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(54) Title: MUSCLE PERFORMANCE IMPROVEMENT COMPOUNDS

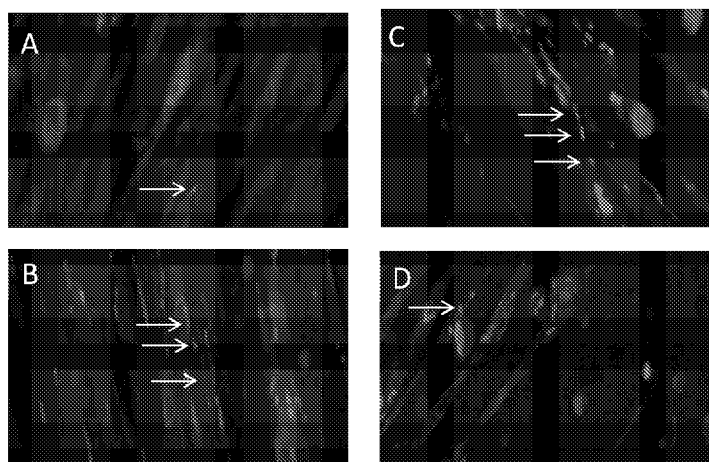


Figure 1

(57) Abstract: A compound comprising at least two components, a first component being the nLG3 domain from the C-terminus of human agrin, and at least one second component, selected from proteins or an antagonistic antibody that inhibit ActR2B-induced signaling activity in the presence of myostatin, the components being linked by means of linking entities. Such compounds are effective treatments for neuromuscular diseases and problems.



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## MUSCLE PERFORMANCE IMPROVEMENT COMPOUNDS

This disclosure relates to a method of treating pathological disorders or diseases affecting the function of muscles and to compounds for use in such treatments. The methods are particularly  
5 suitable for treating, preventing, ameliorating or diagnosing pathological disorders.

Muscles are the contractile tissue responsible for all movement in living organisms. Any loss of muscle function is invariably harmful to a greater or lesser extent. Key elements of muscle function are strength, power and endurance. Muscle strength is the amount of force a muscle, or group of  
10 muscles, can exert upon maximal contraction, generally against an external load. Muscle strength is expressed as the greatest measurable force that can be exerted by a muscle or muscle group to overcome resistance during a single, maximal effort. Muscle power is force developed quickly and combines strength and speed. It is the rate of performing work. Muscular endurance is the ability of a muscle or group of muscles to sustain in a prolonged fashion or repeatedly exert force against  
15 resistance.

These performances can be evaluated by regular assessments of a subject in a testing regime, such as an exercise machine. Most important is the loss of muscle endurance. For example, improvement in a 6-minute walking test or "6MWT" (see, for example, Bautmans *et al* (BMC  
20 Geriatr. 2004 Jul 23;4:6)) and Enright (Respir Care. 2003 Aug;48(8):783-5) is a prerequisite for approval by regulatory authorities such as the US Food and Drug Administration of a drug intended to treat pathological disorders affecting the functioning of the muscle.

There are many pathological disorders that can lead to the loss of muscle function. As used herein,  
25 a "pathological disorder" includes, but is not limited to, neuromuscular diseases. Neuromuscular diseases is a very broad term that encompasses many diseases and ailments that impair the functioning of the muscles, either directly, being pathologies of the voluntary muscle, or indirectly, being pathologies of nerves or neuromuscular junctions.

30 One pathological disorder that can lead to loss of muscle function is muscle atrophy. There are many causes of muscle atrophy, including the result of treatment with a glucocorticoid such as cortisol, dexamethasone, betamethasone, prednisone, methylprednisolone, or prednisolone. The muscle atrophy can also be a result of denervation due to nerve trauma or a result of degenerative, metabolic, or inflammatory neuropathy (e.g., Guillian-Barre syndrome, peripheral neuropathy, or  
35 exposure to environmental toxins or drugs).

In addition, the muscle atrophy can be a result of myopathy, such as myotonia; a congenital myopathy, including nemaline myopathy, multi/minicore myopathy and myotubular (centronuclear) myopathy; mitochondrial myopathy; familial periodic paralysis; inflammatory myopathy; metabolic

myopathy, such as caused by a glycogen or lipid storage disease; dermatomyositis; polymyositis; inclusion body myositis; myositis ossificans; rhabdomyolysis and myoglobinurias.

The myopathy may be caused by a muscular dystrophy syndrome, such as Duchenne, Becker,  
5 myotonic, fascioscapulohumeral, Emery-Dreifuss, oculopharyngeal, scapulohumeral, limb girdle, Fukuyama, a congenital muscular dystrophy, or hereditary distal myopathy. The musculoskeletal disease can also be osteoporosis, a bone fracture, short stature, or dwarfism.

Other pathological disorders that can lead to the loss of muscle function are adult motor neuron  
10 disease, infantile spinal muscular atrophy, amyotrophic lateral sclerosis, juvenile spinal muscular atrophy, autoimmune motor neuropathy with multifocal conductor block, paralysis due to stroke or spinal cord injury, skeletal immobilization due to trauma, prolonged bed rest, voluntary inactivity, involuntary inactivity, metabolic stress or nutritional insufficiency, cancer, AIDS, fasting, a thyroid gland disorder, diabetes, benign congenital hypotonia, central core disease, burn injury, chronic  
15 obstructive pulmonary disease, liver diseases (examples such as fibrosis, cirrhosis), sepsis, renal failure, congestive heart failure, ageing, space travel or time spent in a zero gravity environment.

Examples of age-related conditions that may be treated include, sarcopenia, skin atrophy, muscle wasting, brain atrophy, atherosclerosis, arteriosclerosis, pulmonary emphysema, osteoporosis,  
20 osteoarthritis, immunologic incompetence, high blood pressure, dementia, Huntington's disease, Alzheimer's disease, cataracts, age-related macular degeneration, prostate cancer, stroke, diminished life expectancy, frailty, memory loss, wrinkles, impaired kidney function, and age-related hearing loss; metabolic disorders, including Type II Diabetes, Metabolic Syndrome, hyperglycemia, and obesity. Of course, patients may simultaneously suffer from one or more of these conditions,  
25 for example, sarcopenia and pulmonary emphysema, or sarcopenia and impaired kidney function.

Other conditions that are considered to be "pathological disorders" as recited herein include acute and/or chronic renal disease or failure, liver fibrosis or cirrhosis, cancer such as breast cancer, Parkinson's Disease; conditions associated with neuronal death, such as ALS, brain atrophy, or  
30 dementia and anemia. In addition, there are losses suffered as a consequence of age, trauma or inactivity.

Further conditions include cachexia, cachexia associated with a rheumatoid arthritis and cachexia associated with cancer.

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To date, very few reliable or effective therapies have been developed to treat these disorders. Individual aspects of muscular problems have been addressed. For example, one potential avenue of treatment for loss of muscle mass is the inhibition of myostatin. Myostatin, sometimes referred to as GDF-8 (growth differentiation factor-8), is one of a family of dimeric growth and differentiation  
40 factors which belong to the transforming growth factor-beta (TGF-beta) superfamily of structurally

related signalling proteins. These proteins signal through a heterodimeric complex of receptor serine kinases which include at least two type I receptors, ActRIB (ALK4) and ActRIC (ALK7) and two type II receptors, ActRIIA (ACVR2A) and ActR2B (ACVR2B). These receptors are all transmembrane proteins, composed of a ligand-binding extracellular domain with cysteine-rich region, a transmembrane domain, and a cytoplasmic domain with predicted serine/threonine specificity. Type I receptors are essential for signalling while type II receptors are required for binding ligands and for expression of type I receptors. Type I and II receptors form a stable complex after ligand binding resulting in the phosphorylation of type I receptors by type II receptors.

10 The activin receptor II B (ActR2B) is a receptor for myostatin (GDF8) but numerous other members of the TGF-beta super family, such as Activin B, Activin AB, Inhibin A, Inhibin B, GDF3, GDF1 1 , Nodal, BMP2, BMP4, BMP7, BMP9, and BMP10 bind to and activate ActR2B as well (see, for example Tsuchida *et al* (Endocrine journal 2008 55(1), 11-21). Blocking the interactions of ActR2B with its ligands can lead to beneficial physiological effects. The interaction between myostatin and this receptor regulates the inhibition of skeletal muscle differentiation via the Smad-dependent pathway. (SMADs are intracellular proteins that transduce extracellular signals from transforming growth factor beta ligands, like myostatin to the nucleus where they activate downstream gene transcription). Thus, by inhibiting or preventing myostatin from binding to ActR2B, one can induce the formation of skeletal muscle.

20

Various groups have looked into this. Bogdanovich *et al* (Nature, 2002, 420:418-421) describes that anti-myostatin antibodies were able to block myostatin, resulting in an increase in muscle mass in a mouse model of Duchenne muscular dystrophy. Bradley *et al* (Cell Mol. Life. Sci. 2008, 65:2119-2124) have reviewed the different available approaches for modulating the myostatin/ActR2B interaction, including the aforementioned anti-myostatin antibodies, inhibiting the release of mature myostatin by administering the myostatin propeptide, administering follistatin to block the myostatin receptor, administering HDAC inhibitors to induce follistatin production, administering an altered myostatin peptide which prevents myostatin from binding the receptor and administering a soluble decoy receptor for myostatin.

30

Myostatin acts to inhibit muscle fibre growth and muscle stem cell growth. Studies have shown that animals either lacking myostatin or treated with substances that block the activity of myostatin have significantly larger muscle mass. Furthermore, individuals who have mutations in both copies of the myostatin gene have significantly increased muscle mass and in many cases are stronger than normal. In a particular case, described in N Engl J Med 2004; 350:2682-2688 (June 24, 2004), a baby born with unusually muscular development turned out to have a deficiency of myostatin, believed due to a defect in the gene responsible for its production.

A number of recent publications have sought to exploit this as a treatment, for example:

40

- Myostatin binding proteins such as ActR2B polypeptides (e.g. US7842663 and US8614292),
- follistatin (e.g. US6004937)
- anti-ActR2B antibodies (e.g. US 8,551,482)
- anti-myostatin antibodies (e.g., US7261893, US8063188, US8415159, US8551482 and US8710212)
- anti-activin antibodies (e.g. US 20150037339 A1, WO 2009137075 A1)

Muscular hypertrophy and/or atrophy is not the whole story. Another factor is denervation.

Muscle contraction is triggered by impulse from the nerves through release of electrical and chemical messages. Nerves also provide the muscle fibers with a number of trophic factors, essential to the well-being and proper functioning of the muscle. The connection between nerves and muscle occurs via a highly complex synaptic structure called the neuromuscular junction (NMJ). Loss of NMJs results in the reduction of muscular function, independently of the anatomical and biochemical integrity of the muscle itself.

Some approaches have focussed on preventing loss of neuro-muscular junctions. For correct maintenance of NMJs, it has been shown by, for example, Wu *et al* (Development. 2010 Apr;137(7):1017-33. doi: 10.1242/dev.038711) and Tezuka *et al* (Proc Natl Acad Sci U S A. 2014 Nov 18;111(46):16556-61) that agrin is important for their formation and maintenance of a neuromuscular junction.

It has nevertheless been found that even the combination of increased muscle mass and treatment seeking to retain NMJs does not prevent the loss of muscle performance (measured by the performance of suitably treated mice on a treadmill). In short, while there are a number of methods for improving individual aspects of muscle problems, there currently exists nothing that can counteract the overall loss of muscle performance.

It has now been found that a particular group of linked proteins, polypeptides and monoclonal antibodies has given exceptional results in improving muscle performance. Therefore, there is provided a compound comprising at least two components, a first

component being the nLG3 domain from the C-terminus of human or mouse agrin, and at least one second component, selected from proteins or an antagonistic antibody that inhibit ActR2B-induced signaling activity in the presence of myostatin, the components being linked by means of linking entities.

5

There is additionally provided a method of improving muscle performance, comprising the administration of an effective amount of a compound as hereinabove defined.

10

By "improving muscle performance" is meant that muscular endurance especially is improved. It is a surprising fact that, although the compounds hereinabove defined do not provide much more muscle mass (myostatin) nor do they cluster acetyl choline receptors efficiently (at least 100-1000



times lower than a fully active agrin fragment), the overall muscular performance as measured by the endurance is remarkably improved. This can be demonstrated by the experimental methods hereinafter described.

- 5 With respect to amino acid sequences suitable for use in, and described in, this disclosure, the sequences should be at least 75% identical to those set forth hereinunder. More particularly, they may be 80%, 85%, 90% or 95% identical, most particularly 95% identical. In the case of inserts in the agrin (described in detail hereinunder), the inserts should be at least 95% identical. In a particular embodiment, all amino acid sequences are at least 95% identical.

10

The first component is the nLG3 domain from the C-terminus of agrin. Agrin is a large heparan proteoglycans with a molecular weight of 400-600 kDa. (Database accession number NP—940978). The protein core consists of about 2000 amino acids and its mass is about 225 kDa. It is a multidomain protein composed of 9 K (kunitz-type) domains, 2 LE (laminin-EGF-like) domains,  
 15 one SEA (sperm protein, enterokinase and agrin) domain, 4 EG (epidermal growth factor-like) domains and 3 LG (laminin globular) domains. Agrin is a very important protein and agrin-deficient mice die at birth due to respiratory failure. This is caused by the fact that agrin is strictly required for the proper innervation of muscle fibers and that these mice are not able to build proper NMJs.

- 20 Agrin exists in several splice variants and can be expressed as a secreted protein, containing the N-terminal NtA (N-terminal agrin) domain, which is the most abundant form of agrin and the predominant form expressed in motor neurons. It is produced in the soma of the neurons, transported down the axon and released from the axon ending of the motor nerve into the synaptic cleft of the NMJ. Here it acts as an agonist of LRP4 and may also become a component of the  
 25 basal lamina. In the CNS, most agrin is expressed as a type-II transmembrane protein by alternative splicing at the N-terminus lacking the N-terminal NtA domain (Bezakova and Ruegg, 2003).

- The serine/threonine (S/T) rich segments in agrin are responsible for a high degree of  
 30 glycosylation, containing several glycosylation and glucosaminoglycan attachment sites giving rise to the big mass of the proteoglycan. The C-terminal, 75 kDa moiety of agrin starting with the first EG domain, is required for full activity in acetylcholine receptor (AChR) clustering activity on muscle cells, although the most C-terminal 20 kDa fragment is sufficient to induce AChR aggregation (Bezakova and Ruegg, 2003). Several binding sites for interaction partners of agrin,  
 35 including  $\alpha$ -dystroglycan, heparin, some integrins and LRP4, are mapped to the C-terminal region. The large heparansulfate side chains are binding sites for heparin binding proteins, e.g some growth factors.

- In the C-terminal part of human agrin, there are 2 alternative splice sites y and z. At the y-site, there  
 40 may be inserts of 0, 4, 17 or 21 (4+17) amino acids and at the z site there may be inserts of 0, 8,

11 or 19 (8+11) amino acids. The function of the four inserted amino acids in the  $\gamma$ -site is to create a heparin binding site. Motor neurons express predominantly  $\gamma 4$  agrin. The most important splice site of agrin in respect of NMJ maturation is the  $z$ -site, giving agrin the ability to be active as an acetylcholine-receptor clustering agent. It is well known that full-length agrin containing the insertion of 8 amino acids at the  $z$ -site in presence of the 4-amino acid insert in splice site  $\gamma$  ( $\gamma 4z8$ ) generates an agrin variant with a half maximal AChR clustering activity of 35 pM in cultured myotube clustering assays. The insertion of 11 amino acids gives rise to a half-maximal AChR clustering activity while the 19 amino acid insertion results in a half-maximal AChR clustering activity of 110 pM. Agrin without an insertion at this site is not active in clustering acetylcholine-receptors on the in-vitro cultured myotubes (Bezakova and Ruegg, 2003). Thus, the most active form of agrin in the clustering assay is the  $\gamma 4z8$  variant, which is expressed by motor neurons.

A 40 kDa C-terminal fragment of agrin ( $\gamma 4z8$ ) containing the LG2, EG4 and the LG3 domains was found to be active in AChR clustering with an EC<sub>50</sub> of 130 pM in the AChR clustering activity while shorter fragments have only lower activities. The C-terminal LG3 domain with the  $z8$  insertion, the so-called LG3 $z8$  domain, exhibits a half maximal AChR clustering activity of only 13 nM, which is a factor 100 fold lower than the 40 kDa fragment (Bezakova and Ruegg, 2003).

During the development and maturation of the NMJ, agrin is a key player of molecules involved in the clustering of acetylcholine receptors. While NMJs are destabilized by the neurotransmitter acetyl choline, agrin, which is secreted by the motor neuron, stabilizes and increases the clusters of the AChR's via phosphorylation of MuSK, a membrane bound receptor tyrosine kinase. The interaction of agrin with MuSK is postulated to be mediated via LRP4, a low-density lipoprotein receptor (LDLR)-related protein. It was found that agrin ( $\gamma 4z8$ ) has a 10-fold higher affinity to LRP4 than agrin ( $\gamma 4z0$ ) giving rise to the differential AChR clustering activity of the different agrin splice variants observed in the *in vitro* cultured myotube assays. Upon agrin binding, LRP4 causes self-phosphorylation of MuSK, which then activates the signal cascade for the expression and clustering of acetylcholine receptors. It

has been shown that a 44-kD fragment of agrin leads to the formation of clusters of acetyl choline receptors on the surface of muscle cells (see Hettwer *et al* (PLOS ONE February 2014. Vol.9, Issue 2, e88739)), which is believed to be the initial step in the formation of a NMJ.

