



(86) Date de dépôt PCT/PCT Filing Date: 1999/11/11
(87) Date publication PCT/PCT Publication Date: 2000/05/25
(85) Entrée phase nationale/National Entry: 2001/05/10
(86) N° demande PCT/PCT Application No.: JP 99/06275
(87) N° publication PCT/PCT Publication No.: WO 00/29570
(30) Priorités/Priorities: 1998/11/13 (10/323199) JP;
1998/12/07 (10/346925) JP

(51) Cl.Int.⁷/Int.Cl.⁷ C12N 15/12, A61K 35/76, A61K 31/7088,
A61K 39/395, A61P 25/28, A61P 25/16, A61P 25/14,
A61P 25/08, A01K 67/027, A61K 38/02, A61P 43/00,
A61P 25/00, A61K 48/00, A61K 45/00, C07K 16/18,
C12N 1/21, C07K 14/47

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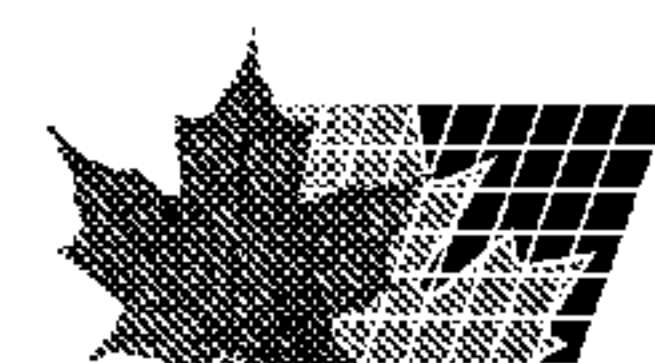
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(54) Titre : NOUVELLE PROTEINE ET SON UTILISATION
(54) Title: NOVEL PROTEIN AND UTILIZATION THEREOF

(57) **Abrégé/Abstract:**

A protein which is expressed particularly in the brain, has a PDZ domain and/or a WW domain and shows an affinity with, in particular, activin receptor and/or activin intracellular signal transducer molecule, its peptide fragment or a salt thereof which are useful in determining a binding protein capable of binding to the above protein and screening a compound inhibiting or promoting the binding of the binding protein to the above-described protein, etc.



ABSTRACT

The protein of the present invention, a partial peptide thereof or a salt thereof, expressed specifically in the brain, having PDZ domains and/or WW domains and
5 having affinity particularly for activin receptors and/or activin intracellular information transmission molecules, is useful for determining a binding protein having an ability to bind to said protein and for screening a compound inhibiting or promoting a binding of the protein
10 etc. of the present invention to the binding protein.

DESCRIPTION

NOVEL PROTEIN AND UTILIZATION THEREOF

5 Technical Field

 The present invention relates to a novel protein having a specific amino acid sequence, having PDZ domains and/or WW domains, and being expressed specifically in the brain, a DNA comprising a DNA region encoding said protein,
10 a process for producing the protein, and use of the protein and the DNA.

Background Art

 Heretofore, a large number of physiologically active
15 substances have been isolated and identified, and their functions are being elucidated. Some of them are known to exhibit various activities in a wide variety of organs or cells. The various activities in a wide variety of organs or cells appear generally via receptors to which the
20 physiologically active substances bind, but there are many cases where whether the combination of the physiologically active substances binding to the receptors is identical in all organs and cells or specific to each organ or cell is not elucidated.

25 Some physiologically active proteins have PDZ domains

and WW domains. These domains are protein-binding domain modules found relatively recently. Accordingly, the in vivo distribution, functions and regulatory mechanism of proteins having PDZ domains and WW domains are still not elucidated in many cases.

The proteins having PDZ domains are considered to play an important role in formation of a backing structure of cell membrane, in formation of a network of cytoskeleton, and in formation of an intracellular signal transmission network developed in an intracellular cortex. In the nervous system, these proteins participate in forming complexes of neurotransmitter receptors, ion channels, etc., thus making them essential for assembly of protein clusters in synapse sites. Further, PDZ proteins whose expression is regulated depending on synaptic plasticity were recently found, and there is the possibility that the PDZ proteins participate in altering the morphology of synapses accompanying plasticity through rearrangement of receptors, and they are considered to participate in construction of a nervous network at a development stage and in in vivo high-order mechanism in the brain. The PDZ domain is similar in certain respects to known peptide-binding domains, but has distinctive features.

The PDZ domain is a domain preserved widely in microorganisms, higher plants and animals, and in many

cases, the PDZ domain recognizes a C-terminal short amino acid sequence of its target proteins, and these target proteins are often membrane-penetrating receptors or channels. The PDZ domain is often found in a form repeated
5 about 2- to 6-times in the same protein, and unlike other domain modules, some of the PDZ domains form homodimers. These are features important for participating in formation of a protein-crosslinked network and microdomains in intercellular adhesion, such as a cellular surface
10 structure and tight junction of synapses etc.

