG ATG GAC ACA GTC CAC AAC CTT GTC
 N   N   S   V   L   Q   K   Q   Q   L   E   L   M   D   T   V   H   N   L   V>
```

Fig.5. (Cont.)

```
      1100        1110        1120        1130        1140        1150
       .           .           .           .           .           .
AAT CTT TGC ACT AAA GAA GTT TTA CTA AAG GGA GGA AAA AGA GAG GAA GAG AAA CCA TTT
 N   L   C   T   K   E   V   L   L   K   G   G   K   R   E   E   E   K   P   F>

      1160        1170        1180        1190        1200        1210
       .           .           .           .           .           .
AGA GAC TGT GCA GAT GTA TAT CAA GCT GGT TTT AAT AAA AGT GGA ATC TAC ACT ATT TAT
 R   D   C   A   D   V   Y   Q   A   G   F   N   K   S   G   I   Y   T   I   Y>

      1220        1230        1240        1250        1260        1270
       .           .           .           .           .           .
ATT AAT AAT ATG CCA GAA CCC AAA AAG GTG TTT TGC AAT ATG GAT GTC AAT GGG GGA GGT
 I   N   N   M   P   E   P   K   K   V   F   C   N   M   D   V   N   G   G   G>

      1280        1290        1300        1310        1320        1330
       .           .           .           .           .           .
TGG ACT GTA ATA CAA CAT CGT GAA GAT GGA AGT CTA GAT TTC CAA AGA GGC TGG AAG GAA
 W   T   V   I   Q   H   R   E   D   G   S   L   D   F   Q   R   G   W   K   E>

      1340        1350        1360        1370        1380        1390
       .           .           .           .           .           .
TAT AAA ATG GGT TTT GGA AAT CCC TCC GGT GAA TAT TGG CTG GGG AAT GAG TTT ATT TTT
 Y   K   M   G   F   G   N   P   S   G   E   Y   W   L   G   N   E   F   I   F>

      1400        1410        1420        1430        1440        1450
       .           .           .           .           .           .
GCC ATT ACC AGT CAG AGG CAG TAC ATG CTA AGA ATT GAG TTA ATG GAC TGG GAA GGG AAC
 A   I   T   S   Q   R   Q   Y   M   L   R   I   E   L   M   D   W   E   G   N>

      1460        1470        1480        1490        1500        1510
       .           .           .           .           .           .
CGA GCC TAT TCA CAG TAT GAC AGA TTC CAC ATA GGA AAT GAA AAG CAA AAC TAT AGG TTG
 R   A   Y   S   Q   Y   D   R   F   H   I   G   N   E   K   Q   N   Y   R   L>

      1520        1530        1540        1550        1560        1570
       .           .           .           .           .           .
TAT TTA AAA GGT CAC ACT GGG ACA GCA GGA AAA CAG AGC AGC CTG ATC TTA CAC GGT GCT
 Y   L   K   G   H   T   G   T   A   G   K   Q   S   S   L   I   L   H   G   A>

      1580        1590        1600        1610        1620        1630
       .           .           .           .           .           .
GAT TTC AGC ACT AAA GAT GCT GAT AAT GAC AAC TGT ATG TGC AAA TGT GCC CTC ATG TTA
 D   F   S   T   K   D   A   D   N   D   N   C   M   C   K   C   A   L   M   L>

      1640        1650        1660        1670        1680        1690
       .           .           .           .           .           .
ACA GGA GGA TGG TGG TTT GAT GCT TGT GGC CCC TCC AAT CTA AAT GGA ATG TTC TAT ACT
 T   G   G   W   W   F   D   A   C   G   P   S   N   L   N   G   M   F   Y   T>

      1700        1710        1720        1730        1740        1750
       .           .           .           .           .           .
GCG GGA CAA AAC CAT GGA AAA CTG AAT GGG ATA AAG TGG CAC TAC TTC AAA GGG CCC AGT
 A   G   Q   N   H   G   K   L   N   G   I   K   W   H   Y   F   K   G   P   S>

      1760        1770        1780        1790        1800        1810
       .           .           .           .           .           .
TAC TCC TTA CGT TCC ACA ACT ATG ATG ATT CGA CCT TTA GAT TTT TGA AAGCGCAATGTCAGAA
 Y   S   L   R   S   T   T   M   M   I   R   P   L   D   F   *>

1820        1830        1840        1850        1860        1870        1880        1890
 .           .           .           .           .           .           .           .
GCGATTATGAAAGCAACAAAGAAATCCGGAGAAGCTGCCAGGTGAGAAACTGTTTGAAAACTTCAGAAGCAAACAATATT

1900        1910        1920        1930        1940        1950        1960        1970
 .           .           .           .           .           .           .           .
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1980        1990        2000        2010        2020        2030        2040        2050
 .           .           .           .           .           .           .           .
AAGAGTCTCTACTTGGGGTGACAGTGCTCACGTGGCTCGACTATAGAAAACTCCACTGACTGTCGGGCTTTAAAAAGGGA

2060        2070        2080        2090        2100        2110        2120        2130
 .           .           .           .           .           .           .           .
AGAAACTGCTGAGCTTGCTGTGCTTCAAACTACTACTGGACCTTATTTTGGAACTATGGTAGCCAGATGATAAATATGGT

2140
 .
TAATTTC
```

Fig.6.