- 5 The term "LG3" as used in this disclosure means the human- or mouse-derived 22 kDa C-terminal agrin fragment of SEQ ID NO: 1 (all sequences are appended to this disclosure and form part thereof). The term "nLG3" as used in this disclosure means the LG3 fragment which further contains an insertion of 8, 11 or 19 amino acids at the z-site. The inserted sequences at the z-site are ELTNEIPA (z8, SEQ ID NO: 2), PETLDSRALFS (z11, SEQ ID NO: 3) or
- 10 ELTNEIPAPETLDSRALFS (z19, SEQ ID NO: 4, a combination of SEQ ID NO: 2 and SEQ ID NO: 3). An example of nLG3 is SEQ ID NO: 5

The term "(h)LG3" as used in this disclosure means the human derived 22 kDa C-terminal agrin fragment of SEQ ID NO: 6. The term "(h)nLG3" as used in this disclosure means the (h)LG3 fragment which further contains an insertion of 8, 11 or 19 amino acids at the z-site. The inserted sequences at the z-site ELANEIPV (z8, SEQ ID NO: 7), PETLD SGALHS (z11, SEQ ID NO: 8) or  
 5 ELANEIPVPETLD SGALHS (z19, SEQ ID NO: 9, a combination of SEQ ID NO: 7 and SEQ ID NO: 9). A particular example of (h)nLG3 is SEQ ID NO: 10

nLG3 may include additional amino acids at the N-terminus or C-terminus. Such additional amino acids at the N-terminus are e.g. present due to the method of preparation by recombinant synthesis  
 10 and expression in suitable cells.

Proteins containing elongations at the N-terminus by one or more domains of agrin up to the natural N-terminus of agrin are also included, as well as glycosylated or in other ways post-translationally, enzymatically or chemically modified protein variants of human agrin.  
 15

The second component is selected from proteins or antagonistic antibodies that inhibit ActR2B-mediated signaling activity in the presence of myostatin. Examples of a second component as used herein refers to a protein or an antagonistic antibody such as actR2B (AcvRIIB, actRIIB), or acvRA (actR2, actRII), alk4, alk5. The term ActR refers to soluble extracellular part of the mouse ActR2B  
 20 receptor as defined in SEQ ID NO: 11. This extracellular part is any part of the transmembrane protein that projects into the environment surrounding a cell. The term (h)ActR refers to extracellular part of the human ActR2B receptor as defined in SEQ ID NO: 12 (AAC64515.1, GI:3769443). Another example is follistatin as defined in SEQ ID NO: 25.

25 An example of a monoclonal antibody (mAb) that inhibits ActR2B-mediated signaling activity is ActRmAb (US8551482). Antibodies consist of a light chain (LC) and a heavy chain (HC). A typical example of the LC of ActRmAb (ActRmAb(LC)) is as defined in SEQ ID NO: 29. The HC of ActRmAb (ActRmAb(HC)) is as defined in SEQ ID NO: 28. The (h)nLG3 connected to ActRmAb(HC) (ActRmAb(HC)-(h)nLG3) is defined in SEQ ID NO: 30 Another example of an  
 30 antibody is MyomAb (US8063188). This antibody is directed against myostatin and prevents binding of myostatin to the (h)ActR receptor. A typical example of the LC of MyomAb (MyomAb(LC)) is as defined in SEQ ID NO: 32. The HC of MyomAb (MyomAb(HC)) is as defined in SEQ ID NO: 31. The (h)nLG3 connected to MyomAb(HC) (MyomAb(HC)-(h)nLG3) is defined in SEQ ID NO: 33.

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This ActR2B-mediated signal-inhibiting activity of mAbs may be readily ascertained by means of an assay. Such an assay can include, for example, a Smad-dependent reporter gene assay, inhibition of myostatin-induced Smad phosphorylation (P-Smad ELISA) and inhibition of myostatin-induced inhibition of skeletal muscle cell differentiation (for instance by a creatine kinase assay).

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In some embodiments, the second component inhibits myostatin-induced signaling as measured in a Smad-dependent reporter gene assay at an IC<sub>50</sub> of 10 nM or less, 1 nM or less, or 100 pM or less.

- 5 In some instances, it is possible for a compound according to this disclosure to comprise an additional component, meaning that there will be three components, joined by two linking entities. The third component acts as a stabiliser component, that is, it increases *in vivo* serum half-life. This may be as a result of decreased destruction, decreased clearance by the kidney, or other pharmacokinetic effect. "*in vivo* serum half-life" refers to the half-life of a protein circulating in the
- 10 blood of an organism. Fusions with the Fc region of an immunoglobulin (IgG molecule) are known to confer desirable pharmacokinetic properties and increase serum half-life on a wide range of proteins. The term "Fc region of an IgG molecule" refers to the Fc domain of an immunoglobulin of the isotype IgG, as is well known to those skilled in the art. The Fc region of an IgG molecule is that portion of IgG molecule (IgG1, IgG2, IgG3, and IgG4) that is responsible for increasing the *in vivo*
- 15 serum half-life of the IgG molecule.

The third component may also be selected so as to confer a desired property. For example, some domains are particularly useful for isolation of the resulting proteins by affinity chromatography. For the purpose of affinity purification, relevant matrices for affinity chromatography, such as

20 glutathione-, amylase-, and nickel- or cobalt-conjugated resins are used. Many of such matrices are available in "kit" form, such as the Pharmacia GST purification system and the QIAexpress.TM. system (Qiagen) useful with (HIS.sub.6) fusion partners. As another example, the third domain may be selected so as to facilitate detection of the ActR2B polypeptides. Examples of such detection domains include the various fluorescent proteins (e.g., GFP) as well as "epitope tags," which are

25 usually short peptide sequences for which a specific antibody is available. Well known epitope tags for which specific monoclonal antibodies are readily available include FLAG, influenza virus haemagglutinin (HA), and c-myc tags. In some cases, the third domain might have a protease cleavage site, such as for Factor Xa or Thrombin, which allows the relevant protease to partially digest the fusion proteins and thereby liberate the recombinant proteins therefrom. The liberated

30 proteins can then be isolated from the third domain by subsequent chromatographic separation. In certain preferred embodiments, the ActR domain and the (h)nLG3 domain are linked to a domain that stabilizes the resulting polypeptide *in vivo*.

A typical example of a third component is an "Fc" domain SEQ ID NO: 14.

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Likewise, fusions to human serum albumin can confer desirable properties. Other types of fusion domains that may be selected include multimerizing (e.g., dimerizing, tetramerizing) domains and functional domains (that confer an additional biological function, such as further stimulation of muscle growth).

40

A linking entity is a short stretch of amino acids, linking two protein components. This unstructured linker may correspond to the roughly 15 amino acid unstructured region at the C-terminal end of the extracellular domain of ActR2B (the "tail"), or it may be an artificial sequence of between 5 and 15, 20, 30, 50 or more amino acids that are relatively free of secondary structure. A linker may be rich in glycine and proline residues and may, for example, contain repeating sequences of threonine/serine and glycines. Often multiple repeats of the sequence gggg are used (Glycin-Glycin-Glycin-Serin). Glycin gives flexibility and Serin is polar. A linking may be rich in glycine and proline residues and may, for example, contain repeating sequences of threonine/serine and glycines (e.g., TG.sub.4 or SG.sub.4 repeats). A typical linker sequence "L" is as defined in SEQ ID NO: 13. A fusion protein may include a purification subsequence, such as an epitope tag, a FLAG tag, a polyhistidine sequence, and a GST fusion.

Examples of ActR domain containing protein linked to a Fc domain are ActR-Fc (SEQ ID NO: 19), Fc-ActR (SEQ ID NO: 20) and (h)ActR-Fc (SEQ ID NO: 21). (Cadena et al. J Appl Physiol 109: 635–642, 2010). Example of a follistatin linked to an Fc domain is defined in SEQ ID NO: 26.

Examples of nLG3 domain-containing protein linked to a Fc domain are nLG3-Fc (SEQ ID NO: 16), Fc-nLG3 (SEQ ID NO: 15) and Fc-(h)nLG3 (SEQ ID NO: 17). An example of a LG3 domain without insert Fc-(h)LG3 (SEQ ID NO: 18) was also constructed

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Particular embodiments include the following components

- A-L-B
- B-L-A
- A-L-C-L-B
- 25 B-L-C-L-A
- C-L-B-L-A
- C-L-A-L-B
- B-L-A-L-C
- A-L-B-L-C
- 30 A-L-D
- D-L-A
- A-L-E
- E-L-A

in which

- 35 A represents an agrin nLG3 domain-containing protein,
- B represents an extracellular domain of the ActRIB, ActRIC, ActRIIA, ActR2B receptor protein and follistatin
- C represents a "stabiliser" domain,
- D represents ActR2B-mediated signal-inhibiting mAb against the ActRIB, ActRIC, ActRIIA, and
- 40 ActR2B receptor protein,

E represents ActR2B-mediated signal-inhibiting mAb against a member of the TGF-beta super family

L represents a linking entity.

5 particular combinations being B-L-C-L-A, D-L-A and E-L-A

Examples of these particular combinations are ActR-Fc-nLG3 as defined in SEQ ID NO: 22, (h)ActR-Fc-(h)nLG3 as defined in SEQ ID NO: 23. Fol-Fc-nLG3 as defined in SEQ ID NO: 27. The ActRmAb(HC) linked to (h)nLG3 (ActRmAb(HC)-(h)nLG3) as defined in SEQ ID NO: 30. The  
10 MyomAb(HC linked to (h)nLG3) (MyomAb(HC)-(h)nLG3) as defined in SEQ ID NO: 33. The corresponding light chains of the antibodies need to be co-expressed with the heavy chains to create a fully functional antibody.

The heavy and light chains of the antibodies according to this disclosure may be expressed as  
15 contiguous single-chain proteins, with the first and second components joined by the linking entity (see e.g. Bird et al., 1988 Science 242:423-426; Huston et al., 1988 Proc. Natl. Acad. Sci. USA 85:5879-5883; McCafferty et al., 1990 Nature 348:552-554). The contiguous single chain protein can be linked to (h)nLG3.

20 The compounds of this disclosure may be made by known methods. For example, the DNA coding for the compound is expressed in suitable expression systems and the resulting protein is subsequently purified. Several prokaryotic and eukaryotic expression systems are suitable for the production and secretion of the compounds of the disclosure. Prokaryotic expression systems include, but are not limited to, expression in E. coli. Eukaryotic expression systems include  
25 expression in mouse myeloma cells, baculovirus-mediated expression in insect cells, as well as expression in human embryonic kidney (HEK) cells, transient expression in Chinese hamster ovary (CHO) cells and stable expression in Pichia pastoris. These systems have the advantage that they can easily be adapted to serum-free conditions to reduce the amount of contaminating proteins in the supernatant and can be adapted for large scale production. In addition, a variety of cell lines  
30 may be used, including HEK293T and HEK293-cells, COS cells, CHO cells, HeLa cells, H9 cells, Jurkat cells, NIH3T3 cells, C127 cells, CV1 cells, CAP cells or SF cells.

The sequence of a component may be adjusted, as appropriate, depending on the type of  
35 expression system used, as mammalian, yeast, insect and plant cells may all introduce differing glycosylation patterns that can be affected by the amino acid sequence of the peptide. In general, proteins for use in humans will be expressed in a mammalian cell line that provides proper glycosylation, such as HEK293 or CHO cell lines, although other mammalian expression cell lines are also expected to be useful.

The compounds of the disclosure may be purified by standard protein purification technologies. Immunoglobulins G may be purified using protein A or G. His-tagged protein can be purified using IMAC, but ion exchange chromatography or affinity purification using a heparin column can be used as well. Purification via an antibody raised against the C-terminal part of agrin can also be  
5 used. The eluted protein can then further be purified using a hydroxyapatite column or by gel filtration.

The compounds of this disclosure are useful in pharmaceutical compositions. The disclosure therefore provides a pharmaceutical composition comprising at least one compound as  
10 hereinabove described, formulated together with a pharmaceutically-acceptable carrier.

The pharmaceutical compositions of this disclosure are particularly useful for the treatment of pathological conditions leading to the loss of muscle function. Non-limiting examples of such conditions include:

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- muscle atrophy as a result of treatment with a glucocorticoid such as cortisol, dexamethasone, betamethasone, prednisone, methylprednisolone, or prednisolone;
- muscle atrophy as a result of denervation due to nerve trauma or a result of degenerative, metabolic, or inflammatory neuropathy (e.g., Guillian-Barre syndrome, peripheral neuropathy,  
20 or exposure to environmental toxins or drugs);
- muscle atrophy as a result of myopathy, such as myotonia; a congenital myopathy, including nemaline myopathy, multi/minicore myopathy and myotubular (centronuclear) myopathy; mitochondrial myopathy; familial periodic paralysis; inflammatory myopathy; metabolic myopathy, such as caused by a glycogen or lipid storage disease; dermatomyositis;  
25 polymyositis; inclusion body myositis; myositis ossificans; rhabdomyolysis and myoglobinurias;
- myopathy caused by a muscular dystrophy syndrome, such as Duchenne, Becker, myotonic, fascioscapulohumeral, Emery-Dreifuss, oculopharyngeal, scapulohumeral, limb girdle, Fukuyama, a congenital muscular dystrophy, or hereditary distal myopathy
- musculoskeletal diseases such as osteoporosis, bone fracture, short stature, or dwarfism;  
30
- adult motor neuron disease, infantile spinal muscular atrophy, amyotrophic lateral sclerosis, juvenile spinal muscular atrophy, autoimmune motor neuropathy with multifocal conductor block, paralysis due to stroke or spinal cord injury, skeletal immobilization due to trauma, prolonged bed rest, voluntary inactivity, involuntary inactivity, metabolic stress or nutritional  
35 insufficiency, cancer, AIDS, fasting, a thyroid gland disorder, diabetes, benign congenital hypotonia, central core disease, burn injury, chronic obstructive pulmonary disease, liver diseases (examples such as fibrosis, cirrhosis), sepsis, renal failure, congestive heart failure, ageing, space travel or time spent in a zero gravity environment.



- age-related conditions such as sarcopenia, skin atrophy, muscle wasting, brain atrophy, atherosclerosis, arteriosclerosis, pulmonary emphysema, osteoporosis, osteoarthritis, immunologic incompetence, high blood pressure, dementia, Huntington's disease, Alzheimer's disease, cataracts, age-related macular degeneration, prostate cancer, stroke, diminished life expectancy, frailty, memory loss, wrinkles, impaired kidney function, and age-related hearing loss; metabolic disorders, including Type II Diabetes, Metabolic Syndrome, hyperglycemia, and obesity. Of course, patients may simultaneously suffer from one or more of these conditions, for example, sarcopenia and pulmonary emphysema, or sarcopenia and impaired kidney function.
- pathological disorders such as acute and/or chronic renal disease or failure, liver fibrosis or cirrhosis, cancer such as breast cancer, Parkinson's Disease; conditions associated with neuronal death, such as ALS, brain atrophy, or dementia and anemia. In addition, there are losses suffered as a consequence of age, trauma or inactivity.
- further conditions such as cachexia, cachexia associated with a rheumatoid arthritis and cachexia associated with cancer.

The pharmaceutical compounds of the disclosure can also be administered in combination therapy, i.e. combined with other agents. For example, the combination therapy can include an anti-ActR2B antibody of the present disclosure combined with at least one other muscle mass/strength increasing agent, for example, IGF-1, IGF-2 or variants of IGF-1 or IGF-2, an anti-myostatin antibody, a myostatin propeptide, a myostatin decoy protein that binds ActR2B but does not activate it, a beta 2 agonist, a Ghrelin agonist, a SARM, GH agonists/mimetics or follistatin. The pharmaceutical compounds of the disclosure also can be administered in combination therapy with Nusinersen or similar compounds. Nusinersen, an antisense oligonucleotide that modulates alternate splicing of the SMN2 gene, functionally converting it into SMN1 gene, is an investigational drug for spinal muscular atrophy.

As used herein, "pharmaceutically acceptable carrier" includes any and all solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like that are physiologically compatible. The carrier should be suitable for intravenous, intramuscular, subcutaneous, parenteral, spinal or epidermal administration (e.g. by injection or infusion). Depending on the route of administration, the compound of this disclosure may be coated in a material to protect the compound from the action of acids and other natural conditions that may inactivate the compound.

The compounds of the disclosure may be in the form of pharmaceutically-acceptable salts. A "pharmaceutically acceptable salt" refers to a salt that retains the desired biological activity of the parent compound and does not impart any undesired toxicological effects (see e.g. Berge, S. M., et

al., 1977 J. Pharm. Sci. 66:1-19). Examples of such salts include acid addition salts and base addition salts. Acid addition salts include those derived from nontoxic inorganic acids, such as hydrochloric, nitric, phosphoric, sulfuric, hydrobromic, hydroiodic, phosphorous and the like, as well as from nontoxic organic acids such as aliphatic mono- and di-carboxylic acids, phenyl-substituted  
5 alkanolic acids, hydroxy alkanolic acids, aromatic acids, aliphatic and aromatic sulfonic acids and the like. Base addition salts include those derived from alkaline earth metals, such as sodium, potassium, magnesium, calcium and the like, as well as from nontoxic organic amines, such as N,N'-dibenzylethylenediamine, N-methylglucamine, chlorprocaine, choline, diethanolamine, ethylenediamine, procaine and the like.

10

A pharmaceutical composition of the disclosure also may include a pharmaceutically acceptable anti-oxidant. Examples of pharmaceutically acceptable antioxidants include: water soluble antioxidants, such as ascorbic acid, cysteine hydrochloride, sodium bisulfate, sodium metabisulfite, sodium sulfite and the like; oil-soluble antioxidants, such as ascorbyl palmitate, butylated  
15 hydroxyanisole (BHA), butylated hydroxytoluene (BHT), lecithin, propyl gallate, alpha-tocopherol, and the like; and metal chelating agents, such as citric acid, ethylenediamine tetraacetic acid (EDTA), sorbitol, tartaric acid, phosphoric acid, and the like.

Examples of suitable aqueous and nonaqueous carriers that may be employed in the  
20 pharmaceutical compositions of the disclosure include water, ethanol, polyols (such as glycerol, propylene glycol, polyethylene glycol, and the like), and suitable mixtures thereof, vegetable oils, such as olive oil, and injectable organic esters, such as ethyl oleate. Proper fluidity can be maintained, for example, by the use of coating materials, such as lecithin, by the maintenance of the required particle size in the case of dispersions, and by the use of surfactants.

25

These compositions may also contain adjuvants such as preservatives, wetting agents, emulsifying agents and dispersing agents. Prevention of presence of microorganisms may be ensured both by sterilization procedures, supra, and by the inclusion of various antibacterial and antifungal agents, for example, paraben, chlorobutanol, phenol sorbic acid, and the like. It may also be desirable to  
30 include isotonic agents, such as sugars, sodium chloride, and the like into the compounds. In addition, prolonged absorption of the injectable pharmaceutical form may be brought about by the inclusion of agents which delay absorption such as, aluminum monostearate and gelatin.