On the other hand, the WW domain is also a domain consisting of about 40 amino acids preserved well in yeasts and mammals, and recognizes and binds to a relatively short amino acid sequence called PY motif rich in proline. As
15 the protein having the WW domains, there are ubiquitin protein ligase, dystrophin, formine-binding protein, etc. On such protein, 1 to 4 WW domains occur in a form linked in series. A majority of those proteins having WW domains have domains participating in interaction with other
20 proteins and enzymatically active domains such as in proteiphostase [phonetic transcription], ras GAP, etc., and it is suggested that the WW domains have very diverse biological functions.

Activin is a regulatory factor promoting secretion of
25 follicle-stimulating hormone (FSH) from the anterior

pituitary. Heretofore, it has been considered that the regulation of secretion of gonadotrophic hormone from the pituitary is based on steroid hormone produced in the gonad, and since activin and its antagonist i.e. inhibin were found, this regulation attracts attention as a new hormone secretion-regulating mechanism in a hypothalamo-hypophysial-germ system. As analysis of the physiological activities of activin proceeds, this system was found to have diverse physiological activities including not only secretory regulation of FSH but also an activity of inducing or inhibiting differentiation of cells in a blood-corpusele system or in reproductive cells, an activity of maintaining the viability of nerve cells, etc., but many features of this mechanism are not elucidated in detail.

As a means of elucidating the physiological activities of a protein having PDZ domains and/or WW domains expressed particularly in the brain, receptors (e.g., activin receptors) having an ability to bind to said protein, and intracellular information transmission molecules (e.g., Smad) as well as of elucidating detailed molecular mechanisms etc. for neurotrophic factor-like activities and for inhibiting differentiation of cells in nervous-system tissues in an activin-activin receptor system, the object of the present invention is to provide a method for

isolating said novel protein, a method for detecting the same, a DNA comprising a gene for said novel protein, a process for producing a protein encoded by the gene for said novel protein, and use of the DNA and the protein.

5

Disclosure of Invention

To solve the problem described above, the present inventors made extensive study, and as a result, they screened a binding protein from a mouse brain cDNA by means
10 of a yeast two-hybrid method in which an intracellular region in a mouse activin IIA-N receptor protein shown in SEQ ID NO: 1 was used as a bait, thus acquiring a cDNA clone which could be confirmed to bind to activin IIA-N receptors even in COS7 cells. Further, they isolated and
15 analyzed a cDNA clone containing a full-length gene encoded by said clone, and found that said gene is a gene containing a region encoding 5 PDZ domains and 2 WW domains. Said protein encoded by this gene is a protein factor having a plurality of domains for protein-protein
20 interaction, and this protein binds, via PDZ domains out of these domains, to 1) intracellular regions in activin receptors, resulting in inhibiting activin from transmitting information to cells, but not to 2) intracellular regions of receptors for other cytokines,
25 thus exerting no influence on their transmission of

information to cells. Further, the present inventors found that this protein binds via other WW domains to activin intracellular transmission molecules thereby inhibiting activin from transmitting information to cells.

5 Further, the present inventors extracted poly(A) ⁺RNA from a wide variety of mouse organs, and used a DNA shown in SEQ ID NO: 2, 3, or 4 as a probe to examine expression of the novel protein of the present invention by Northern hybridization, and as a result, they found that high-level
10 expression thereof occurs particularly in the brain, as shown in Fig. 23. From this, the activin-activin receptor information transmission system to which said novel protein was added is suggested to govern regulation of multiplication and differentiation of cerebral cells and
15 found to be applicable to diagnosis, treatment, etc. of diseases in the cerebral and nerve system.

As further examination on the basis of these findings, the present inventors arrived at the present invention.

That is, the present invention provides:

- 20 1. A protein which has an amino acid sequence identical to or substantially identical to an amino acid sequence represented by SEQ ID NO: 5, or a salt thereof,
2. A protein which has an amino acid sequence identical to or substantially identical to an amino acid sequence
25 represented by SEQ ID NO: 6, or a salt thereof,

3. The protein according to item 1 or 2, which has PDZ domains and WW domains and is expressed specifically in the brain and has an ability to bind to activin receptors and/or activin intracellular information transmission molecules,

4. The protein according to item 3, wherein the activin intracellular information transmission molecule is Smad3,

5. The protein according to item 1 or 2, which has 5 PDZ domains and 2 WW domains and is expressed specifically in the brain and has an ability to bind to activin receptors and Smad3,

6. A partial peptide of the protein according to item 1, a partial peptide of the protein according to item 2, or a salt thereof,

7. A recombinant DNA comprising a DNA which comprises a nucleotide sequence encoding the protein according to item 1 or the protein according to item 2,

8. The DNA according to item 7, which has a nucleotide sequence represented by SEQ ID NO: 7, a nucleotide sequence represented by SEQ ID NO: 8 or a nucleotide sequence hybridizing therewith under high stringent conditions.