```
        10         20         30         40         50         60         70         80
GAATTCCTGGGTTGGTGTTTATCTCCTCCCAGCCTTGAGGGAGGGAACAACACTGTAGGATCTGGGGAGAGAGGAACAAA

        90        100        110        120        130        140        150        160
GGACCGTGAAAGCTGCTCTGTAAAAGCTGACACAGCCCTCCCAAGTGAGCAGGACTGTTCTTCCCACTGCAATCTGACAG

       170        180        190        200        210        220        230        240
TTTACTGCATGCCTGGAGAGAACACAGCAGTAAAAACCAGGTTTGCTACTGGAAAAAGAGGAAAGAGAAGACTTTCATTG

       250        260        270        280        290        300        310        320
ACGGACCCAGCCATGGCAGCGTAGCAGCCCTGCGTTTCAGACGGCAGCAGCTCGGGACTCTGGACGTGTGTTTGCCCTCA

       330        340        350        360        370        380
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                                        M   W   Q   I   V   F   F   T   L   S   C>

390        400        410        420        430        440
GAT CTT GTC TTG GCC GCA GCC TAT AAC AAC TTT CGG AAG AGC ATG GAC AGC ATA GGA AAG
 D   L   V   L   A   A   A   Y   N   N   F   R   K   S   M   D   S   I   G   K>

450        460        470        480        490        500
AAG CAA TAT CAG GTC CAG CAT GGG TCC TGC AGC TAC ACT TTC CTC CTG CCA GAG ATG GAC
 K   Q   Y   Q   V   Q   H   G   S   C   S   Y   T   F   L   L   P   E   M   D>

510        520        530        540        550        560
AAC TGC CGC TCT TCC TCC AGC CCC TAC GTG TCC AAT GCT GTG CAG AGG GAC GCG CCG CTC
 N   C   R   S   S   S   S   P   Y   V   S   N   A   V   Q   R   D   A   P   L>

570        580        590        600        610        620
GAA TAC GAT GAC TCG GTG CAG AGG CTG CAA GTG CTG GAG AAC ATC ATG GAA AAC AAC ACT
 E   Y   D   D   S   V   Q   R   L   Q   V   L   E   N   I   M   E   N   N   T>

630        640        650        660        670        680
CAG TGG CTA ATG AAG CTT GAG AAT TAT ATC CAG GAC AAC ATG AAG AAA GAA ATG GTA GAG
 Q   W   L   M   K   L   E   N   Y   I   Q   D   N   M   K   K   E   M   V   E>

690        700        710        720        730        740
ATA CAG CAG AAT GCA GTA CAG AAC CAG ACG GCT GTG ATG ATA GAA ATA GGG ACA AAC CTG
 I   Q   Q   N   A   V   Q   N   Q   T   A   V   M   I   E   I   G   T   N   L>

750        760        770        780        790        800
TTG AAC CAA ACA GCT GAG CAA ACG CGG AAG TTA ACT GAT GTG GAA GCC CAA GTA TTA AAT
 L   N   Q   T   A   E   Q   T   R   K   L   T   D   V   E   A   Q   V   L   N>

810        820        830        840        850        860
CAG ACC ACG AGA CTT GAA CTT CAG CTC TTG GAA CAC TCC CTC TCG ACA AAC AAA TTG GAA
 Q   T   T   R   L   E   L   Q   L   L   E   H   S   L   S   T   N   K   L   E>

870        880        890        900        910        920
AAA CAG ATT TTG GAC CAG ACC AGT GAA ATA AAC AAA TTG CAA GAT AAG AAC AGT TTC CTA
 K   Q   I   L   D   Q   T   S   E   I   N   K   L   Q   D   K   N   S   F   L>

930        940        950        960        970        980
GAA AAG AAG GTG CTA GCT ATG GAA GAC AAG CAC ATC ATC CAA CTA CAG TCA ATA AAA GAA
 E   K   K   V   L   A   M   E   D   K   H   I   I   Q   L   Q   S   I   K   E>

990        1000       1010       1020       1030       1040
GAG AAA GAT CAG CTA CAG GTG TTA GTA TCC AAG CAA AAT TCC ATC ATT GAA GAA CTA GAA
 E   K   D   Q   L   Q   V   L   V   S   K   Q   N   S   I   I   E   E   L   E>

1050       1060       1070       1080       1090       1100
AAA AAA ATA GTG ACT GCC ACG GTG AAT AAT TCA GTT CTT CAA AAG CAG CAA CAT GAT CTT
 K   K   I   V   T   A   T   V   N   N   S   V   L   Q   K   Q   Q   H   D   L>
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Fig.6. (Cont.)