Pharmaceutically-acceptable carriers include sterile aqueous solutions or dispersions and sterile  
35 powders for the extemporaneous preparation of sterile injectable solutions or dispersion. The use of such media and agents for pharmaceutically active substances is known in the art. Except insofar as any conventional media or agent is incompatible with the active compound, use thereof in the pharmaceutical compounds of the disclosure is contemplated. Supplementary active compounds can also be incorporated into the compounds.

40

Therapeutic compositions typically must be sterile and stable under the conditions of manufacture and storage. The compound can be formulated as a solution, microemulsion, liposome, or other ordered structure suitable to high drug concentration. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyethylene glycol, and the like), and suitable mixtures thereof. The proper fluidity can be maintained, for example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. In many cases, one can include isotonic agents, for example, sugars, polyalcohols such as mannitol, sorbitol, or sodium chloride in the compound. Prolonged absorption of the injectable compounds can be brought about by including in the compound an agent that delays absorption for example, monostearate salts and gelatin.

Sterile injectable solutions can be prepared by incorporating the active compound in the required amount in an appropriate solvent with one or a combination of agents enumerated above, as required, followed by sterilization microfiltration. Generally, dispersions are prepared by incorporating the active compound into a sterile vehicle that contains a basic dispersion medium and the required other agents from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the methods of preparation are vacuum drying and freeze-drying (lyophilization) that yield a powder of the active agent plus any additional desired agent from a previously sterile-filtered solution thereof.

The amount of compound which can be combined with a carrier material to produce a single dosage form will vary depending upon the subject being treated, and the particular mode of administration. The amount of active agent which can be combined with a carrier material to produce a single dosage form will generally be that amount of the compound which produces a therapeutic effect. Generally, out of one hundred percent, this amount will range from about 0.01 percent to about ninety-nine percent of active agent, from about 0.1 percent to about 70 percent, or from about 1 percent to about 30 percent of active agent in combination with a pharmaceutically acceptable carrier.

30

Dosage regimens are adjusted to provide the optimum desired response (e.g. a therapeutic response). For example, a single bolus may be administered, several divided doses may be administered over time or the dose may be proportionally reduced or increased as indicated by the exigencies of the therapeutic situation. It is especially advantageous to formulate parenteral compounds in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to physically discrete units suited as unitary dosages for the subjects to be treated; each unit contains a predetermined quantity of active compound calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. The specification for the dosage unit forms of the disclosure are dictated by and directly dependent on the unique characteristics of the active compound and the particular therapeutic effect to be

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achieved, and the limitations inherent in the art of compounding such an active compound for the treatment of sensitivity in individuals.

For administration of the compound, the dosage ranges from about 0.0001 to 100 mg/kg, and more usually 0.01 to 5 mg/kg, of the host body weight. For example dosages can be 0.3 mg/kg body weight, 1 mg/kg body weight, 3 mg/kg body weight, 5 mg/kg body weight or 10 mg/kg body weight or within the ranges of 1-10 mg/kg or 3-7 mg/kg. An example treatment regime entails administration once per week, once every two weeks, once every three weeks, once every four weeks, once a month, once every 3 months or once every three to 6 months. Alternatively, the compound may be administered about once a year or once only. Such administration may be carried out intravenously or subcutaneously. Dosage regimens for a compound of the disclosure include 1 mg/kg body weight or 3 mg/kg body weight by intravenous administration, with the antibody being given using one of the following dosing schedules: every four weeks for six dosages, then every three months; every three weeks; 3 mg/kg body weight once followed by 1 mg/kg body weight every three weeks.

The dosage should be one that causes an enhancement of muscle performance. In various embodiments the effect is on skeletal muscle. In various embodiments, the dosage causes muscle hypertrophy with no more than a proportional increase in the size of internal organs (e.g. heart, lungs, liver, kidneys). Such a proportional increase may be compared by measuring either mass or volume.

In some methods, two or more compounds according to this disclosure with different binding specificities may be administered simultaneously, in which case the dosage of each compound administered falls within the ranges indicated. Compound is usually administered on multiple occasions. Intervals between single dosages can be, for example, weekly, monthly, every three months, every six months or yearly. Intervals can also be irregular as indicated by measuring blood levels of compound to the target antigen in the patient. In some methods, dosage is adjusted to achieve a plasma antibody concentration of about 1-1000  $\mu\text{g/ml}$  and in some methods about 25-300  $\mu\text{g/ml}$ .

Alternatively, a compound can be administered as a sustained release formulation, in which case less frequent administration is required. Dosage and frequency vary depending on the half-life of the compound in the patient. In general, human antibodies show the longest half-life, followed by humanized antibodies, chimeric antibodies, and nonhuman antibodies. The dosage and frequency of administration can vary depending on whether the treatment is prophylactic or therapeutic. In prophylactic applications, a relatively low dosage is administered at relatively infrequent intervals over a long period of time. Some patients continue to receive treatment for the rest of their lives. In therapeutic applications, a relatively high dosage at relatively short intervals is sometimes required until progression of the disease is reduced or terminated or until the patient shows partial or

complete amelioration of symptoms of disease. Thereafter, the patient can be administered a prophylactic regime.

- Actual dosage levels of the compounds in the pharmaceutical compositions of the present disclosure may be varied so as to obtain an amount of the compound which is effective to achieve the desired therapeutic response for a particular patient, compound, and mode of administration, without being toxic to the patient. The selected dosage level will depend upon a variety of pharmacokinetic factors including the activity of the particular compounds of the present disclosure employed, or the ester, salt or amide thereof, the route of administration, the time of administration, the rate of excretion of the particular compound being employed, the duration of the treatment, other drugs, compounds and/or materials used in combination with the particular compounds employed, the age, sex, weight, condition, general health and prior medical history of the patient being treated, and like factors well known in the medical arts.
- 15 A "therapeutically effective dosage" of a compound of the disclosure can result in a decrease in severity of disease symptoms, an increase in frequency and duration of disease symptom-free periods, or a prevention of impairment or disability due to the disease affliction i.e. an increase in muscle mass and/or strength.
- 20 A compound of the present disclosure can be administered by one or more routes of administration using one or more of a variety of methods known in the art. As will be appreciated by the skilled artisan, the route and/or mode of administration will vary depending upon the desired results. Routes of administration for antibodies of the disclosure include intravenous, intramuscular, intradermal, intraperitoneal, subcutaneous, spinal or other parenteral routes of administration, for example by injection or infusion. The phrase "parenteral administration" as used herein means modes of administration other than enteral and topical administration, usually by injection, and includes, without limitation, intravenous, intramuscular, intraarterial, intrathecal, intracapsular, intraorbital, intracardiac, intradermal, intraperitoneal, transtracheal, subcutaneous, subcuticular, intraarticular, subcapsular, subarachnoid, intraspinal, epidural and intrasternal injection and infusion. In one embodiment the antibody is administered intravenously. In another embodiment the antibody is administered subcutaneously.
- 30

Alternatively, a compound of the present disclosure can be administered by a nonparenteral route, such as a topical, epidermal or mucosal route of administration, for example, intranasally, orally, vaginally, rectally, sublingually or topically.

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The active compounds can be prepared with carriers that will protect the compound against rapid release, such as a controlled release formulation, including implants, transdermal patches, and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides, polyglycolic acid, collagen, polyorthoesters, and polylactic

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acid. Many methods for the preparation of such formulations are patented or generally known to those skilled in the art. See, e.g. Sustained and Controlled Release Drug Delivery Systems, J. R. Robinson, ed., Marcel Dekker, Inc., New York, 1978.

5 The compounds can be administered with medical devices known in the art. For example, in one embodiment, a compound of the disclosure can be administered with a needleless hypodermic injection device, such as the devices shown in U.S. Pat. Nos. 5,399,163; 5,383,851; 5,312,335; 5,064,413; 4,941,880; 4,790,824 or 4,596,556. Examples of well-known implants and modules useful in the present disclosure include: U.S. Pat. No. 4,487,603, which shows an implantable  
10 micro-infusion pump for dispensing medication at a controlled rate; U.S. Pat. No. 4,486,194, which shows a therapeutic device for administering medicants through the skin; U.S. Pat. No. 4,447,233, which shows a medication infusion pump for delivering medication at a precise infusion rate; U.S. Pat. No. 4,447,224, which shows a variable flow implantable infusion apparatus for continuous drug delivery; U.S. Pat. No. 4,439,196, which shows an osmotic drug delivery system having multi-  
15 chamber compartments; and U.S. Pat. No. 4,475,196, which shows an osmotic drug delivery system. Many other such implants, delivery systems, and modules are known to those skilled in the art and include those made by MicroCHIPS.TM. (Bedford, Mass.).

In certain embodiments, the compounds of the disclosure can be formulated to ensure proper  
20 distribution *in vivo*. For example, the blood-brain barrier (BBB) excludes many highly hydrophilic compounds. To ensure that the compounds of the disclosure cross the BBB (if desired), they can be formulated, for example, in liposomes. For methods of manufacturing liposomes, see, e.g. U.S. Pat. Nos. 4,522,811; 5,374,548; and 5,399,331. The liposomes may comprise one or more moieties which are selectively transported into specific cells or organs, thus enhance targeted drug  
25 delivery (see, e.g. V. V. Ranade, 1989 J. Clin Pharmacol. 29:685). Example targeting moieties include folate or biotin (see, e.g. U.S. Pat. No. 5,416,016); mannosides (Umezawa et al., 1988 Biochem. Biophys. Res. Commun 153:1038); antibodies (P. G. Bloeman et al., 1995 FEBS Lett. 357:140; M. Owais et al., 1995 Antimicrob. Agents Chemother. 39:180); surfactant protein A receptor (Briscoe et al., 1995 Am. J. Physiol. 1233:134); p120 (Schreier et al., 1994 J. Biol. Chem.  
30 269:9090); see also K. Keinänen; M. L. Laukkanen, 1994 FEBS Lett. 346:123; J. J. Killian; I. J. Fidler, 1994 Immunomethods 4:273.

A further surprising and beneficial effect of the compounds of this disclosure is that they are much more specific myostatin inhibitors than those known to the art.

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It is well known that myostatin blocking agents have multiple effects, not only on muscle fibers but also on satellite cells. Satellite cells are a heterogeneous population of stem and progenitor cells that are required for the growth, maintenance and regeneration of skeletal muscle. Myostatin

blocking agents switch on growth and differentiation of these muscle stem cells (McCroskery et al. (2003) J. Cell Biol. 162, 1135–1147).

Myostatin blocking agents have a different effect on muscle fibers; they switch off protein  
 5 degradation of muscle filaments (responsible for muscle contraction) and turn on protein synthesis of muscle filaments (Curr Opin Support Palliat Care. 2011 Dec; 5(4): 334–341). Myosin is the name for a family of muscle protein filaments known for their role in muscle contraction – they comprise a family of ATP-dependent motor proteins and are known for their role in muscle contraction. During muscle contraction, muscle filaments, like myosin, can be damaged and need to be degraded and  
 10 replaced by new filaments. It is known that a human mutation in muscle protein degradation leads to proximal muscle weakness and hypertrophic cardiomyopathy. In a paper by Olivé *et al* (Human Molecular Genetics, 2015, 1–13) it was demonstrated that the muscle fibers of a patient contained inclusions formed by myosin and myosin-associated proteins.

15 Myostatin blocking agents also block protein degradation of muscle filaments and it is therefore feasible that prolonged exposure to a myostatin inhibitor leads to an accumulation of damaged muscle filaments. This could be an unwanted side effect of myostatin blocking agents such as ActR-Fc, ActRmAb and MyomAb. This may explain the relatively low activity of these proteins in performance assays such as the treadmill.

20

The coupling of nLG3 to ActR-Fc, ActRmAb and MyomAb described in this disclosure has resulted in novel compounds that are more specific in their mode of action. Proteins such as ActR-Fc-nLG3, ActRmAb-nLG3 and MyomAb-nLG3 activate only satellite cells and do not have a direct effect on muscle fibers. Presently, it is unclear how such proteins reach this new level of specificity in their  
 25 mode of action. One explanation, without limiting the scope of this disclosure in any way, could be that ActRIIB receptors use also LRP-proteins as co-receptor. It is very well known that nLG3 binds to LRP4. Binding to LRP4 is very important. The protein ActR-Fc-LG3, (note the difference between LG3 and nLG3) used as a control, leads to similar weight and muscle increase as ActR-Fc does, so the addition of a LG3 domain is not sufficient for this new activity (see figure 12). It is  
 30 essential that the LG3 domain has an appropriate insert which makes it capable of binding to LRP4. This result also shows that a bulky residue at the C-terminus of ActR-Fc is not likely to be the reason for the novel effects of ActR-Fc-nLG3.

This disclosure therefore also provides a method of specifically activating muscle satellite cells of  
 35 skeletal muscle in the absence of direct effect on muscle fibers, comprising the treatment of the muscle with a compound as hereinabove described.

The disclosure is further described with reference to the following examples and associated Figures, which depict particular embodiments and which are not in any way limiting.

A more detailed exposition of the Figures 1-11 is provided below, but the basic details are as follows:

**Figure 1** shows the formation of acetyl Choline receptor clusters (dots).

5 **Figure 2** shows the coomassie-stained SDS-PAGE gel of a number of compositions

**Figure 3** shows the relative body weight increase over time.

**Figure 4** shows the relative muscle weights for mice treated with vehicle and a number of compounds.

**Figure 5** shows the rotarod performance of treated mice.

10 **Figure 6** shows relative body weight increase over time.

**Figure 7** shows relative muscle wet weights.

**Figure 8** shows the treadmill performance of the aged mice.

**Figure 9** shows the number of motivational electrical pulses per minute during the treadmill runs.

**Figure 10** shows the mean grip strength (GS) performance of the mice during week three of  
15 vehicle dosing (GS before treatment).

**Figure 11** Summary of muscle pathology.

**Figure 12** shows the relative body weight increase over time.

### Synthesis of proteins

20 cDNAs were obtained commercially. The cDNAs were cloned via restriction enzymes NotI and HindIII into the mammalian gene expression vector pEvi3 (evitria AG, Switzerland). Plasmid DNA was prepared under low-endotoxin conditions using commercially-available DNA purification kits (Macherey Nagel, Germany). The protein Fc-nLG3 was obtained using SEQ ID NO: 34. The protein nLG3-Fc was constructed using SEQ ID NO: 35. Fc-(h)nLG3 was constructed using SEQ  
25 ID NO: 36. Fc-(h)LG3 was constructed using SEQ ID NO: 37. Fc-ActR was constructed using SEQ ID NO: 38, ActR-Fc was constructed using SEQ ID NO: 39. (h)ActR-Fc was constructed using SEQ ID NO: 40. ActR-Fc-nLG3 was constructed using SEQ ID NO: 41.

(h)ActR-Fc-(h)nLG3 was constructed using SEQ ID NO: 42, (h)ActR-Fc-(h)LG3 was constructed  
30 using SEQ ID NO: 43, ActRmAb(LC) was constructed using SEQ ID NO: 44 and ActRmAb(HC) was constructed using SEQ ID NO: 45. ActRmAb(HC)-(h)nLG3 was constructed using SEQ ID NO: 46. MyomAb(LC) was constructed using SEQ ID NO: 47, and MyomAb(HC) was constructed using SEQ ID NO: 48. MyomAb(HC)-(h)nLG3 was constructed using SEQ ID NO: 49. Fol-Fc and Fol-Fc-nLG3 were constructed using SEQ ID NO: 50 and SEQ ID NO: 51 respectively.

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### Production and Purification of the proteins

All proteins were produced by in CHO K1 cells. The seed was grown in eviGrow™ medium (evitria AG, Switzerland), a chemically defined, animal-component free, serum-free medium. Transfection and production were carried out in eviMake™ (evitria AG, Switzerland), an animal-component free,  
40 serum-free medium, at 37°C and 5% CO<sub>2</sub>. ActRmAb and MyomAb are generated by simultaneous



transfection with IgG heavy and light chain expression vector DNA. The resulting antibody are named ActmAb and MyomAb, respectively. ActRmAb-(h)nLG3 was made by simultaneous transfection of vector DNA generated with ActRmAb(HC)-(h)nLG3 and ActRmAb(LC). The resulting antibody is named ActmAb-(h)nLG3. MyomAb-(h)nLG3 was made by simultaneous transfection of  
5 vector DNA generated with MyomAb(HC)-(h)nLG3 and MyomAb(LC). The resulting antibody is named MyomAb-(h)nLG3.

Supernatants were harvested by centrifugation and sterile filtered (0.2 µm) at day 8 after transfection. The target proteins were subsequently purified via Protein A affinity chromatography  
10 on a Bio-Rad BioLogic DuoFlow FPLC system with PBS as wash buffer, 0.1 mol/l glycine pH 3.0 as elution buffer and 1 mol/l TRIS pH 10 as neutralization buffer.

#### Identification of proteins by SDS-PAGE Gel Electrophoresis

Each compound was eluted in 4X LDS Sample Buffer (Invitrogen) and 10X reducing agent  
15 (Invitrogen) to reach the concentration of 1µg. Samples were heated at 70°C for 10 minutes, and subsequently run on 4–12% Bis-Tris Plus gel (Invitrogen). Gels were run at 200V voltage for 35 minutes. Target protein fractions were identified by Coomassie staining of gel. Gels were left in Coomassie staining solution (0.26% Coomassie Blue, 10% Acetic Acid, 25% Methanol) for 4 hours. After removing Coomassie solutions, gels were then incubated in the destaining solution (10%  
20 Acetic Acid, 25% Methanol) overnight, in order to eliminate the excess dye. Gels were scanned and images were taken, using a densitometer (BioRad).