9. A recombinant DNA which comprises a DNA having a nucleotide sequence encoding the partial peptide according to item 6.

10. A recombinant vector which comprises the DNA according

to item 7,

11. A transformant comprising the recombinant vector according to item 10,

12. A method for producing the protein according to item 1,
5 the protein according to item 2 or a salt thereof, which comprises culturing the transformant according to item 11 to produce and accumulate the protein according to item 1 or the protein according to item 2 is formed and accumulated, and then recovering the product,

10 13. An antibody against the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof,

14. A method for quantifying the protein according to item 1, the protein according to item 2, the partial peptide
15 according to item 6 or a salt thereof, which comprises allowing a test solution containing the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof and the labeled protein according to item 1, the labeled protein
20 according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof to react competitively with the antibody according to item 13,

15. A method for determining a binding protein to the protein according to item 1, the protein according to item
25 2, the partial peptide according to item 6 or a salt

thereof, which comprises using the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof,

16. The method according to item 15, which comprises
5 introducing (1) an expression vector which fuses the protein according to item 1, the protein according to item 2 or the partial peptide according to item 6 with a DNA-binding region of a transcriptional factor and (2) a fusion library between a gene encoding a test protein and a
10 transcription-activating region, into a host cell having a reporter gene maintaining a region binding to the transcriptional factor on a promoter, and measuring a change in the amount of the expressed reporter gene which is increased by a binding of the protein according to item
15 1, the protein according to item 2 or the partial peptide according to item 6 to the test compound,

17. A protein or a salt thereof which binds to the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof,
20 which is obtained by the method according to item 15,

18. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to the
25 protein according to item 17 or a salt thereof, activin

receptors or activin intracellular information transmission molecules, which comprises using the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof,

- 5 19. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to activin receptors or activin intracellular information transmission
- 10 molecules, which comprises comparing the case where the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof is contacted with activin receptors or activin intracellular information
- 15 transmission molecules, with the case where the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof and a test compound are contacted with activin receptors or activin intracellular information
- 20 transmission molecules, by measuring the amount of the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof bound to the activin receptors or activin intracellular information transmission
- 25 molecules in both the cases,

20. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to the protein according to item 17 or a salt thereof, which comprises comparing the case where the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof is contacted with the protein according to item 17 or a salt thereof, with the case where the labeled protein according to item 1, the labeled protein according to item 2, [the labeled partial peptide according to item 6] or a labeled salt thereof and a test compound are contacted with the protein according to item 17 or a salt thereof, by measuring the amount of the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof bound to the protein according to item 17 or a salt thereof in both the cases,

21. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission

molecules, which comprises comparing the case where the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof is introduced into cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, with the case where the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof and a test compound are introduced into cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the amount of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof bound to the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules in the cells in both the cases,

22. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof to the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which comprises

comparing the case where the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof is contacted with a membrane fraction of cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, with the case where the labeled protein according to item 1, the labeled protein according to item 2, the labeled partial peptide according to item 6 or a labeled salt thereof and a test compound are contacted with a membrane fraction of cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the amount of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof bound to the membrane fraction of the cells in both the cases,

23. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which comprises comparing the case where the protein according to item 1, the protein according to item

2, [the partial peptide according to item 6] or a salt thereof is introduced into cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, with the case where the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof and a test compound are introduced into cells expressing the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the cell-stimulating activity via the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules in both the cases,

24. The method for determining a protein according to item 16 or the screening method according to any one of items 18 to 23, which comprises using the two-hybrid method,

25. A kit for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to item 1, the protein according to item 2, the partial peptide according to item 6 or a salt thereof to the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which comprises the protein according to item 1, the protein according to item 2, the partial peptide

according to item 6 or a salt thereof,

26. A compound or a salt thereof which inhibits or promotes
a binding of the protein according to item 1, the protein
according to item 2, the partial peptide according to item
5 6 or a salt thereof to the protein according to item 17 or
a salt thereof, activin receptors or activin intracellular
information transmission molecules, which is obtained by
the screening method according to any one of items 18 to 23
or by the screening kit according to item 25,

10 27. A pharmaceutical composition comprising the protein
according to item 17, the compound according to item 26 or
a salt thereof,

28. The pharmaceutical composition according to item 27,
which is an agent for preventing or treating Alzheimer's
15 disease, Parkinson's disease, epilepsy or Huntington's
chorea,

29. A method for preventing and treating abnormalities in
nerve cells or cerebral diseases correlated with the
protein according to item 17 or a salt thereof, activin
20 receptors or activin intracellular information transmission
molecules, which administering an effective amount of the
protein according to item 17, the compound according to
item 26 or a salt thereof to mammals, and

30. Use of the protein according to item 17, the compound
25 according to item 26 or a salt thereof for preparing an

agent for preventing and treating abnormalities in nerve cells or cerebral diseases correlated with the protein according to item 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules.