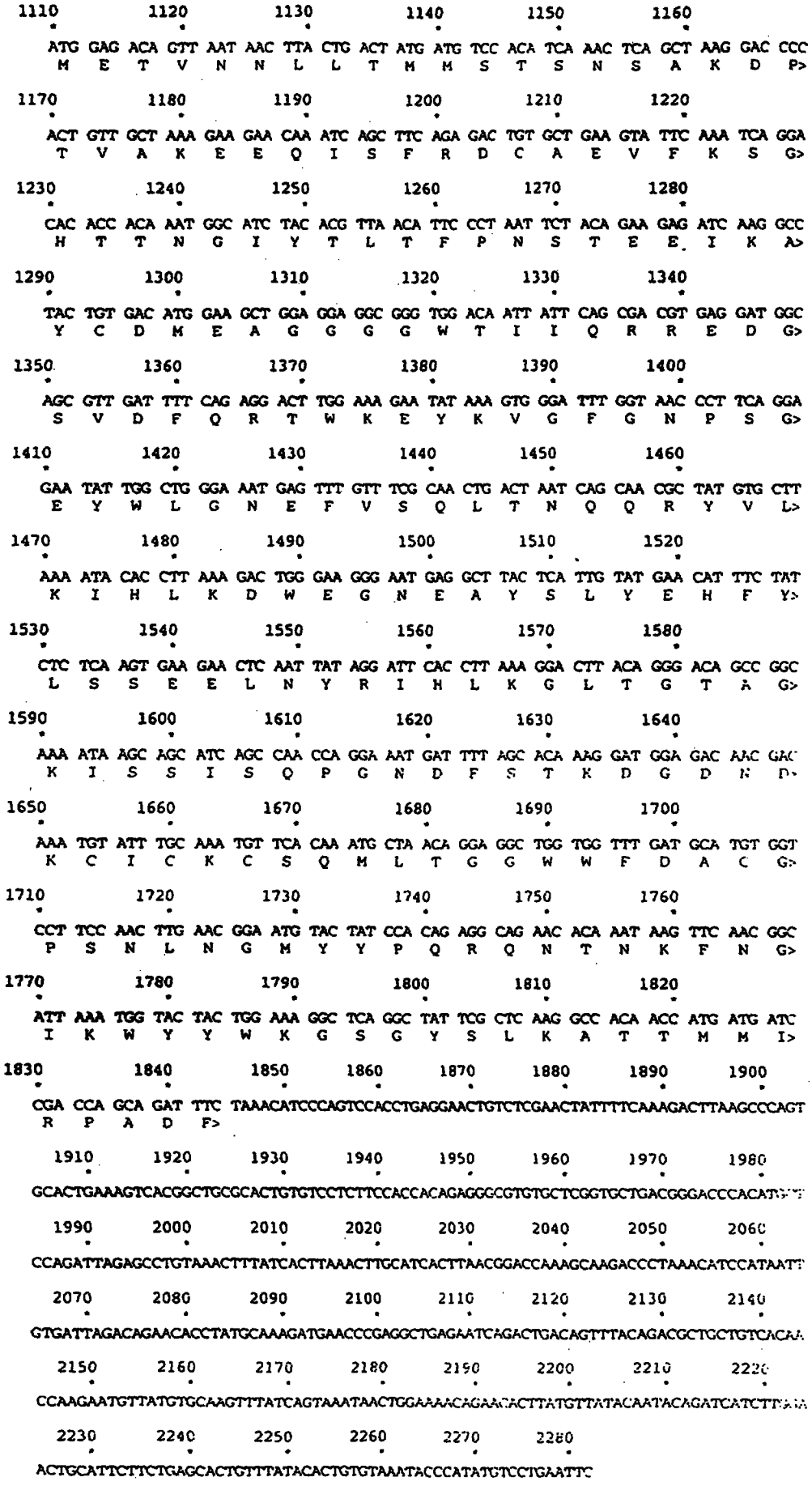

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T V A K E E Q I S F R D C A E V F K S G>
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H T T N G I Y T L T F P N S T E E I K A>
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TAC TGT GAC ATG GAA GCT GGA GGA GGC GGG TGG ACA ATT ATT CAG CGA CGT GAG GAT GGC
Y C D M E A G G G G W T I I Q R R E D G>
1350 1360 1370 1380 1390 1400
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S V D F Q R T W K E Y K V G F G N P S G>
1410 1420 1430 1440 1450 1460
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E Y W L G N E F V S Q L T N Q Q R Y V L>
1470 1480 1490 1500 1510 1520
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K I H L K D W E G N E A Y S L Y E H F Y>
1530 1540 1550 1560 1570 1580
CTC TCA AGT GAA GAA CTC AAT TAT AGG ATT CAC CTT AAA GGA CTT ACA GGG ACA GCC GGC
L S S E E L N Y R I H L K G L T G T A G>
1590 1600 1610 1620 1630 1640
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K I S S I S Q P G N D F S T K D G D N D>
1650 1660 1670 1680 1690 1700
AAA TGT ATT TGC AAA TGT TCA CAA ATG CTA ACA GGA GGC TGG TGG TTT GAT GCA TGT GGT