#### Acetyl Choline Receptor Clustering on C2C12 mouse cells.

C2C12 mouse muscle cells were cultured skeletal myoblasts from ATCC (ATCC-LGC Standards  
25 S.r.l., Italy) which were cultured in Dulbecco's Modified Eagle's Medium (DMEM) high glucose (Sigma, Italy) with 10% FBS (Sigma), containing 2 mM L-glutamine, 100 U/ml penicillin and 100 µg/ml streptomycin (all purchased from Invitrogen-Gibco). They were cultured for 2-3 days on 8 well chamber slides in the previous medium, then replaced with DMEM and 3% FBS, to obtain myotubes . Myotubes were incubated with agrin constructs at 10 µM (microM) for 24h and fixed  
30 with 2% paraformaldehyde for 20 min at RT. The samples were stained for the AChR by incubating the cells with Alexafluor 555-conjugated α-Bungarotoxin (1:500; Invitrogen, Italy) at RT for 1 h. The cells were then rinsed and coverslips were mounted with a drop of PB 0.1M. The levels of AChR clustering were compared by determining the average AChR cluster number in random fields, at a magnification of 40x with a fluorescent microscope.

#### 35 Animal studies

##### Ethic Statement

All procedures involving the use of laboratory animals were performed in accordance with the Italian national (DL n. 116, G.U., Supp. 40, February 18, 1992; permit number 17/2010-B, June 30, 2010) and European Communities Council Directive 24 November 1986 (86/609/EEC).

## Animals

### Nine week-old animals

In one experiment nine-week-old male C57BL/6 mice (n=5 per group, Harlan, Italy) were  
5 randomized with body weight and then treated subcutaneously with the proteins. The proteins used  
are indicated in the figures. Phosphate Buffered Saline (PBS), pH 7.4 was used as vehicle control.  
The dose was 10 mg/kg and is administrated three times per week, on day 1, 3, 5, 8, 10, 12 for a  
two week treatment. The total dose for the mixture was 20 mg/kg, consisting of a 1:1 mix of ActR-  
Fc and Fc-nLG3 so that each protein is given at 10 mg/kg. Body weights are determined three  
10 times per week prior to dosing 25 days after start of administration, mice are euthanized with CO<sub>2</sub>.  
Gastrocnemius, quadriceps femoris and triceps brachii are collected and weighed.

### 22 month-old animals

In the aged mice experiment 24 male mice, strain C57/BL6 (purchase at Charles River, France) are  
15 used. At the beginning of the experimental procedures, mice are 22 months old. Animals are  
weighed, ear punched; mice are kept in regular cages, 5 per cage, under 12/12-h light/dark cycle,  
with food and water available ad libitum. Injections were performed subcutaneously (10 mg/kg) 3  
times per week during five consecutive weeks. For the 5 week treatment the compounds were  
injected on day 1, 3, 5, 8, 10, 12, 15, 17, 19, 22, 24, 26, 29, 31, 33. Animals were split randomly  
20 into three experimental groups: control group (PBS), which received injections of PBS, treated  
groups AcrR-Fc-nLG3, and ActR-Fc.

### Body Weight

Mice body weight was measured 5 times per week throughout the experiment.  
25

### Rotarod.

Rotarod measurements were done on a 7650 accelerating model of a Rotarod™ apparatus (Ugo  
Basile, Italy)). The mouse is placed on the rod of a Rotarod. The rod slowly accelerates from 4 to  
32 rpm. The time that a mouse stays on the rod is recorded and the test terminates when the  
30 mouse is no longer able to remain on the rod. Maximum trial duration in the standard trial is 5  
minutes. In the extended trial the maximum trial duration is 30 minutes. Rotarod performance of  
different treatment groups is indicated. Standard deviations are indicated as error bars. (N = 5 for  
each group). The standard trial was done on day 18 and day 21 of the treatment after two exercise  
trials. The data is the average of four on two days. The extended trial is done on day 21 of the  
35 treatment.

### Treadmill Exercise

Mice were trained on a treadmill apparatus (Panlab, Harvard Apparatus) three times per week, in  
the afternoon. The instrument has the capability of exercising up to five mice simultaneously in  
40 individual lanes. Mice were trained on treadmill for 3 weeks before starting compound injections,

then for 3 weeks during compound/PBS dosing. Each mouse was tested using an accelerating treadmill protocol. Briefly, mice were properly acclimated to the treadmill prior to any experimentation. In the days before experimental runs, mice were placed on the treadmill in their respective lanes with shocking grids off and the belt moving and they were let to explore the instrument for minutes. During experimentation, mice were warmed up before running. For this, the belt was started at a low speed (16 cm/sec) and the shocking grids were gradually turned on to 0.2 mA. The duration of warm-up period was 2 minutes. After warm-up period, mice were tested for their running performances. The treadmill speed starts at 16 cm/sec and accelerated 1cm every minute. The acceleration continues until the mice reach exhaustion state. If a mouse received 10 or more shock per minute, this level is considered the exhaustion state and the experiment is stopped for that particular mouse. After exhaustion, shocking grid is deactivated and the mouse is returned to its cage. The running distance, the number of shocks taken in every minute, and the total number of shocks are evaluated for each mouse.

#### 15 **Grip Strength test**

Forelimb grip strength was measured using a Grip Strength Meter (Ugo Basile, Varese, Italy). The control and treated mice were tested twice a week during the first six weeks of the experiments, and were tested 5 times per week in the last two weeks of experimentation. Mice were held by the tail and allowed to grasp a T-shaped bar with their forepaws. Once the mouse grasped the bar with both paws, the mouse was pulled away from the bar until the mouse released the bar. The digital meter displays the level of tension (in grams) exerted on the bar by the mouse. Each animal was given five consecutive tests, the lowest and the highest values were excluded by the analysis, and the average value was taken.

#### 25 **Muscle isolation and storage**

Mice were sacrificed by cervical dislocation. After dislocation, the fresh skeletal muscles (triceps, quadriceps and gastrocnemius), were quickly dissected out from the skin and bones by forceps and scissors. The wet muscle weight was determined immediately after isolation. Then, muscles (3 for each mouse) were placed to the Peel-A-Way embedding molds (Sigma-Aldrich; E6032-1cs) containing Killik, embedding medium for criostate neutral (Bio-Optica, Milan; 05-9801), for cryosectioning. The minimal amount of Killik possible to cover the muscles was used, thus allowing rapid freezing to occur. Then the molds were immediately transferred in beaker filled with Isopentane (1-Methylbutane; Sigma-Aldrich; M32631) and dry ice (-80°C) for 20-40 seconds (longer contact times can result in the formation of cracks in the samples; insufficient time can result in freezing artifacts) and then were transferred the muscle sample to dry ice. For long-term storage samples were kept in freezer at -80°C.

The other three muscles for each mouse were quickly placed into a tube and covered with at least 1ml of RNAlater (Sigma-Aldrich), in order to stabilize and protect RNA with immediate RNase inactivation. Samples were kept at 4°C for 24 hours, then RNAlater were removed from tubes and samples were stored at -80°C until use.

### Cryosectioning

Before cryosectioning, samples were placed into the cryostat for at least 20 minutes before further processing. The sample was mounted on the round metallic mount of the cryostat with Killik  
 5 embedding medium. 20  $\mu$ m-thick cross sections were made and collected on warm (RT) gelatinated Superfrost slides (ThermoScientific Menzel Gläser (217655)). The sections were dried at RT for 1 hour and then stored at -20°C.

### Morphometrical analysis of muscles

10 The cross-sections of mice muscles were stained with Hematoxylin Gill № 2 (Sigma-Aldrich (GHS232)) and Eosin Y 1% aqueous solution (H/E staining procedure see ). Morphometrical analysis was performed on 3 cross-sections from each experimental group. The following parameters were evaluated: 1) area and perimeter of peripherally and centrally nucleated fibers, 2) the total number of nuclei referred to the number of fibers , 3) percentage of central nuclei referred  
 15 to the total number.

### Data analysis and statistics

Data are presented as means  $\pm$  S.D. (standard deviation of the mean). Student's unpaired t-test  
 20 was used to determine significant differences between the experimental groups. Values of \*p < 0.05 were considered significant, \*\*p < 0.01 very significant and \*\*\* p < 0.001 extremely significant.

The results obtained are explained with reference to the Figures.

**Figure 1** shows the formation of acetylcholine receptor clusters (dots). 1A, Vehicle treated control;  
 25 1B, Fc-nLG3; 1C, nLG3-Fc; 1D, ActR-Fc-nLG3. As expected at high concentration (10  $\mu$ M) of Fc-agrin (1B) and agrin-Fc (1C) AChR clusters were visible. However no clear clusters were visible using ActR-Fc-nLG3 (1D) and PBS (1A). Only incidental, probably spontaneous clusters, were visible. In addition, nLG3-Fc-ActR treated cells did also not show clusters on C2C12 treated cells. (results not shown). As expected AChR clusters appeared on nLG3-Fc and Fc-nLG3 treated  
 30 C2C12 cells albeit only at high concentrations. No clusters were visible at 1  $\mu$ M. It is not clear why actR-Fc-nLG3 did not show AChR clusters. The "ActR" part of ActR-Fc-nLG3 might be inhibiting the formation of clusters. This might be caused by steric hindering making proper agrin binding impossible or the agrin and myostatin signaling pathways interfere.

35 **Figure 2** shows the coomassie stained SDS-PAGE gel of: Fc-nLG3 (Lane 1); nLG3-Fc (Lane 2); Fc-ActR (Lane 3); ActR-Fc (Lane 4); ActR-Fc-nLG3 (Lane 5); (h)ActR-Fc (Lane 6); Fc-(h)nLG3 (Lane 7); (h)ActR-Fc-(h)nLG3 (Lane 8); (h)ActR-Fc-(h)LG3 (Lane 9); ActRmAb (Lane10); ActRmAb-(h)nLG3 (Lane 11); MyomAb (Lane 12); MyomAb-(h)nLG3 (Lane 13). All observed protein bands are as expected. The protein bands of ActR and (h)ActR derivatives are fuzzy

because this protein is glycosylated and the degree of glycosylation generates multiple bands of the same protein.

**Figure 3** shows the relative body weight increase over time. Nine weeks old mice were treated with vehicle, ActR-Fc, Fc-nLG3, ActR-Fc-nLG3 and a 1:1 mixture of ActR-Fc and Fc-nLG3 (Figure 2A); vehicle, (h)ActR-Fc, (h)Fc-(h)nLG3, and (h)ActR-Fc-(h)nLG3 (Figure 2B); vehicle, ActmAb and ActmAb-nLG3 (Figure 3C); vehicle, MyomAb and MyomAB-(h)nLG3 (Figure 2D). As expected ActR-Fc, ActR-Fc-nLG3, and the ActR-Fc + Fc-nLG3 mixture treated mice have significantly increased body weights compared to vehicle treated mice at day15. Surprisingly, ActR-Fc-nLG3 treated mice have a significantly lower body weight compared to ActR-Fc, ActR-Fc-nLG3, and the ActR-Fc + Fc-nLG3 mixture. Also (h)Fc-(h)nLG3, ActmAb-nLG3 MyomAb-(h)nLG3 have significantly lower body weight compared to their relative control compounds (h)ActR-Fc, ActmAb and MyomAb.

**Figure 4** shows the relative muscle weights for mice treated with vehicle, ActR-Fc, Fc-nLG3, ActR-Fc-nLG3, a 1:1 mixture of ActR-Fc, Fc-nLG3, (h)ActR-Fc, Fc-(h)nLG3, (h)ActR-Fc-(h)nLG3, ActmAb, ActmAb-nLG3, MyomAb and MyomAB-(h)nLG3. The relative mean muscle weights for the Gastrocnemius, Quadriceps and Triceps was calculated compared to muscles of vehicle treated mice. The results of the relative muscle weights resemble the results of the total body weights. As expected all compounds except Fc-nLG3 have significantly increased relative muscle weights. Surprisingly, compounds carrying in addition nLG3, or the human version of nLG3 (h)nLG3, ActR-Fc-nLG3, (h)ActR-Fc-(h)nLG3 ActmAb-(h)nLG3 MyomAb-(h)nLG3 have significantly lower body weights compared to their control compounds.

**Figure 5** shows the rotarod performance of the mice. The performance of ActR-Fc, Fc-nLG3, a 1:1 mixture of ActR-Fc and Fc-nLG3, ActmAb, MyomAb treated mice were not significantly increased. Surprisingly, the performance of the nLG3 resp (h)nLG3 containing compounds ActR-Fc-nLG3, ActmAb-nLG3 and MyomAB-(h)nLG3 were significantly increased compared to their control compounds ActR-Fc, ActmAb, MyomAb.

30

**Figure 6** shows the relative body weight increase over time. The relative mean body weight for every week was calculated. All 22 old mice were treated with vehicle (PBS) during the first three weeks of the experiment. In the following five weeks the aged mice were treated with vehicle, ActR-Fc, and ActR-Fc-nLG3. After week 3 the ActR-Fc dosed animals reach highly significant levels of weight increased compared to vehicle. After week 3 ActR-Fc-nLG3 dosed animals have significantly increased body weight compared to vehicle but significantly lower than the body weights of ActR-Fc.

**Figure 7** shows the relative muscle wet weights. The relative mean muscle weights for the Gastrocnemius, Quadriceps and Triceps was calculated compared to muscles of vehicle treated

40

mice. ActR-Fc dosed animals have highly significant levels of muscle weight increased compared to vehicle. ActR-Fc-nLG3 dosed animals are significantly increased in muscle weight compared to vehicle but significantly lower than the body weights of ActR-Fc.

5 **Figure 8** shows the treadmill performance of the aged mice. Figure 8A shows the treadmill performance during week three of vehicle dosing (before treatment). At this time point the performances of all groups are very similar. Figure 8B shows the mean treadmill performance during week 5 and 6 (after treatment). The performance of ActR-Fc and vehicle treated mice were lower after treatment than before treatment (not significant). It is likely that with increasing age the  
10 treadmill performance of these mice is declining. Surprisingly, the performance of ActR-Fc-nLG3 treated mice was improved after treatment compared to before treatment ( $p = X$ ). This shows that in spite of the mice being older, the treadmill performance improved. The performance of ActR-Fc-nLG3 in treated mice was significantly higher than vehicle and ActR-Fc treated mice. This shows that treatment with ActR-Fc-nLG3 improves the muscle endurance of the aged mice.

15

**Figure 9** shows the mean number of motivational electrical pulses per minute during the treadmill runs at week 5 and 6. Mice require more electrical pulses when they get exhausted. ActR-Fc and vehicle treated mice needed more pulses than ActR-Fc-nLG3 treated mice. This was highly significant  $p < 0.001$ . Interestingly, in the first nine minutes all  
20 three groups of mice required about the same number of pulses with no statistical differences. With increasing time on the treadmill, the performance of ActR-Fc-nLG3 treated mice was much better and the mice required fewer pulses than vehicle and ActR-Fc treated mice. This also clearly shows that mice treated with ActR-Fc-nLG3 have improved muscle endurance.

25

**Figure 10** shows the mean grip strength (GS) performance of the mice during week three of vehicle dosing (GS before treatment). At this time point the performances of all groups are very similar. Figure 7B shows the mean grip strength (GS) performance during week 5 and 6 (GS after treatment). The performance of vehicle treated mice was lower after treatment than before  
30 treatment (significant). The performance of ActR-Fc and ActR-Fc-nLG3 treated mice were higher after treatment than before treatment (significant). The GS performance of ActR-Fc-nLG3 and ActR-Fc treated mice were very similar after treatment and both are significantly increased compared to vehicle treated mice. So administration of the compound ActR-Fc-nLG3 has retained the increased muscle strength performance as ActR-Fc.

35 **Figure 11** Summary of muscle pathology. Cross sectional area (CSA), and number of nuclei per muscle fiber were determined for vehicle, ActR-Fc and ActR-Fc-nLG3 treated mice. From these results, the number of nuclei per CSA was calculated. In Figure 11 the relative values for CSA, and number of nuclei per nuclei per fiber and number of nuclei per CSA are depicted. ActR-Fc ( $p < 0.001$ ) and ActR-Fc-nLG3 ( $p < 0.05$ ) treated mice have

a statistically significantly increased CSA and number of nuclei per fiber compared to vehicle treated mice. In addition, compared to ActR-Fc-nLG3, ActR-Fc treated mice have a significantly increased CSA ( $p<0.01$ ) and number of nuclei ( $p<0.05$ ). However, compared to vehicle or ActR-Fc-nLG3, ActR-Fc treated mice have a significantly  
 5 decreased number of nuclei per fiber area ( $p<0.05$ ). Nuclei formation is promoted by the activity of the satellite cells. As satellite cells grow and differentiate they will fuse with an existing muscle fiber leading to more nuclei the muscle fiber.

**Figure 12** shows the relative body weight increase over time. Nine weeks old mice were  
 10 treated with vehicle, (h)ActR-Fc, (h)ActR-Fc-(h)nLG3, and (h)ActR-Fc-(h)LG3. At day 19 ActR-Fc, (h)ActR-Fc-(h)nLG3, and (h)ActR-Fc-(h)LG3 treated mice have significantly increased relative body weights ( $p<0.001$ ,  $p<0.05$ ,  $p<0.001$  respectively) compared to vehicle treated mice. The relative body weights of (h)ActR-Fc-(h)LG3 and (h)ActR-Fc are not significantly different. Notably, (h)ActR-Fc-(h)LG3 treated mice have a significantly  
 15 ( $p>0.01$ ) higher body weight compared to (h)ActR-Fc-(h)nLG3. The two proteins differ only by an 8 amino acid sequence insert in (h)nLG3. This insert is responsible for binding to the LRP4 receptor, so (h)nLG3 binds to the LRP4 receptor whereas (h)LG3 does not bind. Mice treated with (h)ActR-Fc-(h)nLG3 show a similar growth curve as ActR-Fc-nLG3 (Figure 3).

20

From these results it seems likely that muscle growth of ActR-Fc-nLG3 is solely caused by growth of muscle stem cells, (i.e. satellite cells) which fuse with the muscle fiber leading to more nuclei. More nuclei will lead to higher protein synthesis in the muscle fiber leading to a modest increase in muscle and body weight increase in ActR-Fc-nLG3,  
 25 (h)ActR-Fc-(h)nLG3, ActRmAb-nLG3 and MyomAb-nLG3 treated animals. Treatment with ActR-Fc also leads to more nuclei but the fiber growth is over proportional leading in fact to a lower nuclei density.