5 Further, the present invention provides:

31. The protein according to item 1 or a salt thereof, wherein the protein comprises an amino acid sequence represented by SEQ ID NO: 5, an amino acid sequence in which 1 or 2 or more (preferably 2 to 10) amino acid
10 sequence in an amino acid sequence represented by SEQ ID NO: 5 excluding an amino acid sequence represented by SEQ ID NO: 6 are deleted, an amino acid sequence in which 1 or 2 or more (preferably 2 to 10) amino acid sequence are added or inserted to an amino acid sequence represented by
15 SEQ ID NO: 5 excluding an amino acid sequence represented by NO: 6, or an amino acid sequence in which 1 or 2 or more (preferably 2 to 10) amino acids in an amino acid sequence represented by SEQ ID NO: 5 excluding an amino acid sequence represented by NO: 6 are substituted with other
20 amino acids,

32. The two-hybrid method according to item 24, wherein yeasts are used,

33. A compound obtained by the screening method according to any one of items 19 to 23 or a salt thereof,

25 34. A pharmaceutical composition comprising the compound

according to item 33 or a salt thereof,

35. Non-human mammals having the extraneous DNA according to item 7 or a mutated DNA thereof,

36. The non-human mammals according to item 35 wherein the
5 non-human mammals are rodents,

37. The non-human mammals according to item 35 wherein the rodents are mice or rats, and

38. A recombinant vector comprising the DNA according to item 7 or a mutated DNA thereof and capable of expression
10 in mammals.

Brief Description of Drawings

Fig. 1 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as
15 well as an amino acid sequence (SEQ ID NO: 5) encoded thereby.

Fig. 2 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 5) encoded
20 thereby. Fig. 2 is a continuation of Fig. 1.

Fig. 3 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 5) encoded
thereby. Fig. 3 is a continuation of Fig. 2.

25 Fig. 4 shows the entire nucleotide sequence of cDNA

encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 5) encoded thereby. Fig. 4 is a continuation of Fig. 3.

Fig. 5 shows the entire nucleotide sequence of cDNA
5 encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 5) encoded thereby. Fig. 5 is a continuation of Fig. 4.

Fig. 6 shows the entire nucleotide sequence of cDNA
encoding the novel protein of the present invention, as
10 well as an amino acid sequence (SEQ ID NO: 5) encoded thereby. Fig. 6 is a continuation of Fig. 5.

Fig. 7 shows the entire nucleotide sequence of cDNA
encoding the novel protein of the present invention, as
well as an amino acid sequence (SEQ ID NO: 5) encoded
15 thereby. Fig. 7 is a continuation of Fig. 6.

Fig. 8 shows the entire nucleotide sequence of cDNA
encoding the novel protein of the present invention, as
well as an amino acid sequence (SEQ ID NO: 5) encoded
thereby. Fig. 8 is a continuation of Fig. 7.

20 Fig. 9 shows the entire nucleotide sequence of cDNA
encoding the novel protein of the present invention, as
well as an amino acid sequence (SEQ ID NO: 5) encoded
thereby. Fig. 9 is a continuation of Fig. 8.

Fig. 10 shows the entire nucleotide sequence of cDNA
25 encoding the novel protein of the present invention, as

well as an amino acid sequence (SEQ ID NO: 5) encoded thereby. Fig. 10 is a continuation of Fig. 9.

Fig. 11 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 5) encoded thereby. Fig. 11 is a continuation of Fig. 10.

Fig. 12 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby.

Fig. 13 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 13 is a continuation of Fig. 12.

Fig. 14 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 14 is a continuation of Fig. 13.

Fig. 15 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 15 is a continuation of Fig. 14.

Fig. 16 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded

thereby. Fig. 16 is a continuation of Fig. 15.

Fig. 17 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded
5 thereby. Fig. 17 is a continuation of Fig. 16.

Fig. 18 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 18 is a continuation of Fig. 17.

10 Fig. 19 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 19 is a continuation of Fig. 18.

Fig. 20 shows the entire nucleotide sequence of cDNA
15 encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 20 is a continuation of Fig. 19.

Fig. 21 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as
20 well as an amino acid sequence (SEQ ID NO: 6) encoded thereby. Fig. 21 is a continuation of Fig. 20.

Fig. 22 shows the entire nucleotide sequence of cDNA encoding the novel protein of the present invention, as well as an amino acid sequence (SEQ ID NO: 6) encoded
25 thereby. Fig. 22 is a continuation of Fig. 21.