K C I C K C S Q M L T G G W W F D A C G>
1710 1720 1730 1740 1750 1760
CCT TCC AAC TTG AAC GGA ATG TAC TAT CCA CAG AGG CAG AAC ACA AAT AAG TTC AAC GGC
P S N L N G M Y Y P Q R Q N T N K F N G>
1770 1780 1790 1800 1810 1820
ATT AAA TGG TAC TAC TGG AAA GGC TCA GGC TAT TCG CTC AAG GCC ACA ACC ATG ATG ATC
I K W Y Y W K G S G Y S L K A T T M M I>
1830 1840 1850 1860 1870 1880 1890 1900
CGA CCA GCA GAT TTC TAAACATCCCAGTCCACCTGAGGAACTGTCTCGAACTATTTTCAAAGACTTAAGCCCAGT
R P A D F>
1910 1920 1930 1940 1950 1960 1970 1980
GCACTGAAAGTCACGGCTGCGCACTGTGTCCTCTTCCACCACAGAGGGCGTGTGCTCGGTGCTGACGGGACCCACATG
1990 2000 2010 2020 2030 2040 2050 2060
CCAGATTAGAGCCTGTAAACTTTATCACTTAAACTTGCATCACTTAACGGACCAAAGCAAGACCCTAAACATCCATAATT
2070 2080 2090 2100 2110 2120 2130 2140
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2150 2160 2170 2180 2190 2200 2210 2220
CCAAGAATGTTATGTGCAAGTTTATCAGTAAATAACTGGAAAACAGAACACTTATGTTATACAATACAGATCATCTTAAA
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Fig.7.

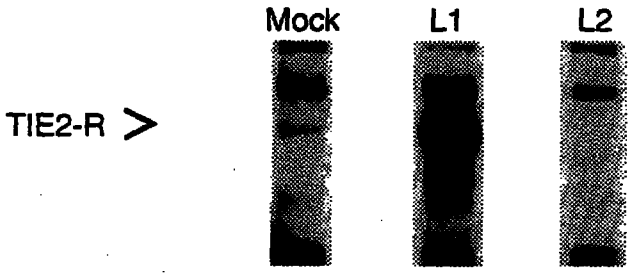


Fig.8.

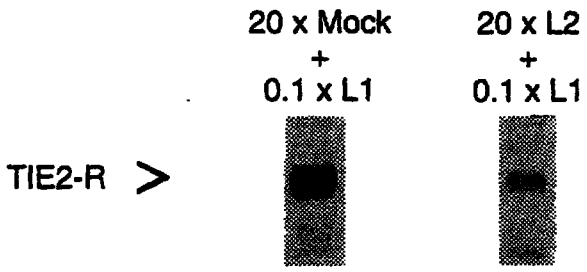


Fig.9.