The claims defining the invention are as follows:

1. A compound comprising at least two components, a first component being the nLG3 domain from the C-terminus of human or mouse agrin, and at least one second component, selected from proteins or an antagonistic antibody that inhibit ActR2B-induced signaling activity in the presence of myostatin, the components being linked by means of linking entities.
2. A compound according to claim 1, selected from the group consisting of combinations of components, as follows:
  - A-L-B
  - B-L-A
  - A-L-C-L-B
  - B-L-C-L-A
  - C-L-B-L-A
  - C-L-A-L-B
  - B-L-A-L-C
  - A-L-B-L-C
  - A-L-D
  - D-L-A
  - A-L-E
  - E-L-A

in which

A represents an agrin nLG3 domain-containing protein,

B represents an extracellular domain of the ActRIB, ActRIC, ActRIIA, ActR2B receptor protein and follistatin

C represents a "stabilizer" domain,



D represents ActR2B-mediated signal-inhibiting mAb against the ActRIB, ActRIC, ActRIIA, and ActR2B receptor protein,

E represents ActR2B-mediated signal-inhibiting mAb against a member of the TGF-beta super family

L represents a linking entity.

3. A compound according to claim 2, in which the combinations is chosen from B-L-C-L-A, D-L-A and E-L-A.
4. A compound according to claim 2, in which A is selected from (h)nLG3 and proteins that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
5. A compound according to claim 4, in which A is at least 75%, 80%, 85%, 90%, 95% and completely identical with SEQ ID NO:10.
6. A compound according to claim 2, in which B is selected from the extracellular domain from ActRIB, ActRIC, ActRIIA, ActR2B and follistatin, and proteins that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
7. A compound according to claim 6, in which B is selected from human ActR2B receptor as defined in SEQ ID NO: 12 (AAC64515.1, GI:3769443) and follistatin as defined in SEQ ID NO: 25.
8. A compound according to claim 2, in which C is an "Fc" domain selected from an IGG1, IGG2, IGG3 and IGG4 and "Fc" domains that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
9. A compound according to claim 2, in which C is an "Fc" domain as defined in SEQ ID NO: 14 and proteins that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
10. A compound according to claim 2, in which D is selected from signalling blocking mAbs against ActRIB, ActRIC, ActRIIA, ActR2B, or mAbs that are at least 75%, 80%, 85%, 90% and 95% identical thereto.

11. A compound according to claim 6, in which D is a mAb as defined by ActRmAb(LC), (SEQ ID NO: 44) and ActRmAb(HC) (SEQ ID NO: 45) and mAbs that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
12. A compound according to claim 2, in which E is selected from signalling blocking mAbs against the TGF-beta super family, such as Activin B, Activin AB, Inhibin A, Inhibin B, GDF3, GDF1 1, Nodal, BMP2, BMP4, BMP7, BMP9, and BMP10 and that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
13. A compound according to claim 2, in which the linking entity is selected the roughly 15 amino acid unstructured region at the C-terminal end of the extracellular domain of ActR2B (the "tail"), and an artificial sequence of between 5 and 15, 20, 30, 50 or more amino acids that are relatively free of secondary structure.
14. A compound according to claim 6, in which E is a mAb as defined MyomAb(LC) SEQ ID NO: 32 and MyomAb(HC) SEQ ID NO: 31. and mAbs that are at least 75%, 80%, 85%, 90% and 95% identical thereto.
15. A compound according to claim 1, selected from the group consisting of ActR-Fc-nLG3 as defined in SEQ ID NO: 22, (h)ActR-Fc-(h)nLG3 as defined in SEQ ID NO: 23. Fol-Fc-nLG3 as defined in SEQ ID NO: 27, ActRmAb(HC) linked to (h)nLG3 (ActRmAb(HC)-(h)nLG3) as defined in SEQ ID NO: 30, MyomAb(HC linked to (h)nLG3) and (MyomAb(HC)-(h)nLG3) as defined in SEQ ID NO: 33.
16. A method of improving muscle performance, comprising the administration of an effective amount of a compound according to claim 1.
17. A pharmaceutical composition comprising at least one compound according to claim 1, formulated together with a pharmaceutically-acceptable carrier.
18. A method of treatment of a pathological condition leading to the loss of muscle function, comprising the administration of an effective amount of the compound according to claim 1, the pathological condition being one or more of:

- muscle atrophy as a result of treatment with a glucocorticoid such as cortisol, dexamethasone, betamethasone, prednisone, methylprednisolone, or prednisolone;
- muscle atrophy as a result of denervation due to nerve trauma or a result of degenerative, metabolic, or inflammatory neuropathy (e.g., Guillian-Barre syndrome, peripheral neuropathy, or exposure to environmental toxins or drugs);
- muscle atrophy as a result of myopathy, such as myotonia; a congenital myopathy, including nemaline myopathy, multi/minicore myopathy and myotubular (centronuclear) myopathy; mitochondrial myopathy; familial periodic paralysis; inflammatory myopathy; metabolic myopathy, such as caused by a glycogen or lipid storage disease; dermatomyositis; polymyositis; inclusion body myositis; myositis ossificans; rhabdomyolysis and myoglobinurias;
- myopathy caused by a muscular dystrophy syndrome, such as Duchenne, Becker, myotonic, fascioscapulohumeral, Emery-Dreifuss, oculopharyngeal, scapulohumeral, limb girdle, Fukuyama, a congenital muscular dystrophy, or hereditary distal myopathy
- musculoskeletal diseases such as osteoporosis, bone fracture, short stature, or dwarfism;
- adult motor neuron disease, infantile spinal muscular atrophy, amyotrophic lateral sclerosis, juvenile spinal muscular atrophy, autoimmune motor neuropathy with multifocal conduction block, paralysis due to stroke or spinal cord injury, skeletal immobilization due to trauma, prolonged bed rest, voluntary inactivity, involuntary inactivity, metabolic stress or nutritional insufficiency, cancer, AIDS, fasting, a thyroid gland disorder, diabetes, benign congenital hypotonia, central core disease, burn injury, chronic obstructive pulmonary disease, liver diseases (examples such as fibrosis, cirrhosis), sepsis, renal failure, congestive heart failure, ageing, space travel or time spent in a zero gravity environment.
- age-related conditions such as sarcopenia, skin atrophy, muscle wasting, brain atrophy, atherosclerosis, arteriosclerosis, pulmonary emphysema, osteoporosis, osteoarthritis, immunologic incompetence, high blood pressure, dementia, Huntington's disease, Alzheimer's disease, cataracts, age-related macular

5 degeneration, prostate cancer, stroke, diminished life expectancy, frailty, memory loss, wrinkles, impaired kidney function, and age-related hearing loss; metabolic disorders, including Type II Diabetes, Metabolic Syndrome, hyperglycemia, and obesity. Of course, patients may simultaneously suffer from one or more of these conditions, for example, sarcopenia and pulmonary emphysema, or sarcopenia and impaired kidney function.

- pathological disorders such as acute and/or chronic renal disease or failure, liver fibrosis or cirrhosis, cancer such as breast cancer, Parkinson's Disease; conditions associated with neuronal death, such as ALS, brain atrophy, or dementia and anemia.

10 In addition, there are losses suffered as a consequence of age, trauma or inactivity.

- cachexia, particularly cachexia associated with a rheumatoid arthritis and cachexia associated with cancer.

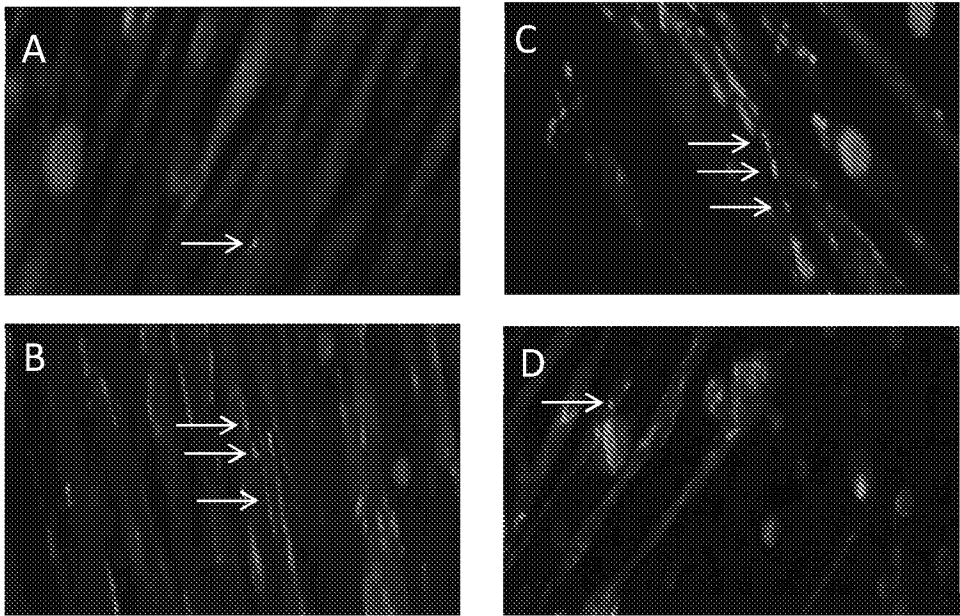


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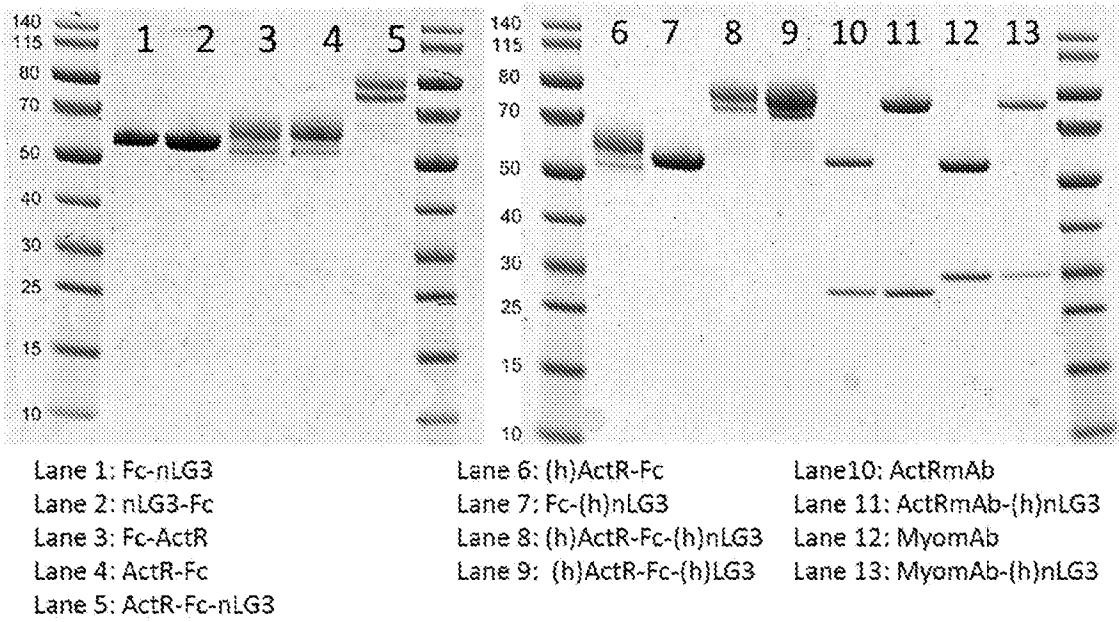


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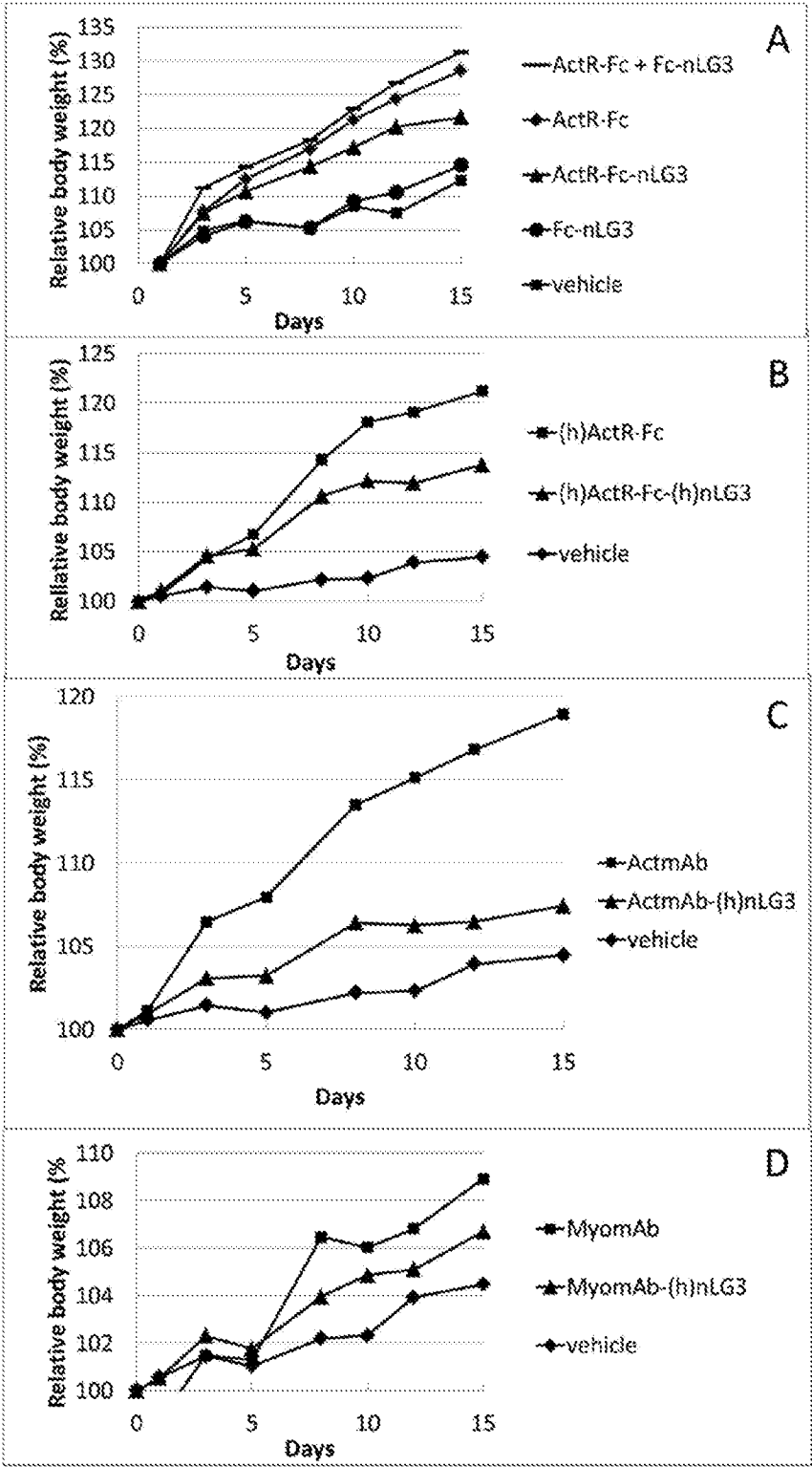


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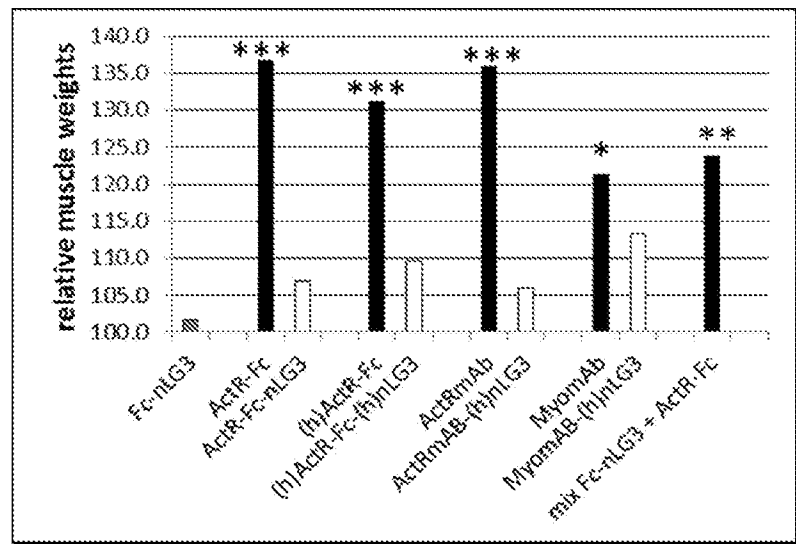


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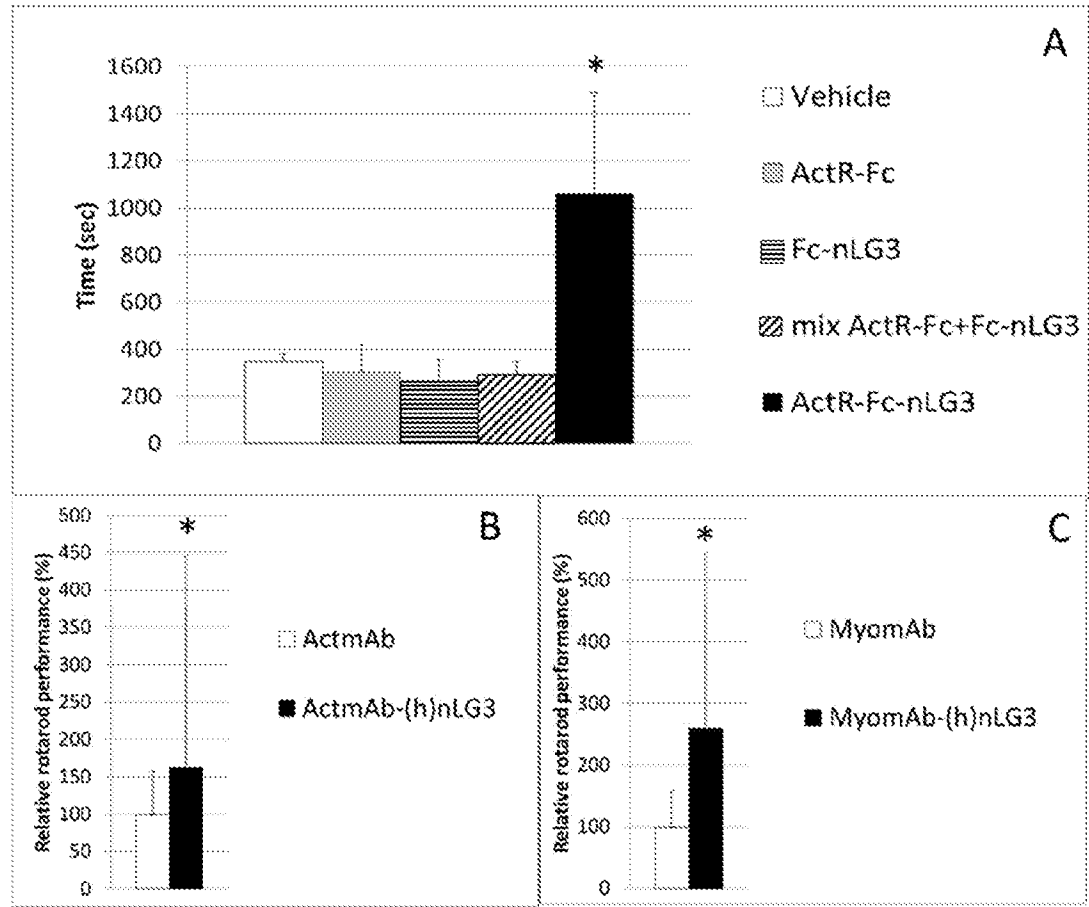