Fig. 23 shows the analysis result of expression by Northern hybridization.

Fig. 24 shows the result of examination of the specific interaction of the novel protein of the present invention with Smad3. ARIP1 on the abscissa indicates the novel protein of the present invention. The interaction on the ordinate shows the strength of luciferase activity.

Best Mode for Carrying Out the Invention

10 The protein comprising an amino acid sequence substantially identical to the amino acid sequence represented by SEQ ID NO: 5 or NO: 6 of the present invention may be a protein derived from cells (e.g., hepatic cells, spleen cells, nerve cells, glia cells, 15 pancreatic β -cells, bone marrow cells, mesangial cells, Langerhans' cells, epidermal cells, epithelial cells, endothelial cells, fibroblasts, fibrous cells, muscular cells, fat cells, immune cells (e.g., macrophage, T cell, B cell, natural killer cell, mast cell, neutrophils, 20 basophils, eosinophils, monocyte), megakaryocyte, synovial membrane cells, soft-bone cells, bone cells, osteoblast, osteoclast, mammary gland cells or interstitial cells, or their precursor cells, stem cells or cancer cells) or any tissues having such cells, for example, the brain and each 25 site of brain (e.g., olfactory bulb, tonsil nuclei,

cerebral basal bulb, hippocampus, thalamus, hypothalamus, cerebral cortex, medulla bulb, cerebellum), spinal cord, pituitary, stomach, pancreas, kidney, liver, gonads, thyroid glands, galls, bone marrow, adrenals, skin, muscles, lungs, digestive tracts (e.g., large and small intestines), blood vessels, heart, thymus, spleen, salivary glands, peripheral blood, prostate, testicles, ovary, placenta, uterus, bone, joints, skeletal muscles, or cells in the hemocyte system or their cultured cell strain (particularly in the brain), from humans or warm-blooded animals (e.g., guinea pig, rat, mouse, chicken, rabbit, pig, sheep, cow, monkey, etc.), as well as a recombinant protein or a synthetic protein.

As the protein comprising an amino acid sequence substantially identical to an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, an amino acid sequence having about 50 % or more, preferably about 70 % or more, more preferably about 90 % or more, and most preferably about 95 % or more homology with an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, etc., are exemplified.

The protein of the present invention comprising an amino acid sequence substantially identical to an amino acid sequence represented by SEQ ID NO: 5 or NO: 6 is preferably, for example, a protein having an amino acid

sequence substantially identical to the amino acid sequence represented by SEQ ID NO: 5 or NO: 6 and having substantially identical physiological properties with those of a protein having the amino acid sequence represented by
5 SEQ ID NO: 5 or NO: 6.

As the substantially identical physiological properties, for example, qualitative elements such as receptor affinity, signal transmission ability, specificity of distribution of expression in organs, etc. are
10 exemplified. The terms "substantially identical" mean that these physiological activities are biologically or physiologically homogenous. Therefore, although it is preferable that the activities such as the strength of receptor affinity are equal (e.g., about 0.1- to 20-fold,
15 preferably about 0.5- to 2-fold), quantitative elements such as a degree of these activities and the molecular weight of the protein may be different.

Measurement of the receptor affinity can be carried out according to the per se known method, for example, by
20 the screening method described later.

In addition, as the protein of the present invention, for example, a mutein such as a protein comprising an amino acid sequence in which 1 or 2 or more (preferably about 1 to 30, more preferably about 1 to 10, further preferably a
25 few) amino acids are deleted, an amino acid sequence in

which 1 or 2 or more (preferably about 1 to 30, more preferably about 1 to 10, further preferably a few) amino acids are added or inserted to an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, or an amino acid sequence in which 1 or 2 or more (preferably about 1 to 30, more preferably about 1 to 10, further preferably a few) amino acids are substituted with other amino acids, or a protein containing a combination of these amino acid sequences, etc. are exemplified.

10 In proteins in the present specification, a left end is an N-terminal (amino terminal) and a right end is a C-terminal (carboxyl terminal) according to the convention of the peptide display. Although the proteins of the present invention including a protein comprising an amino acid
15 sequence represented by SEQ ID NO: 5 or NO: 6 have usually carboxyl group ($-\text{COOH}$) or carboxylate ($-\text{COO}^-$) as a C-terminal, they may have an amide ($-\text{CONH}_2$) or an ester ($-\text{COOR}$) as a C-terminal.

As R in ester, for example C_{1-6} alkyl group such as
20 methyl, ethyl, n-propyl, isopropyl and n-butyl, C_{3-8} cycloalkyl group such as cyclopentyl and cyclohexyl, C_{6-12} aryl group such as phenyl and α -naphthyl, and C_{6-12} aryl- C_{1-2} alkyl group such as benzyl, phenetyl and α -naphthylmethyl, as well as a pivaloyloxymethyl group used generally in an
25 oral ester are used.