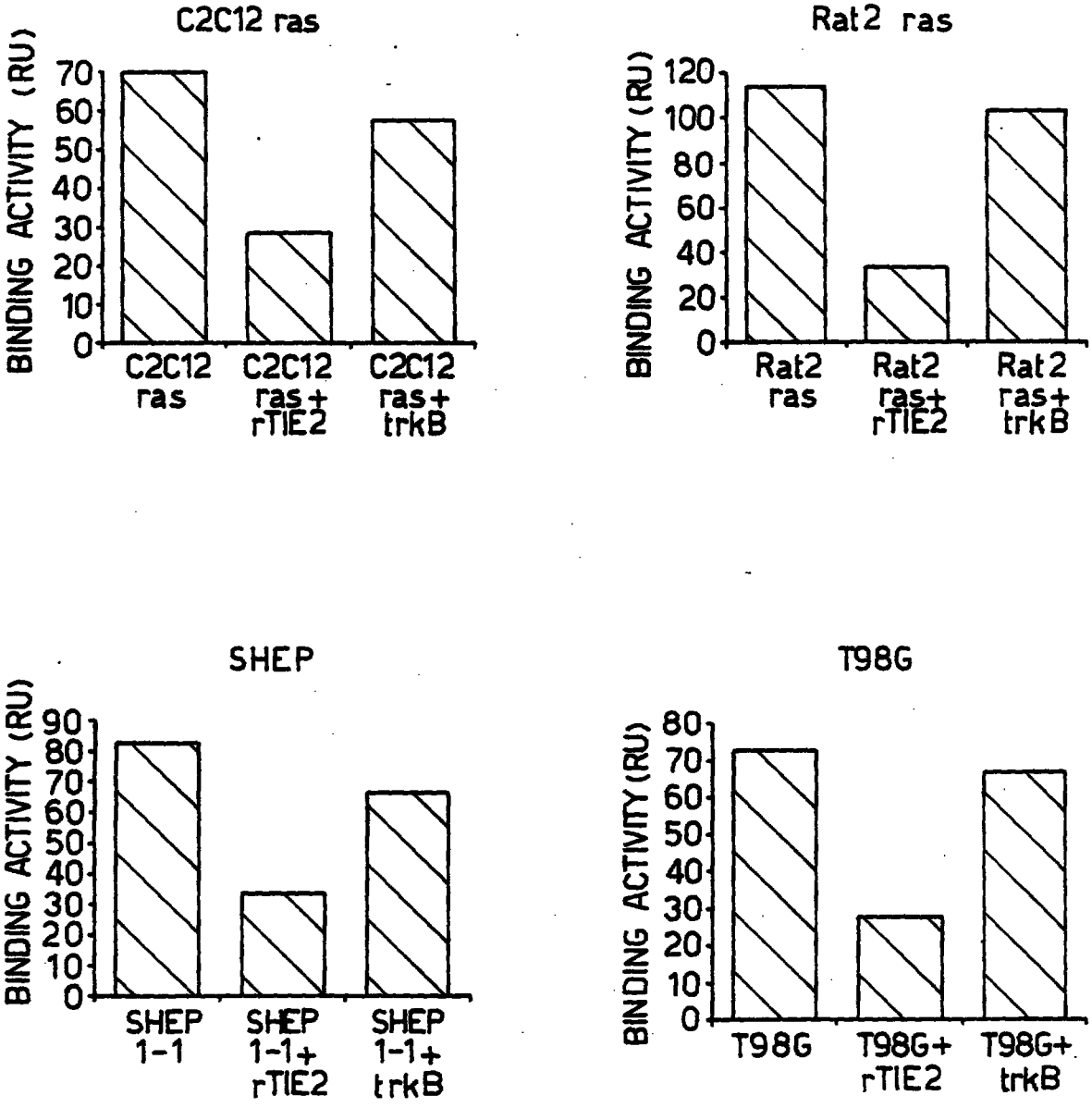
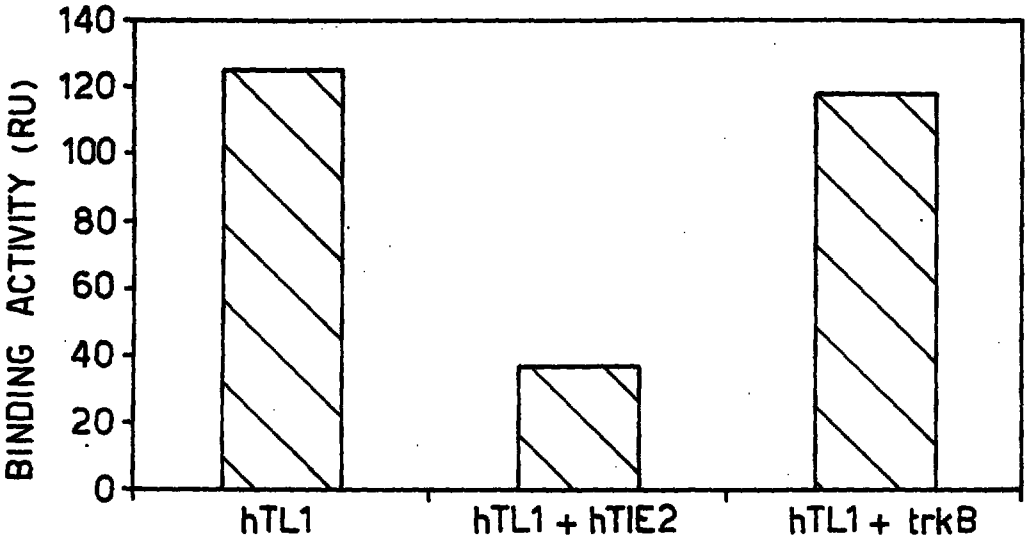

C2C12 ras
BINDING ACTIVITY (RU)
0
10
20
30
40
50
60
70
C2C12 ras
C2C12 ras+ rTIE2
C2C12 ras+ trkB
Rat2 ras
BINDING ACTIVITY (RU)
0
20
40
60
80
100
120
Rat2 ras
Rat2 ras+ rTIE2
Rat2 ras+ trkB
SHEP
BINDING ACTIVITY (RU)
0
10
20
30
40
50
60
70
80
90
SHEP 1-1
SHEP 1-1+ rTIE2
SHEP 1-1+ trkB
T98G
BINDING ACTIVITY (RU)
0
10
20
30
40
50
60
70
80
T98G
T98G+ rTIE2
T98G+ trkB

Fig.10.