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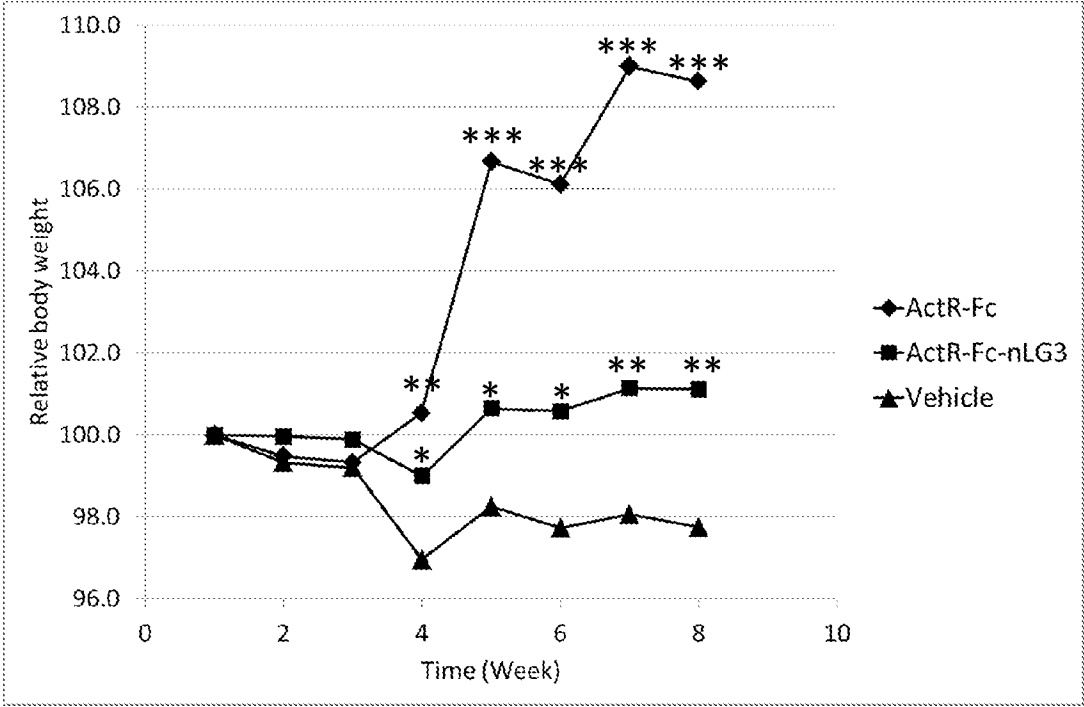


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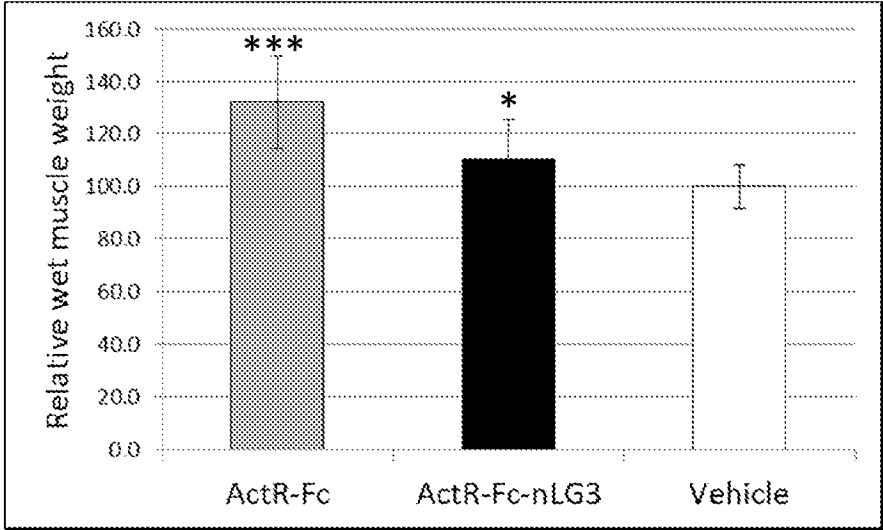


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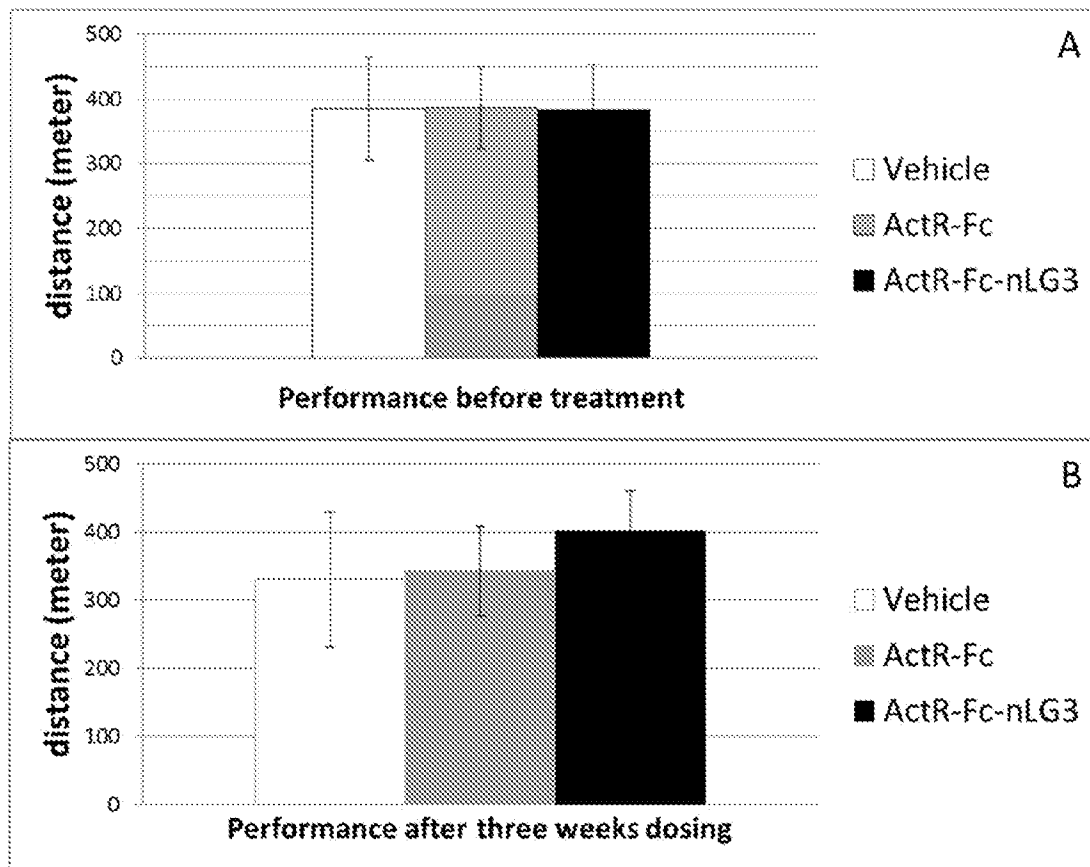


Figure 8

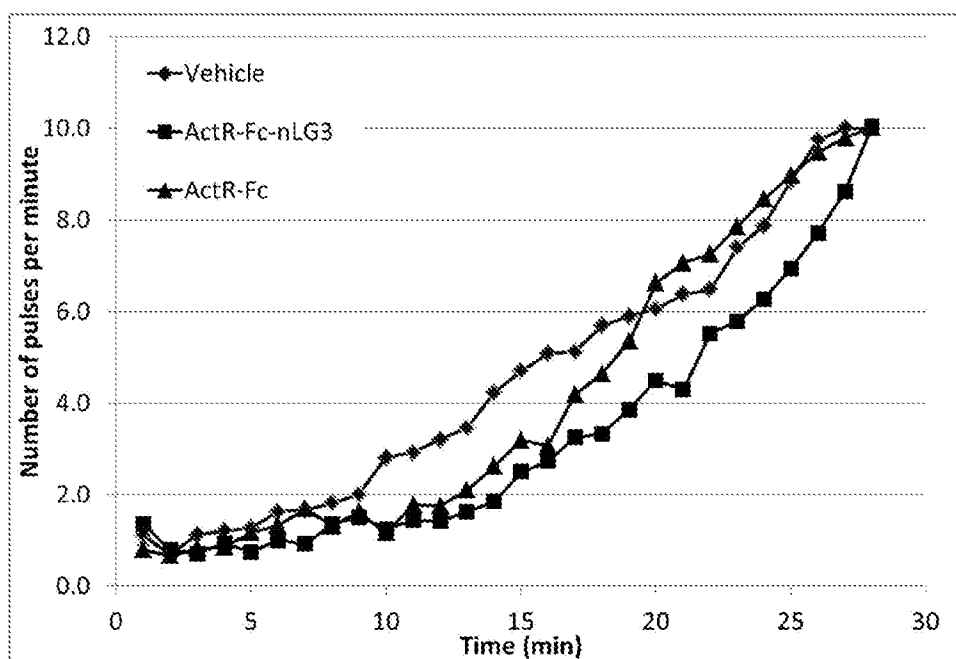


Figure 9

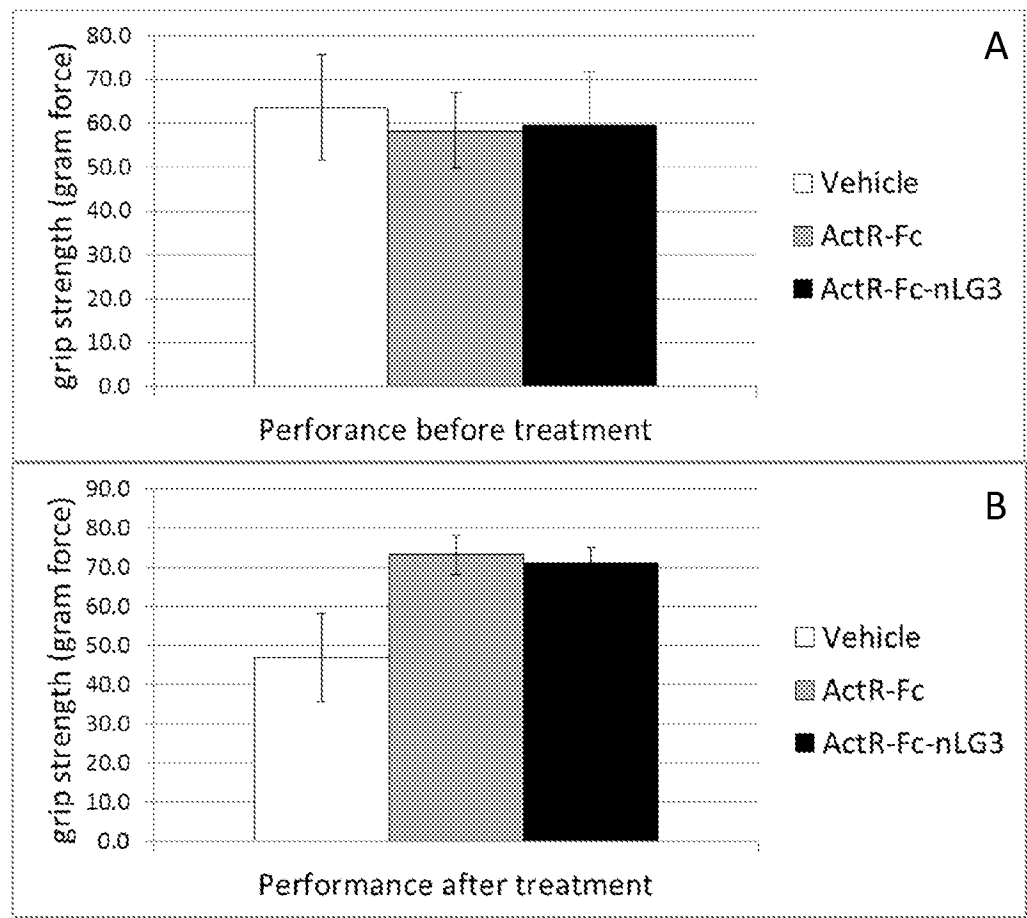


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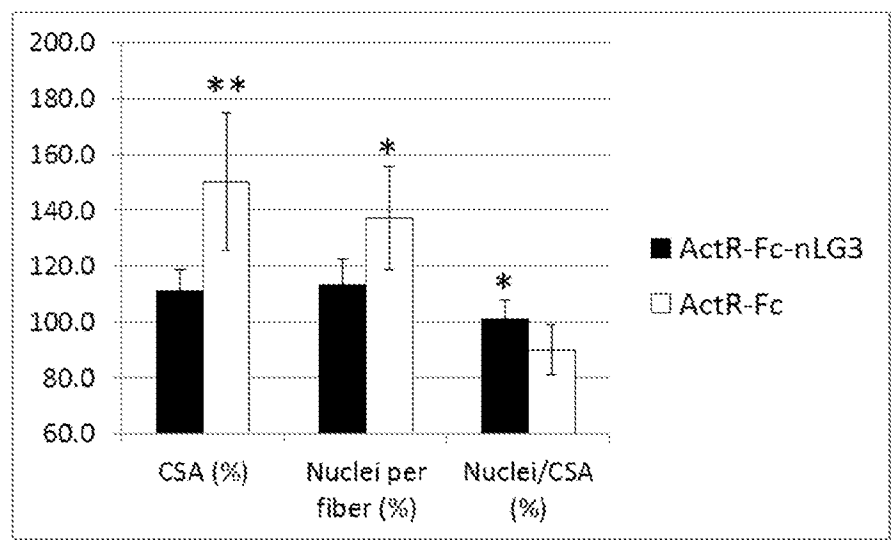


Figure 11

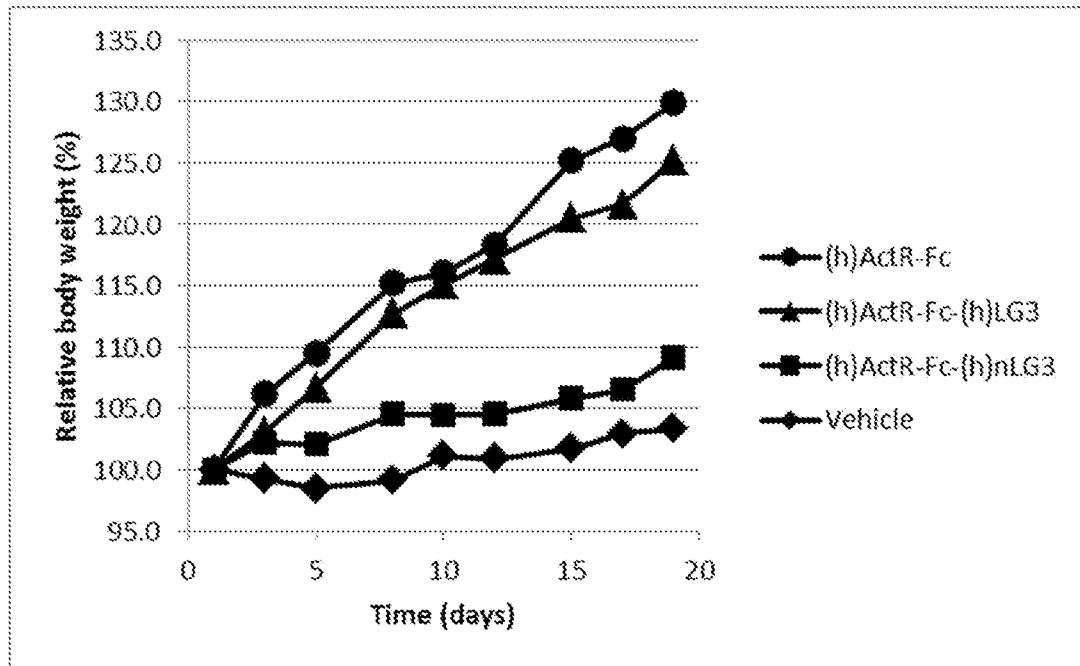


Figure 12

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Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly His Arg Gln Leu  
 165 170 175

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His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro  
180 185 190

Thr Leu

<210> 6  
<211> 186  
<212> PRT  
<213> human

<400> 6

Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg Thr Phe Val  
1 5 10 15

Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Lys Ala Leu Gln Ser Asn  
20 25 30

His Phe Glu Leu Ser Leu Arg Thr Glu Ala Thr Gln Gly Leu Val Leu  
35 40 45

Trp Ser Gly Lys Ala Thr Glu Arg Ala Asp Tyr Val Ala Leu Ala Ile  
50 55 60

Val Asp Gly His Leu Gln Leu Ser Tyr Asn Leu Gly Ser Gln Pro Val  
65 70 75 80

Val Leu Arg Ser Thr Val Pro Val Asn Thr Asn Arg Trp Leu Arg Val  
85 90 95

Val Ala His Arg Glu Gln Arg Glu Gly Ser Leu Gln Val Gly Asn Glu  
100 105 110

Ala Pro Val Thr Gly Ser Ser Pro Leu Gly Ala Thr Gln Leu Asp Thr  
115 120 125

Asp Gly Ala Leu Trp Leu Gly Gly Leu Pro Glu Leu Pro Val Gly Pro  
130 135 140

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Ala Leu Pro Lys Ala Tyr Gly Thr Gly Phe Val Gly Cys Leu Arg Asp  
145 150 155 160