When the protein of the present invention has carboxyl group (or carboxylate) at a position other than a C-terminal, proteins in which carboxyl group is amidated or esterified are also included in the protein of the present invention. As an ester used in this case, for example, an ester at a C-terminal described above is used.

Further, the protein of the present invention includes proteins in which an amino group of a methionine residue of a N-terminal amino acid residue is protected with a protecting group (e.g., C₁₋₆ acyl group such as formyl group and acetyl group) in the aforementioned proteins, proteins in which a N-terminal glutamic acid residue formed by cutting in the living body is converted into pyroglutamine, proteins in which for example -OH, COOH, NH₂, -SH, etc. on a side chain of an intramolecular amino acid is protected with a suitable protecting group (e.g., C₁₋₆ acyl group such as formyl group and acetyl group), and conjugated proteins such as glycoprotein having sugar chains bound thereto.

As an embodiment of the protein of the present invention, for example, a protein comprising an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, etc. are used.

The protein of the present invention has PDZ domains and/or WW domains, preferably 5 PDZ domains and/or 2 WW domains, and is expressed specifically in the brain. Then,

the protein of the present invention binds via the PDZ domains to an activin receptor isolated as a serine/threonine kinase-type receptor, or via the WW domains to an activin intracellular information transmission molecule.

The activin receptor may be of any subtype of activin receptor, but particularly activin receptor IIA-N etc. are preferably used.

As the activin intracellular information transmission molecule, for example, Smad such as Smad1, 2, 3, 4, 5, 6, 7, etc. is exemplified, and the protein of the present invention strongly binds particularly to Smad3.

As the protein of the present invention, in particular, a protein having 5 PDZ domains and 2 WW domains, being expressed specifically in the brain and having an ability to bind to activin receptors and/or Smad3 is preferably used.

As a partial peptide of the protein of the present invention, any peptides may be used as long as they are partial peptides of the protein of the present invention described above, having the physiological activities possessed by the protein of the present invention, for example activities such as receptor affinity and an ability to transmit signal information, and specificity of distribution of expression thereof in organs. For example,

a peptide having 100 or more, preferably 250 or more, more preferably 350 or more, further preferably 500 or more and most preferably 800 or more amino acids from the amino acid sequence constituting the protein of the present invention and having receptor affinity and an ability to transmit signal information, etc., are used.

Further, in the partial peptide of the present invention, 1 or 2 or more (preferably about 1 to 10, more preferably a few) amino acids in the amino acid sequence thereof may be deleted, 1 or 2 or more (preferably about 1 to 20, more preferably about 1 to 10, further preferably a few) amino acids may be added to the amino acid sequence thereof, or 1 or 2 or more (preferably about 1 to 10, more preferably a few, further preferably about 1 to 5) amino acids in the amino acid sequence thereof may be substituted with other amino acids.

Although the partial peptide of the present invention has usually carboxyl group ($-\text{COOH}$) or carboxylate ($-\text{COO}^-$) as a C-terminal, it may have an amide ($-\text{CONH}_2$) or an ester ($-\text{COOR}$) as a C-terminal, as with the protein of the present invention described above.

Further, the partial peptide of the present invention, similar to the protein of the present invention described above, includes peptides in which an amino group of a methionine residue as a N-terminal amino acid residue is

protected with a protecting group, peptides in which a N-terminal glutamyl group formed by cutting in the living body is converted into pyroglutamine, peptides in which a substituent group on a side chain of an intramolecular amino acid is protected with a suitable protecting group, and conjugated peptides such as glycopeptides having sugar chains bound thereto.

The partial peptide of the present invention preferably has an ability to bind to activin receptors and/or the above activin intracellular information transmission molecules (particularly Smad3).

As a salt of the protein of the present invention or of a partial peptide thereof, physiologically acceptable acid addition salts are particularly preferable. As such salts, for example, salts with inorganic acids (e.g., hydrochloric acid, phosphoric acid, hydrobromic acid, sulfuric acid) or salts with organic acids (e.g., acetic acid, formic acid, propionic acid, fumaric acid, maleic acid, succinic acid, tartaric acid, citric acid, malic acid, oxalic acid, benzoic acid, methanesulfonic acid, benzenesulfonic acid), etc., are used. Further, salts with inorganic bases (e.g., salts with alkali metals such as sodium and potassium, alkaline earth metals such as calcium and magnesium, or aluminum, ammonium, etc.), salts with organic bases (e.g., trimethylamine, triethylamine,

pyridine, picoline, 2,6-lutidine, ethanolamine, diethanolamine, triethanolamine, cyclohexyl amine, dicyclohexyl amine, N,N'-dibenzylethylene diamine, etc.), etc., are also used.