BINDING ACTIVITY (RU)
140
120
100
80
60
40
20
0
hTL1
hTL1 + hTIE2
hTL1 + trkB

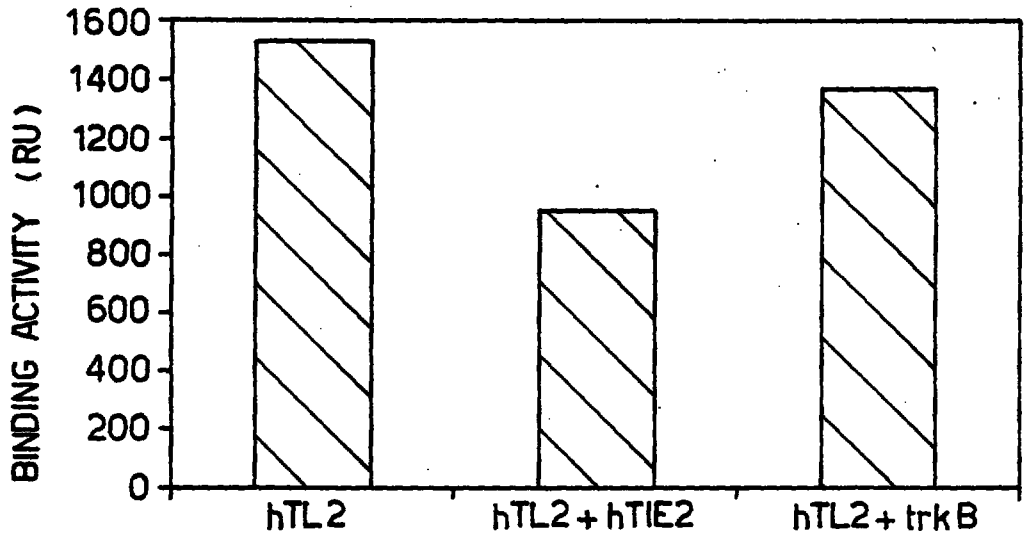
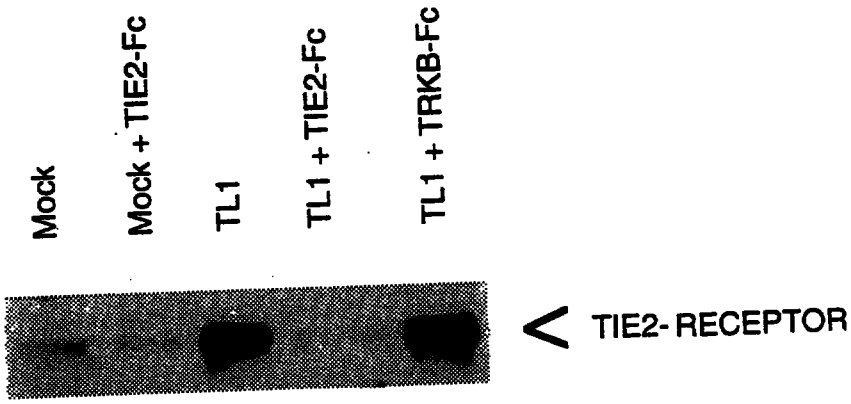

BINDING ACTIVITY (RU)
1600
1400
1200
1000
800
600
400
200
0
hTL2
hTL2 + hTIE2
hTL2 + trkB

Fig.11.



CLAIMS

1. An isolated nucleic acid molecule encoding a TIE-2 ligand.

2. An isolated nucleic acid molecule according to claim 1 wherein the nucleic acid sequence is:

(a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4 or Figure 5;

(b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or

(c) a nucleic acid sequence which, but for the degeneracy of the genetic code, would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

3. An isolated nucleic acid molecule according to claim 2 wherein the encoded TIE-2 ligand is a TIE-2 agonist.

4. An isolated nucleic acid molecule according to claim 1 wherein the nucleic acid sequence is

(a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 6;

(b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or

(c) a nucleic acid sequence which, but for the degeneracy of the genetic code would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

5. An isolated nucleic acid molecule according to claim 4 wherein the encoded TIE-2 ligand is a TIE-2 antagonist.

6. A nucleic acid molecule which is degenerate of a sequence of claim 2 or 4 and encodes a ligand that binds the TIE-2 receptor.

7. A nucleic acid molecule which hybridizes under moderately stringent conditions to a sequence encoding a TIE-2 ligand as set out in Figure 4, 5 or 6.