Val Val Val Gly Arg His Pro Leu His Leu Leu Glu Asp Ala Val Thr  
165 170 175

Lys Pro Glu Leu Arg Pro Cys Pro Thr Pro  
180 185

<210> 7  
<211> 8  
<212> PRT  
<213> human

<400> 7

Glu Leu Ala Asn Glu Ile Pro Val  
1 5

<210> 8  
<211> 11  
<212> PRT  
<213> human

<400> 8

Pro Glu Thr Leu Asp Ser Gly Ala Leu His Ser  
1 5 10

<210> 9  
<211> 19  
<212> PRT  
<213> human

<400> 9

Glu Leu Ala Asn Glu Ile Pro Val Pro Glu Thr Leu Asp Ser Gly Ala  
1 5 10 15

Leu His Ser

<210> 10  
<211> 194  
<212> PRT



&lt;213&gt; human

&lt;400&gt; 10

Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg Thr Phe Val  
 1 5 10 15

Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Ala Asn Glu Ile Pro  
 20 25 30

Val Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser Leu Arg Thr  
 35 40 45

Glu Ala Thr Gln Gly Leu Val Leu Trp Ser Gly Lys Ala Thr Glu Arg  
 50 55 60

Ala Asp Tyr Val Ala Leu Ala Ile Val Asp Gly His Leu Gln Leu Ser  
 65 70 75 80

Tyr Asn Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr Val Pro Val  
 85 90 95

Asn Thr Asn Arg Trp Leu Arg Val Val Ala His Arg Glu Gln Arg Glu  
 100 105 110

Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly Ser Ser Pro  
 115 120 125

Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly Gly  
 130 135 140

Leu Pro Glu Leu Pro Val Gly Pro Ala Leu Pro Lys Ala Tyr Gly Thr  
 145 150 155 160

Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly Arg His Pro Leu  
 165 170 175

His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro  
 180 185 190

Thr Pro

<210> 11  
 <211> 115  
 <212> PRT  
 <213> mouse

<400> 11

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
 1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
 20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
 35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
 50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
 65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
 85 90 95

Pro Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
 100 105 110

Leu Leu Thr  
 115

<210> 12  
 <211> 115  
 <212> PRT  
 <213> human

<400> 12

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
 1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
85 90 95

Ala Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
100 105 110

Leu Leu Thr  
115

<210> 13  
<211> 15  
<212> PRT  
<213> recombinant

<400> 13

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser  
1 5 10 15

<210> 14  
<211> 227  
<212> PRT  
<213> human

<400> 14

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
1 5 10 15

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Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
20 25 30

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
35 40 45

Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
50 55 60

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
65 70 75 80

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
85 90 95

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
100 105 110

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
115 120 125

Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser  
130 135 140

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
145 150 155 160

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro  
165 170 175

Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
180 185 190

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
195 200 205

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
210 215 220

Pro Gly Lys  
225

<210> 15  
<211> 435  
<212> PRT  
<213> recombinant

<400> 15

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
1 5 10 15

Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
20 25 30

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
35 40 45

Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
50 55 60

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
65 70 75 80

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
85 90 95

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
100 105 110

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
115 120 125

Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser  
130 135 140

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
145 150 155 160

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro

165

170

175

Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
 180 185 190

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
 195 200 205

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
 210 215 220

Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
 225 230 235 240

Gly Ser Val Gly Asp Leu Glu Thr Leu Ala Phe Asp Gly Arg Thr Tyr  
 245 250 255

Ile Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Thr Asn Glu Ile  
 260 265 270

Pro Ala Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser Leu Arg  
 275 280 285

Thr Glu Ala Thr Gln Gly Leu Val Leu Trp Ile Gly Lys Val Gly Glu  
 290 295 300

Arg Ala Asp Tyr Met Ala Leu Ala Ile Val Asp Gly His Leu Gln Leu  
 305 310 315 320

Ser Tyr Asp Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr Val Lys  
 325 330 335

Val Asn Thr Asn Arg Trp Leu Arg Val Arg Ala His Arg Glu His Arg  
 340 345 350

Glu Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly Ser Ser  
 355 360 365

Pro Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly

370

375

380

Gly Leu Gln Lys Leu Pro Val Gly Gln Ala Leu Pro Lys Ala Tyr Gly  
 385 390 395 400

Thr Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly His Arg Gln  
 405 410 415

Leu His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys  
 420 425 430

Pro Thr Leu  
 435

<210> 16  
 <211> 436  
 <212> PRT  
 <213> recombinant

<400> 16

Ser Val Gly Asp Leu Glu Thr Leu Ala Phe Asp Gly Arg Thr Tyr Ile  
 1 5 10 15

Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Thr Asn Glu Ile Pro  
 20 25 30

Ala Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser Leu Arg Thr  
 35 40 45

Glu Ala Thr Gln Gly Leu Val Leu Trp Ile Gly Lys Val Gly Glu Arg  
 50 55 60

Ala Asp Tyr Met Ala Leu Ala Ile Val Asp Gly His Leu Gln Leu Ser  
 65 70 75 80

Tyr Asp Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr Val Lys Val  
 85 90 95

Asn Thr Asn Arg Trp Leu Arg Val Arg Ala His Arg Glu His Arg Glu  
 100 105 110

Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly Ser Ser Pro  
115 120 125

Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly Gly  
130 135 140

Leu Gln Lys Leu Pro Val Gly Gln Ala Leu Pro Lys Ala Tyr Gly Thr  
145 150 155 160

Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly His Arg Gln Leu  
165 170 175

His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro  
180 185 190

Thr Leu Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly  
195 200 205

Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu  
210 215 220

Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu  
225 230 235 240

Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser  
245 250 255

His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu  
260 265 270

Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr  
275 280 285

Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn  
290 295 300

Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro  
305 310 315 320



Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln  
325 330 335

Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val  
340 345 350

Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val  
355 360 365

Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro  
370 375 380

Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr  
385 390 395 400

Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val  
405 410 415

Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu  
420 425 430

Ser Pro Gly Lys  
435

<210> 17  
<211> 435  
<212> PRT  
<213> recombinant

<400> 17

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
1 5 10 15

Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
20 25 30

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
35 40 45

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Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
50 55 60

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
65 70 75 80

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
85 90 95

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
100 105 110

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
115 120 125

Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser  
130 135 140

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
145 150 155 160

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro  
165 170 175

Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
180 185 190

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
195 200 205

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
210 215 220

Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
225 230 235 240

Gly Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg Thr Phe  
245 250 255

TP0063W0-seq1-000001.txt

Val Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Ala Asn Glu Ile  
260 265 270

Pro Val Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser Leu Arg  
275 280 285

Thr Glu Ala Thr Gln Gly Leu Val Leu Trp Ser Gly Lys Ala Thr Glu  
290 295 300

Arg Ala Asp Tyr Val Ala Leu Ala Ile Val Asp Gly His Leu Gln Leu  
305 310 315 320

Ser Tyr Asn Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr Val Pro  
325 330 335

Val Asn Thr Asn Arg Trp Leu Arg Val Val Ala His Arg Glu Gln Arg  
340 345 350

Glu Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly Ser Ser  
355 360 365

Pro Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly  
370 375 380

Gly Leu Pro Glu Leu Pro Val Gly Pro Ala Leu Pro Lys Ala Tyr Gly  
385 390 395 400

Thr Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly Arg His Pro  
405 410 415

Leu His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys  
420 425 430

Pro Thr Pro  
435

<210> 18  
<211> 427  
<212> PRT

&lt;213&gt; recombinant

&lt;400&gt; 18

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
 1 5 10 15

Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
 20 25 30

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
 35 40 45

Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
 50 55 60

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
 65 70 75 80

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
 85 90 95

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
 100 105 110

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
 115 120 125

Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser  
 130 135 140

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
 145 150 155 160

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro  
 165 170 175

Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
 180 185 190

TP0063W0-seq1-000001.txt

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
195 200 205

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
210 215 220

Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
225 230 235 240

Gly Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg Thr Phe  
245 250 255

Val Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Lys Ala Leu Gln Ser  
260 265 270

Asn His Phe Glu Leu Ser Leu Arg Thr Glu Ala Thr Gln Gly Leu Val  
275 280 285

Leu Trp Ser Gly Lys Ala Thr Glu Arg Ala Asp Tyr Val Ala Leu Ala  
290 295 300

Ile Val Asp Gly His Leu Gln Leu Ser Tyr Asn Leu Gly Ser Gln Pro  
305 310 315 320

Val Val Leu Arg Ser Thr Val Pro Val Asn Thr Asn Arg Trp Leu Arg  
325 330 335

Val Val Ala His Arg Glu Gln Arg Glu Gly Ser Leu Gln Val Gly Asn  
340 345 350

Glu Ala Pro Val Thr Gly Ser Ser Pro Leu Gly Ala Thr Gln Leu Asp  
355 360 365

Thr Asp Gly Ala Leu Trp Leu Gly Gly Leu Pro Glu Leu Pro Val Gly  
370 375 380

Pro Ala Leu Pro Lys Ala Tyr Gly Thr Gly Phe Val Gly Cys Leu Arg  
385 390 395 400

TP0063W0-seq1-000001.txt

Asp Val Val Val Gly Arg His Pro Leu His Leu Leu Glu Asp Ala Val  
405 410 415

Thr Lys Pro Glu Leu Arg Pro Cys Pro Thr Pro  
420 425

<210> 19  
<211> 357  
<212> PRT  
<213> Recombinant

<400> 19

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
85 90 95

Pro Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
100 105 110

Leu Leu Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
115 120 125

Gly Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
130 135 140

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr

145					150						155					160
Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	
				165					170					175		
Ser	His	Glu	Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	
			180					185					190			
Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	
		195					200					205				
Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	
	210					215					220					
Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	
225					230					235					240	
Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	
				245					250					255		
Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	
			260					265					270			
Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	
		275					280					285				
Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	
	290					295					300					
Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	
305					310					315					320	
Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	
				325					330					335		
Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	
			340				345						350			
Leu	Ser	Pro	Gly	Lys												

355

<210> 20  
 <211> 357  
 <212> PRT  
 <213> recombinant  
 <400> 20

Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
 1 5 10 15

Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
 20 25 30

Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
 35 40 45

Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
 50 55 60

His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
 65 70 75 80

Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
 85 90 95

Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
 100 105 110

Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
 115 120 125

Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser  
 130 135 140

Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
 145 150 155 160

Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro  
 165 170 175



Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
180 185 190

Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
195 200 205

His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
210 215 220

Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
225 230 235 240

Gly Ser Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp  
245 250 255

Glu Leu Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu  
260 265 270

Gln Asp Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly  
275 280 285

Thr Ile Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys  
290 295 300

Tyr Asp Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr  
305 310 315 320

Phe Cys Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu  
325 330 335

Pro Glu Pro Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala  
340 345 350

Pro Thr Leu Leu Thr  
355

<210> 21  
<211> 357

&lt;212&gt; PRT

&lt;213&gt; recombinant

&lt;400&gt; 21

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
 1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
 20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
 35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
 50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
 65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
 85 90 95

Ala Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
 100 105 110

Leu Leu Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
 115 120 125

Gly Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
 130 135 140

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr  
 145 150 155 160

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val  
 165 170 175

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val  
 180 185 190

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Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser  
195 200 205

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu  
210 215 220

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala  
225 230 235 240

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro  
245 250 255

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln  
260 265 270

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala  
275 280 285

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr  
290 295 300

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu  
305 310 315 320

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser  
325 330 335

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser  
340 345 350

Leu Ser Pro Gly Lys  
355

<210> 22  
<211> 565  
<212> PRT  
<213> recombinant

<400> 22

TP0063W0-seq1-000001.txt

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
85 90 95

Pro Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
100 105 110

Leu Leu Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
115 120 125

Gly Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
130 135 140

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr  
145 150 155 160

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val  
165 170 175

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val  
180 185 190

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser  
195 200 205

TP0063W0-seq1-000001.txt

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu  
210 215 220

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala  
225 230 235 240

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro  
245 250 255

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln  
260 265 270

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala  
275 280 285

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr  
290 295 300

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu  
305 310 315 320

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser  
325 330 335

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser  
340 345 350

Leu Ser Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly  
355 360 365

Gly Gly Gly Ser Val Gly Asp Leu Glu Thr Leu Ala Phe Asp Gly Arg  
370 375 380

Thr Tyr Ile Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Thr Asn  
385 390 395 400

Glu Ile Pro Ala Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser  
405 410 415

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Leu Arg Thr Glu Ala Thr Gln Gly Leu Val Leu Trp Ile Gly Lys Val  
420 425 430

Gly Glu Arg Ala Asp Tyr Met Ala Leu Ala Ile Val Asp Gly His Leu  
435 440 445

Gln Leu Ser Tyr Asp Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr  
450 455 460

Val Lys Val Asn Thr Asn Arg Trp Leu Arg Val Arg Ala His Arg Glu  
465 470 475 480

His Arg Glu Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly  
485 490 495

Ser Ser Pro Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp  
500 505 510

Leu Gly Gly Leu Gln Lys Leu Pro Val Gly Gln Ala Leu Pro Lys Ala  
515 520 525

Tyr Gly Thr Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly His  
530 535 540

Arg Gln Leu His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg  
545 550 555 560

Pro Cys Pro Thr Leu  
565

<210> 23  
<211> 565  
<212> PRT  
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<400> 23

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp

20

25

30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
           35                          40                          45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
       50                          55                          60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
   65                          70                          75                          80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
                           85                          90                          95

Ala Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
                           100                          105                          110

Leu Leu Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
       115                          120                          125

Gly Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
       130                          135                          140

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr  
   145                          150                          155                          160

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val  
                           165                          170                          175

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val  
                           180                          185                          190

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser  
       195                          200                          205

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu  
       210                          215                          220

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala

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225                               230                               235                               240

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
      245                               250                               255

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
      260                               265                               270

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
      275                               280                               285

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
      290                               295                               300

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
305                               310                               315                               320

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
      325                               330                               335

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
      340                               345                               350

Leu Ser Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
      355                               360                               365

Gly Gly Gly Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg
      370                               375                               380

Thr Phe Val Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Ala Asn
385                               390                               395                               400

Glu Ile Pro Val Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser
      405                               410                               415

Leu Arg Thr Glu Ala Thr Gln Gly Leu Val Leu Trp Ser Gly Lys Ala
      420                               425                               430

Thr Glu Arg Ala Asp Tyr Val Ala Leu Ala Ile Val Asp Gly His Leu

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435

440

445

Gln Leu Ser Tyr Asn Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr  
 450 455 460

Val Pro Val Asn Thr Asn Arg Trp Leu Arg Val Val Ala His Arg Glu  
 465 470 475 480

Gln Arg Glu Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly  
 485 490 495

Ser Ser Pro Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp  
 500 505 510

Leu Gly Gly Leu Pro Glu Leu Pro Val Gly Pro Ala Leu Pro Lys Ala  
 515 520 525

Tyr Gly Thr Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly Arg  
 530 535 540

His Pro Leu His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg  
 545 550 555 560

Pro Cys Pro Thr Pro  
 565

<210> 24  
 <211> 557  
 <212> PRT  
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<400> 24

Glu Ala Glu Thr Arg Glu Cys Ile Tyr Tyr Asn Ala Asn Trp Glu Leu  
 1 5 10 15

Glu Arg Thr Asn Gln Ser Gly Leu Glu Arg Cys Glu Gly Glu Gln Asp  
 20 25 30

Lys Arg Leu His Cys Tyr Ala Ser Trp Arg Asn Ser Ser Gly Thr Ile  
 35 40 45

Glu Leu Val Lys Lys Gly Cys Trp Leu Asp Asp Phe Asn Cys Tyr Asp  
50 55 60

Arg Gln Glu Cys Val Ala Thr Glu Glu Asn Pro Gln Val Tyr Phe Cys  
65 70 75 80

Cys Cys Glu Gly Asn Phe Cys Asn Glu Arg Phe Thr His Leu Pro Glu  
85 90 95

Ala Gly Gly Pro Glu Val Thr Tyr Glu Pro Pro Pro Thr Ala Pro Thr  
100 105 110

Leu Leu Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly  
115 120 125

Gly Ser Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
130 135 140

Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr  
145 150 155 160

Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val  
165 170 175

Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val  
180 185 190

Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser  
195 200 205

Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu  
210 215 220

Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala  
225 230 235 240

Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro  
245 250 255

Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln  
260 265 270

Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala  
275 280 285

Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr  
290 295 300

Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu  
305 310 315 320

Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser  
325 330 335

Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser  
340 345 350

Leu Ser Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly  
355 360 365

Gly Gly Gly Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly Arg  
370 375 380

Thr Phe Val Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Lys Ala Leu  
385 390 395 400

Gln Ser Asn His Phe Glu Leu Ser Leu Arg Thr Glu Ala Thr Gln Gly  
405 410 415

Leu Val Leu Trp Ser Gly Lys Ala Thr Glu Arg Ala Asp Tyr Val Ala  
420 425 430

Leu Ala Ile Val Asp Gly His Leu Gln Leu Ser Tyr Asn Leu Gly Ser  
435 440 445

Gln Pro Val Val Leu Arg Ser Thr Val Pro Val Asn Thr Asn Arg Trp  
450 455 460

Leu Arg Val Val Ala His Arg Glu Gln Arg Glu Gly Ser Leu Gln Val  
465 470 475 480

Gly Asn Glu Ala Pro Val Thr Gly Ser Ser Pro Leu Gly Ala Thr Gln  
485 490 495

Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly Gly Leu Pro Glu Leu Pro  
500 505 510

Val Gly Pro Ala Leu Pro Lys Ala Tyr Gly Thr Gly Phe Val Gly Cys  
515 520 525

Leu Arg Asp Val Val Val Gly Arg His Pro Leu His Leu Leu Glu Asp  
530 535 540

Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro Thr Pro  
545 550 555

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<211> 288  
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<213> recombinant