5 The protein of the present invention or a salt thereof can be produced from cells or tissues of a human being or a warm blood mammal described above by the method of purifying proteins known per se, or can be prepared by culturing a transformant containing a DNA encoding a
10 protein described below. Alternatively, it can also be prepared according to a method of synthesizing a peptide described below.

For production from human or mammalian tissues or cells, the human or mammalian tissues or cells are
15 homogenized and then extracted with e.g. an acid, and the extract can be subjected to purification and isolation by a combination of chromatographic techniques such as reverse-phase chromatography, ion-exchange chromatography, etc.

For synthesis of the protein of the present invention
20 or a partial peptide thereof or a salt thereof or amide derivatives thereof, usually commercially available resin for protein synthesis can be used. Such resin includes, for example, chloromethyl resin, hydroxymethyl resin, benzhydryl amine resin, aminomethyl resin,
25 4-benzyloxybenzyl alcohol resin, 4-methylbenzhydryl amine

resin, PAM resin, 4-hydroxymethyl phenylacetamidemethyl resin, polyacrylamide resin, 4-(2',4'-dimethoxyphenylhydroxymethyl)phenoxy resin, 4-(2',4'-dimethoxyphenyl-Fmoc aminoethyl)phenoxy resin, etc.

5 On the resin described above, each amino acid with the α -amino group and side-chain functional group properly protected is condensed sequentially in accordance with the sequence of the desired protein by the per se known condensation methods. At the end of the reaction, the
10 protein is cleaved off from the resin while various protecting groups are removed, and the product is subjected to a reaction of forming intramolecular disulfide bonds in a highly dilute solution to give the desired protein or an amide thereof.

15 A wide variety of activating reagents usable for protein synthesis can be used for condensation of the protected amino acids described above, and carbodiimides are particularly preferable. Examples of such carbodiimides include DCC, N,N'-diisopropylcarbodiimide, N-ethyl-N'-(3-dimethylaminopropyl)carbodiimide, etc.
20 For activation by these reagents, the protected amino acids along with racemization inhibitors (e.g., HOBt, HOObt) can be added to the resin directly or after the protected amino acids were previously activated as symmetric acid
25 anhydrides or HOBt esters or HOObt esters.

The solvent used for activation of each protected amino acid or for condensation thereof with the resin can be selected as necessary from those solvents known to be usable in protein condensation reaction. Examples of such solvent include acid amides such as N,N-dimethylformamide, N,N-dimethylacetamide and N-methylpyrrolidone, halogenated hydrocarbons such as methylene chloride and chloroform, alcohols such as trifluoroethanol, sulfoxides such as dimethyl sulfoxide, ethers such as pyridine, dioxane and tetrahydrofuran, nitriles such as acetonitrile and propionitrile, esters such as methyl acetate and ethyl acetate, or a suitable mixture thereof. The reaction temperature is usually selected as necessary within the range known to be usable in the reaction of forming protein bonds, and usually the reaction temperature is selected within the range of about -20 °C to 50 °C. The activated amino acid derivatives are used usually in excess (1.5- to 4-fold). When their condensation is insufficient, their sufficient condensation is achieved as a result of a test using ninhydrin reaction by repeatedly carrying out the condensation reaction without conducting elimination of the protecting groups. When their sufficient condensation is not achieved even by repeatedly carrying out the reaction, the unreacted amino acids are acetylated with acetic anhydride or acetyl imidazole.

The protecting groups for amino groups in the starting materials include, for example, Z, Boc, t-pentyloxycarbonyl, isobornyloxycarbonyl, 4-methoxybenzyloxycarbonyl, Cl-Z, Br-Z, adamantyloxycarbonyl, trifluoroacetyl, phthaloyl, formyl,
5 2-nitrophenylsulphenyl, diphenylphosfinothioyl, Fmoc, etc.

The carboxyl group can be protected by, for example, alkyl esterification (e.g., straight-chain, branched or cyclic alkyl esterification such as methyl, ethyl, propyl, butyl, tert-butyl, cyclopentyl, cyclohexyl, cycloheptyl,
10 cyclooctyl or 2-adamantyl esterification), aralkyl esterification (e.g., benzyl esterification, 4-nitrobenzyl esterification, 4-methoxybenzyl esterification, 4-chlorobenzyl esterification, benzhydryl esterification), phenacyl esterification, benzyloxycarbonylhydrazidation,
15 tert-butoxycarbonylhydrazidation, tritylhydrazidation, etc.

The hydroxyl group in serine can be protected by, for example, esterification or etherification. A suitable group used in this esterification includes, for example, lower alkanoyl groups such as acetyl group, allyl groups
20 such as benzoyl group, and carbonic acid-derived groups such as benzyloxycarbonyl group and ethoxycarbonyl group. A suitable group for etherification includes, for example, a benzyl group, tetrahydropyranyl group, t-butyl group, etc.