8. A vector which comprises a nucleic acid molecule of any one of the preceding claims.

9. A vector according to claim 8 wherein the nucleic acid molecule is operatively linked to an expression control sequence capable of directing its expression in a host cell.

10. A vector according to claim 8 or 9 which is a plasmid.

11. A plasmid according to claim 10 designated pJFE14 encoding TIE-2 ligand (ATCC Accession No. 75910).

12. A plasmid according to claim 10 designated pBluescript KS encoding human TIE-2 ligand 2 (ATCC Accession No. 75963).

13. A vector according to claim 8 or 9 designated as λgtl0 encoding hTIE-2 ligand 1 (ATCC Accession No. 75928).

14. An isolated TIE-2 ligand substantially free of other proteins.

15. An isolated TIE-2 ligand according to claim 14 encoded by a nucleic acid molecule according to claim 1.

16. An isolated TIE-2 ligand according to claim 14 encoded by a nucleic acid according to claim 2 or 3.

17. An isolated TIE-2 ligand according to claim 14 encoded by a nucleic acid according to claim 4 or 5.

18. A host-vector system for the production of a ligand according to any one of claims 14 to 17 which comprises a vector according to any one of claims 8 to 13 in a host cell.

19. A host-vector system according to claim 18 wherein the host cell is a bacterial, yeast, insect or mammalian cell.

20. A host vector system comprising the host vector system of claim 18 or 19 and a nucleic acid encoding the TIE-2 receptor.

21. A method of producing a ligand as defined in any one of claims 14 to 17 which comprises growing cells of a host-vector system according to any one of claims 18 to 20 under conditions permitting production of the ligand, and recovering the ligand so produced.

22. An antibody which specifically binds the ligand of any one of claims 14 to 17.

23. An antibody according to claim 22 which is a monoclonal antibody.

24. A receptorbody which specifically binds the ligand of any one of claims 14 to 17.

25. An isolated nucleic acid molecule encoding a receptorbody according to claim 24.

26. A vector comprising a nucleic acid molecule according to claim 25.

27. A vector according to claim 26 which is a plasmid.

28. A plasmid according to claim 27 designated vTIE-2 receptorbody (ATCC Deposit VR2484).

29. A conjugate comprising a ligand according to any one of claims 14 to 17 and, conjugated thereto, a cytotoxic agent.

30. A conjugate according to claim 29 wherein the cytotoxic agent is a radioisotope or toxin.

31. A pharmaceutical composition comprising a TIE-2 ligand according to any one of claims 14 to 17 and a pharmaceutically acceptable carrier.

32. A pharmaceutical composition comprising an antibody according to claim 22 or 23 and a pharmaceutically acceptable carrier.

33. A pharmaceutical composition comprising a receptorbody according to claim 24 and a pharmaceutically acceptable carrier.

34. A pharmaceutical composition comprising a conjugate according to claim 29 or 30 and a pharmaceutically acceptable carrier.

35. A ligand according to any one of claims 14 to 17, an antibody according to claim 22 or 23, a receptorbody according to claim 24, a conjugate according to claim 29 or 30. or a composition according to any

* C L M S *

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one of claims 31 to 34 for use in a method of treatment of the human or animal body, or in a method of diagnosis.

36. A ligand according to claim 16 for use in a method of treatment of the human or animal body.

37. A ligand according to claim 17 for use in a method of treatment of the human or animal body.

38. An antibody or receptorbody according to claim 35, which antibody or receptorbody specifically binds the ligand of claim 16, for use in a method of blocking blood vessel growth in a mammal.

39. An antibody or receptorbody according to claim 38 for use in a method wherein the mammal is a human.

40. A ligand according to claim 36 for use in a method of promoting neovascularization in a mammal.

41. A ligand according to claim 40 for use in the promotion of wound healing.

42. A ligand according to claim 40 for use in the treatment of ischemia.

43. A TIE-2 antagonist for use in a method of inhibiting TIE-2 ligand activity in a mammal.

44. An antagonist according to claim 43 which is an antibody capable of specifically binding TIE-2 receptor.

AMENDED SHEET (ARTICLE 19)

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45. An antibody according to claim 43 which is an antibody according to claim 22 or 23.

46. An antagonist according to claim 43 which is a receptorbody according to claim 24.

47. An antagonist according to claim 43 which is a ligand according to claim 17.

48. An antagonist according to any one of claims 43 to 47 for use in a method wherein the mammal is a human.

49. An antagonist according to any one of claims 43 to 48 for use in a method of attenuating or preventing tumour growth in a human.

50. A method of maintaining a TIE-2 receptor expressing cell in culture, which method comprises administering to the TIE-2 receptor expressing cell an effective amount of the ligand of claim 16.