<400> 25

Gly Asn Cys Trp Leu Arg Gln Ala Lys Asn Gly Arg Cys Gln Val Leu  
1 5 10 15

Tyr Lys Thr Glu Leu Ser Lys Glu Glu Cys Cys Ser Thr Gly Arg Leu  
20 25 30

Ser Thr Ser Trp Thr Glu Glu Asp Val Asn Asp Asn Thr Leu Phe Lys  
35 40 45

Trp Met Ile Phe Asn Gly Gly Ala Pro Asn Cys Ile Pro Cys Lys Glu  
50 55 60

Thr Cys Glu Asn Val Asp Cys Gly Pro Gly Lys Lys Cys Arg Met Asn  
65 70 75 80

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Lys Lys Asn Lys Pro Arg Cys Val Cys Ala Pro Asp Cys Ser Asn Ile  
85 90 95

Thr Trp Lys Gly Pro Val Cys Gly Leu Asp Gly Lys Thr Tyr Arg Asn  
100 105 110

Glu Cys Ala Leu Leu Lys Ala Arg Cys Lys Glu Gln Pro Glu Leu Glu  
115 120 125

Val Gln Tyr Gln Gly Lys Cys Lys Lys Thr Cys Arg Asp Val Phe Cys  
130 135 140

Pro Gly Ser Ser Thr Cys Val Val Asp Gln Thr Asn Asn Ala Tyr Cys  
145 150 155 160

Val Thr Cys Asn Arg Ile Cys Pro Glu Pro Ser Ser Ser Glu Gln Tyr  
165 170 175

Leu Cys Gly Asn Asp Gly Val Thr Tyr Ser Ser Ala Cys His Leu Arg  
180 185 190

Lys Ala Thr Cys Leu Leu Gly Arg Ser Ile Gly Leu Ala Tyr Glu Gly  
195 200 205

Lys Cys Ile Lys Ala Lys Ser Cys Glu Asp Ile Gln Cys Gly Gly Gly  
210 215 220

Lys Lys Cys Leu Trp Asp Ser Lys Val Gly Arg Gly Arg Cys Ser Leu  
225 230 235 240

Cys Asp Glu Leu Cys Pro Asp Ser Lys Ser Asp Glu Pro Val Cys Ala  
245 250 255

Ser Asp Asn Ala Thr Tyr Ala Ser Glu Cys Ala Met Lys Glu Ala Ala  
260 265 270

Cys Ser Ser Gly Val Leu Leu Glu Val Lys His Ser Gly Ser Cys Asn  
275 280 285

<210> 26  
 <211> 530  
 <212> PRT  
 <213> recombinant

<400> 26

Gly Asn Cys Trp Leu Arg Gln Ala Lys Asn Gly Arg Cys Gln Val Leu  
 1 5 10 15

Tyr Lys Thr Glu Leu Ser Lys Glu Glu Cys Cys Ser Thr Gly Arg Leu  
 20 25 30

Ser Thr Ser Trp Thr Glu Glu Asp Val Asn Asp Asn Thr Leu Phe Lys  
 35 40 45

Trp Met Ile Phe Asn Gly Gly Ala Pro Asn Cys Ile Pro Cys Lys Glu  
 50 55 60

Thr Cys Glu Asn Val Asp Cys Gly Pro Gly Lys Lys Cys Arg Met Asn  
 65 70 75 80

Lys Lys Asn Lys Pro Arg Cys Val Cys Ala Pro Asp Cys Ser Asn Ile  
 85 90 95

Thr Trp Lys Gly Pro Val Cys Gly Leu Asp Gly Lys Thr Tyr Arg Asn  
 100 105 110

Glu Cys Ala Leu Leu Lys Ala Arg Cys Lys Glu Gln Pro Glu Leu Glu  
 115 120 125

Val Gln Tyr Gln Gly Lys Cys Lys Lys Thr Cys Arg Asp Val Phe Cys  
 130 135 140

Pro Gly Ser Ser Thr Cys Val Val Asp Gln Thr Asn Asn Ala Tyr Cys  
 145 150 155 160

Val Thr Cys Asn Arg Ile Cys Pro Glu Pro Ser Ser Ser Glu Gln Tyr  
 165 170 175

TP0063W0-seq1-000001.txt

Leu Cys Gly Asn Asp Gly Val Thr Tyr Ser Ser Ala Cys His Leu Arg  
180 185 190

Lys Ala Thr Cys Leu Leu Gly Arg Ser Ile Gly Leu Ala Tyr Glu Gly  
195 200 205

Lys Cys Ile Lys Ala Lys Ser Cys Glu Asp Ile Gln Cys Gly Gly Gly  
210 215 220

Lys Lys Cys Leu Trp Asp Ser Lys Val Gly Arg Gly Arg Cys Ser Leu  
225 230 235 240

Cys Asp Glu Leu Cys Pro Asp Ser Lys Ser Asp Glu Pro Val Cys Ala  
245 250 255

Ser Asp Asn Ala Thr Tyr Ala Ser Glu Cys Ala Met Lys Glu Ala Ala  
260 265 270

Cys Ser Ser Gly Val Leu Leu Glu Val Lys His Ser Gly Ser Cys Asn  
275 280 285

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp  
290 295 300

Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
305 310 315 320

Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
325 330 335

Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
340 345 350

Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
355 360 365

Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
370 375 380

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Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
385 390 395 400

Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
405 410 415

Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
420 425 430

Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu  
435 440 445

Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
450 455 460

Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
465 470 475 480

Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
485 490 495

Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
500 505 510

Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
515 520 525

Gly Lys  
530

<210> 27  
<211> 738  
<212> PRT  
<213> recombinant

<400> 27

Gly Asn Cys Trp Leu Arg Gln Ala Lys Asn Gly Arg Cys Gln Val Leu  
1 5 10 15

Tyr Lys Thr Glu Leu Ser Lys Glu Glu Cys Cys Ser Thr Gly Arg Leu



20

25

30

Ser Thr Ser Trp Thr Glu Glu Asp Val Asn Asp Asn Thr Leu Phe Lys  
           35                          40                          45

Trp Met Ile Phe Asn Gly Gly Ala Pro Asn Cys Ile Pro Cys Lys Glu  
       50                          55                          60

Thr Cys Glu Asn Val Asp Cys Gly Pro Gly Lys Lys Cys Arg Met Asn  
  65                          70                          75                          80

Lys Lys Asn Lys Pro Arg Cys Val Cys Ala Pro Asp Cys Ser Asn Ile  
                           85                          90                          95

Thr Trp Lys Gly Pro Val Cys Gly Leu Asp Gly Lys Thr Tyr Arg Asn  
           100                          105                          110

Glu Cys Ala Leu Leu Lys Ala Arg Cys Lys Glu Gln Pro Glu Leu Glu  
           115                          120                          125

Val Gln Tyr Gln Gly Lys Cys Lys Lys Thr Cys Arg Asp Val Phe Cys  
       130                          135                          140

Pro Gly Ser Ser Thr Cys Val Val Asp Gln Thr Asn Asn Ala Tyr Cys  
  145                          150                          155                          160

Val Thr Cys Asn Arg Ile Cys Pro Glu Pro Ser Ser Ser Glu Gln Tyr  
                           165                          170                          175

Leu Cys Gly Asn Asp Gly Val Thr Tyr Ser Ser Ala Cys His Leu Arg  
           180                          185                          190

Lys Ala Thr Cys Leu Leu Gly Arg Ser Ile Gly Leu Ala Tyr Glu Gly  
       195                          200                          205

Lys Cys Ile Lys Ala Lys Ser Cys Glu Asp Ile Gln Cys Gly Gly Gly  
       210                          215                          220

Lys Lys Cys Leu Trp Asp Ser Lys Val Gly Arg Gly Arg Cys Ser Leu

225		230		235		240									
Cys	Asp	Glu	Leu	Cys	Pro	Asp	Ser	Lys	Ser	Asp	Glu	Pro	Val	Cys	Ala
				245					250					255	
Ser	Asp	Asn	Ala	Thr	Tyr	Ala	Ser	Glu	Cys	Ala	Met	Lys	Glu	Ala	Ala
			260					265					270		
Cys	Ser	Ser	Gly	Val	Leu	Leu	Glu	Val	Lys	His	Ser	Gly	Ser	Cys	Asn
		275					280					285			
Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	Asp
	290					295					300				
Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly
305					310					315					320
Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile
				325				330						335	
Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu
			340					345					350		
Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His
		355					360					365			
Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg
	370					375					380				
Val	Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys
385					390				395						400
Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu
				405					410					415	
Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr
			420					425					430		
Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu

435

440

445

Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 450 455 460

Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 465 470 475 480

Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 485 490 495

Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 500 505 510

Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 515 520 525

Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly  
 530 535 540

Ser Val Gly Asp Leu Glu Thr Leu Ala Phe Asp Gly Arg Thr Tyr Ile  
 545 550 555 560

Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Thr Asn Glu Ile Pro  
 565 570 575

Ala Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu Ser Leu Arg Thr  
 580 585 590

Glu Ala Thr Gln Gly Leu Val Leu Trp Ile Gly Lys Val Gly Glu Arg  
 595 600 605

Ala Asp Tyr Met Ala Leu Ala Ile Val Asp Gly His Leu Gln Leu Ser  
 610 615 620

Tyr Asp Leu Gly Ser Gln Pro Val Val Leu Arg Ser Thr Val Lys Val  
 625 630 635 640

Asn Thr Asn Arg Trp Leu Arg Val Arg Ala His Arg Glu His Arg Glu

645

650

655

Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr Gly Ser Ser Pro  
 660 665 670

Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly Gly  
 675 680 685

Leu Gln Lys Leu Pro Val Gly Gln Ala Leu Pro Lys Ala Tyr Gly Thr  
 690 695 700

Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly His Arg Gln Leu  
 705 710 715 720

His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro  
 725 730 735

Thr Leu

<210> 28  
 <211> 445  
 <212> PRT  
 <213> recombinant

<400> 28

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala  
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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Ser  
 20 25 30

Tyr Ile Asn Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met  
 35 40 45

Gly Thr Ile Asn Pro Val Ser Gly Ser Thr Ser Tyr Ala Gln Lys Phe  
 50 55 60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr  
 65 70 75 80

Met Glu Leu Ser Arg Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Arg Gly Gly Trp Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr  
100 105 110

Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro  
115 120 125

Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val  
130 135 140

Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
145 150 155 160

Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly  
165 170 175

Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly  
180 185 190

Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys  
195 200 205

Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys  
210 215 220

Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu  
225 230 235 240

Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu  
245 250 255

Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys  
260 265 270

Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys  
275 280 285

Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu  
290 295 300

Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys  
305 310 315 320

Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys  
325 330 335

Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser  
340 345 350

Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys  
355 360 365

Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln  
370 375 380

Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly  
385 390 395 400

Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln  
405 410 415

Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn  
420 425 430

His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
435 440 445

<210> 29  
<211> 217  
<212> PRT  
<213> recombinant

<400> 29

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln  
1 5 10 15

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Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Ser Asp Val Gly Ser Tyr  
20 25 30

Asn Tyr Val Asn Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu  
35 40 45

Met Ile Tyr Gly Val Ser Lys Arg Pro Ser Gly Val Ser Asn Arg Phe  
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu  
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Thr Phe Ala Gly Gly  
85 90 95

Ser Tyr Tyr Gly Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly  
100 105 110

Gln Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser Glu  
115 120 125

Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser Asp Phe  
130 135 140

Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser Ser Pro Val  
145 150 155 160

Lys Ala Gly Val Glu Thr Thr Thr Pro Ser Lys Gln Ser Asn Asn Lys  
165 170 175

Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro Glu Gln Trp Lys Ser  
180 185 190

His Arg Ser Tyr Ser Cys Gln Val Thr His Glu Gly Ser Thr Val Glu  
195 200 205

Lys Thr Val Ala Pro Thr Glu Cys Ser  
210 215

<210> 30  
 <211> 653  
 <212> PRT  
 <213> recombinant

<400> 30

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala  
 1 5 10 15

Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Ser  
 20 25 30

Tyr Ile Asn Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met  
 35 40 45

Gly Thr Ile Asn Pro Val Ser Gly Ser Thr Ser Tyr Ala Gln Lys Phe  
 50 55 60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr  
 65 70 75 80

Met Glu Leu Ser Arg Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95

Ala Arg Gly Gly Trp Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr  
 100 105 110

Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro  
 115 120 125

Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val  
 130 135 140

Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
 145 150 155 160

Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly  
 165 170 175



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Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly  
180 185 190

Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys  
195 200 205

Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys  
210 215 220

Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu  
225 230 235 240

Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu  
245 250 255

Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys  
260 265 270

Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys  
275 280 285

Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu  
290 295 300

Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys  
305 310 315 320

Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys  
325 330 335

Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser  
340 345 350

Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys  
355 360 365

Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln  
370 375 380

TP0063W0-seq1-000001.txt

Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly  
385 390 395 400

Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln  
405 410 415

Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn  
420 425 430

His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys Gly Gly Gly  
435 440 445

Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Ala Gly Asp Val  
450 455 460

Asp Thr Leu Ala Phe Asp Gly Arg Thr Phe Val Glu Tyr Leu Asn Ala  
465 470 475 480

Val Thr Glu Ser Glu Leu Ala Asn Glu Ile Pro Val Glu Lys Ala Leu  
485 490 495

Gln Ser Asn His Phe Glu Leu Ser Leu Arg Thr Glu Ala Thr Gln Gly  
500 505 510

Leu Val Leu Trp Ser Gly Lys Ala Thr Glu Arg Ala Asp Tyr Val Ala  
515 520 525

Leu Ala Ile Val Asp Gly His Leu Gln Leu Ser Tyr Asn Leu Gly Ser  
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Gln Pro Val Val Leu Arg Ser Thr Val Pro Val Asn Thr Asn Arg Trp  
545 550 555 560

Leu Arg Val Val Ala His Arg Glu Gln Arg Glu Gly Ser Leu Gln Val  
565 570 575

Gly Asn Glu Ala Pro Val Thr Gly Ser Ser Pro Leu Gly Ala Thr Gln  
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Leu Asp Thr Asp Gly Ala Leu Trp Leu Gly Gly Leu Pro Glu Leu Pro  
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Val Gly Pro Ala Leu Pro Lys Ala Tyr Gly Thr Gly Phe Val Gly Cys  
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Ala Val Thr Lys Pro Glu Leu Arg Pro Cys Pro Thr Pro  
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<400> 31

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Gly Ser Ser Val Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly Leu Glu  
35 40 45

Trp Ile Gly His Ile Tyr Trp Asp Asp Asp Lys Arg Leu Asn Pro Ser  
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Leu Arg Asn Arg Val Thr Ile Ser Val Asp Thr Ser Lys Asn Gln Phe  
65 70 75 80

Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr  
85 90 95

Cys Ala Arg Arg Ala Ile Thr Thr Val Ile Gly Gly Gly Thr Phe Asp  
100 105 110

Leu Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys

115

120

125

Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly  
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Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro  
 145 150 155 160

Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr  
 165 170 175

Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val  
 180 185 190

Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn  
 195 200 205

Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro  
 210 215 220

Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu  
 225 230 235 240

Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp  
 245 250 255

Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp  
 260 265 270

Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly  
 275 280 285

Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn  
 290 295 300

Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp  
 305 310 315 320

Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro

325

330

335

Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu  
 340 345 350

Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn  
 355 360 365

Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile  
 370 375 380

Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr  
 385 390 395 400

Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys  
 405 410 415

Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys  
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Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu  
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Ser Leu Ser Pro Gly Lys  
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 35 40 45

Asp Thr Ser Lys Leu Ala Arg Gly Val Pro Ser Arg Phe Ser Gly Ser  
50 55 60

Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu  
65 70 75 80

Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Trp Tyr Leu His Pro Leu Thr  
85 90 95

Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala Pro  
100 105 110

Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr  
115 120 125

Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys  
130 135 140

Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu  
145 150 155 160

Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser  
165 170 175

Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala  
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Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe  
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Asn Arg Gly Glu Cys  
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Gly Ser Ser Val Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly Leu Glu  
35 40 45

Trp Ile Gly His Ile Tyr Trp Asp Asp Asp Lys Arg Leu Asn Pro Ser  
50 55 60

Leu Arg Asn Arg Val Thr Ile Ser Val Asp Thr Ser Lys Asn Gln Phe  
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Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr  
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Cys Ala Arg Arg Ala Ile Thr Thr Val Ile Gly Gly Gly Thr Phe Asp  
100 105 110

Leu Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys  
115 120 125

Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly  
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Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro  
145 150 155 160

Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr  
165 170 175

Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val  
180 185 190

Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn  
195 200 205

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Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro  
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Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu  
225 230 235 240

Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp  
245 250 255

Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp  
260 265 270

Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly  
275 280 285

Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn  
290 295 300

Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp  
305 310 315 320

Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro  
325 330 335

Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu  
340 345 350

Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn  
355 360 365

Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile  
370 375 380

Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr  
385 390 395 400

Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys  
405 410 415



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Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys  
420 425 430

Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu  
435 440 445

Ser Leu Ser Pro Gly Lys Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser  
450 455 460

Gly Gly Gly Gly Ser Ala Gly Asp Val Asp Thr Leu Ala Phe Asp Gly  
465 470 475 480

Arg Thr Phe Val Glu Tyr Leu Asn Ala Val Thr Glu Ser Glu Leu Ala  
485 490 495

Asn Glu Ile Pro Val Glu Lys Ala Leu Gln Ser Asn His Phe Glu Leu  
500 505 510

Ser Leu Arg Thr Glu Ala Thr Gln Gly Leu Val Leu Trp Ser Gly Lys  
515 520 525

Ala Thr Glu Arg Ala Asp Tyr Val Ala Leu Ala Ile Val Asp Gly His  
530 535 540

Leu Gln Leu Ser Tyr Asn Leu Gly Ser Gln Pro Val Val Leu Arg Ser  
545 550 555 560

Thr Val Pro Val Asn Thr Asn Arg Trp Leu Arg Val Val Ala His Arg  
565 570 575

Glu Gln Arg Glu Gly Ser Leu Gln Val Gly Asn Glu Ala Pro Val Thr  
580 585 590

Gly Ser Ser Pro Leu Gly Ala Thr Gln Leu Asp Thr Asp Gly Ala Leu  
595 600 605

Trp Leu Gly Gly Leu Pro Glu Leu Pro Val Gly Pro Ala Leu Pro Lys  
610 615 620

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Ala Tyr Gly Thr Gly Phe Val Gly Cys Leu Arg Asp Val Val Val Gly  
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Arg His Pro Leu His Leu Leu Glu Asp Ala Val Thr Lys Pro Glu Leu  
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Arg Pro Cys Pro Thr Pro  
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 <212> DNA  
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