51. A method according to claim 50 wherein the TIE-2 receptor expressing cell is an endothelial cell.

52. A method of identifying a TIE-2 receptor antagonist comprising contacting cells expressing the TIE-2 receptor with:

a) a test compound; and

b) a ligand according to claim 16 or 17;

under conditions permitting binding of the ligand to the receptor and determining whether the test compound is capable ofinterfering with the binding of the ligand to the receptor.

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AMENDED SHEET (ARTICLE 19)

53. A polypeptide produced by the method of claim 21.

54. A nucleic acid according to claim 1 or 25, substantially as hereinbefore described with reference to any one of the foregoing Examples.

55. A vector according to claim 8 or 26, substantially as hereinbefore described with reference to any one of the foregoing Examples.

56. A ligand according to any one of claims 14 to 17, substantially as hereinbefore described with reference to any one of the foregoing Examples.

57. A host-vector system according to claim 18, substantially as hereinbefore described with reference to any one of the foregoing Examples.

58. A method according to claim 21, 50 or 52, substantially as hereinbefore described with reference to any one of the foregoing Examples.

59. An antibody according to claim 22 or 35, substantially as hereinbefore described with reference to any one of the foregoing Examples.

60. A receptorbody according to claim 24 or 35, substantially as hereinbefore described with reference to any one of the foregoing Examples.

61. A conjugate according to claim 29 or 35, substantially as hereinbefore described with reference to any one of the foregoing Examples.

62. A composition according to any one of claims 31 to 35, substantially as hereinbefore described with reference to any one of the foregoing Examples.

63. A TIE-2 antagonist according to claim 43, substantially as hereinbefore described with reference to any one of the foregoing Examples.

64. A ligandbody which specifically binds the TIE-2 receptor or the receptorbody of claim 24, 35 or 60.

65. A ligandbody which comprises a TIE-2 ligand fused to an immunoglobulin constant region.

66. The ligandbody of claim 65 wherein the TIE-2 ligand is TIE-2 ligand 1 or TIE-2 ligand 2 and the immunoglobulin constant region is the Fc portion of human IgG1.

67. A ligandbody according to any one of claims 64 to 66 for use in a method of treatment of the human or animal body, or in a method of diagnosis.

68. A method of treating a human or animal subject comprising administering to the subject an effective amount of a ligand according to any one of claims 14 to 17, an antibody according to claim 22 or 23, a receptorbody according to claim 24, a conjugate according to claim 29 or 30, a composition according to any one of claims 31 to 34, or a ligandbody according to any one of claims 64 to 67.

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21 MAY 1996

69. A method according to claim 68, the method being as defined in any one of claims 38 to 49, or 67.

AMENDED SHEET (ARTICLE 19)

CORRECTED VERSION*

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Without international search report and to be republished upon receipt of that report.

With an indication in relation to a deposited microorganism furnished under Rule 13bis separately from the description.

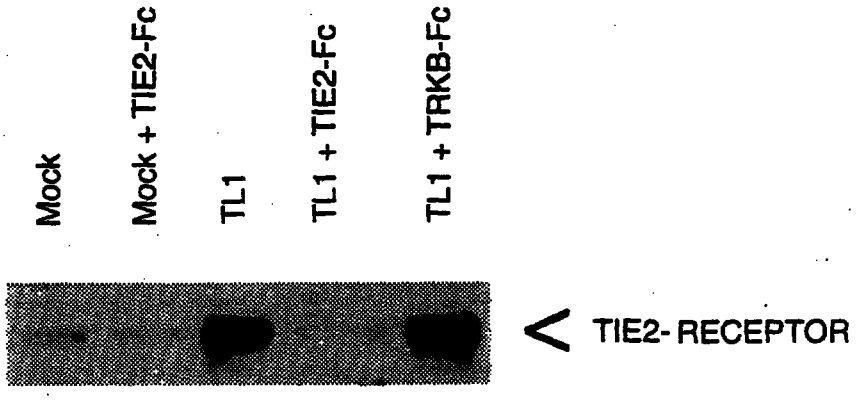
Date of receipt by the International Bureau:
15 March 1996 (15.03.96)

(54) Title: TIE-2 LIGANDS, METHODS OF MAKING AND USES THEREOF

RTMP copy

(57) Abstract

The present invention provides for an isolated nucleic acid molecule encoding a human TIE-2 ligand. In addition, the invention provides for a receptor body which specifically binds a human TIE-2 ligand. The invention also provides an antibody which specifically binds a human TIE-2 ligand. The invention further provides for an antagonist of human TIE-2. The invention also provides for therapeutic compositions as well as a method of blocking blood vessel growth, a method of promoting neovascularization, a method of promoting the growth or differentiation of a cell expressing the TIE-2 receptor, a method of blocking the growth or differentiation of a cell expressing the TIE-2 receptor and a method of attenuating or preventing tumor growth in a human.



* (Referred to in PCT Gazette No. 26/1996, Section II)