The protecting group used for the phenolic hydroxyl group in tyrosine includes, for example, Bzl, Cl₂-Bzl, 2-

25

nitrobenzyl, Br-Z, tert-butyl, etc.

The protecting group used for imidazole in histidine includes, for example, Tos, 4-methoxy-2,3,6-trimethylbenzenesulfonyl, DNP, benzyloxymethyl, Bum, Boc, 5 Trt, Fmoc, etc.

Activated carboxyl groups in the starting materials include, for example, the corresponding acid anhydrides, azides and active esters (i.e. esters with alcohols such as pentachlorophenol, 2,4,5-trichlorophenol, 2,4-dinitrophenol, 10 cyanomethyl alcohol, p-nitrophenol, HONB, N-hydroxysuccinamide, N-hydroxyphthalimide and HOBt). The activated amino groups in the starting materials include, for example, the corresponding phosphoric acid amides.

Examples of methods for removing (leaving) the 15 protecting groups include catalytic reduction in a hydrogen stream in the presence of a catalyst such as Pd-black or PC-carbon, acid treatment with anhydrous hydrogen fluoride, methane sulfonic acid, trifluoromethane sulfonic acid, trifluoroacetic acid or a mixed solution thereof, base 20 treatment using diisopropylethylamine, triethylamine, piperidine or piperazine, and reduction using sodium in liquid ammonia. The leaving reaction by the acid treatment described above is carried out generally at a temperature of about -20 °C to 40 °C, and it is effective in the acid 25 treatment to add a cation scavenger such as anisole, phenol,

thioanisole, m-cresol, p-cresol, dimethylsulfide, 1,4-butanedithiol and 1,2-ethanedithiol. A 2,4-dinitrophenyl group used as a protecting group for imidazole in histidine can also be removed by treatment with thiophenol, while a
5 formyl group used as a protecting group for indole in tryptophan can be removed not only by deprotection by acid treatment in the presence of 1,2-ethanedithiol or 1,4-butanedithiol, but also by alkali treatment using dilute sodium hydroxide solution or dilute ammonia.

10 Protection and protecting groups for functional groups which should not participate in the reaction of the starting materials, elimination of the protecting groups, and activation of functional groups participating in the reaction can be selected as necessary from known groups or
15 known means.

Another method of obtaining an amide derivative of the protein or a partial peptide thereof includes, for example, protecting the α -carboxyl group of a C-terminal amino acid by amidation, then extending a peptide (protein) chain at
20 the side of the amino group until it attains desired chain length, and thereafter producing a protein (partial peptide) of said peptide chain from which only the protecting group for the N-terminal α -amino group was removed and a protein (partial peptide) of said peptide
25 chain from which only the protecting group for the C-

terminal carboxyl group was removed, followed by condensation both the proteins in the mixed solvent as described above. The details of the condensation reaction are the same as described above. The protected protein
5 obtained by condensation is purified, and every protecting group is removed by the method described above, whereby the desired crude protein (partial peptide) can be obtained. This crude protein (partial peptide) is purified by a wide variety of known purification techniques, and by
10 lyophilizing its major fraction, the desired amide derivative of the protein (partial peptide) can be obtained.

To obtain an ester derivative of the protein or a partial peptide thereof, for example the α -carboxyl group of a C-terminal amino acid is condensed with desired
15 alcohol to form an amino acid ester from which the desired ester derivative of the protein (partial peptide) can be obtained in the same manner as for the amide derivative of the protein (partial peptide).

The partial peptide of the present invention or a salt
20 thereof can be produced according to a per se known peptide synthesis method or by cleaving the protein of the present invention with a suitable peptidase. For example, the peptide synthesis method may be the solid- or liquid-phase synthesis method. That is, the desired peptide can be
25 obtained by condensation of a partial peptide or amino

acids capable of constituting the partial peptide of the present invention with the remainder, followed by elimination of protecting groups if any from the product. As the known condensation method and the elimination of the protecting groups, mention is made of e.g. the methods described in (1) to (5) below:

(1) M. Bodanszky and M. A. Ondetti, Peptide Synthesis, Interscience Publisher, New York (1966);

(2) Schroeder and Luebke, The Peptide, Academic Press, New York (1965);

(3) Nobuo Izumiya et al., Basis and Experiments in Peptide Synthesis (in Japanese), Maruzen Co., Ltd. (1975);

(4) Haruaki Yajima and Shunpei Sakakibara, Biochemical Experimental Course 1, Protein Chemistry IV, 205 (1977);

and

(5) Haruaki Yajima (supervisor), Development of Medicines, a second series, vol. 14, Peptide Synthesis, Hirokawashoten.

After the reaction, the protein of the present invention or a partial peptide thereof can be isolated and purified by a combination of conventional purification techniques such as solvent extraction, distillation, column chromatography, liquid chromatography and recrystallization. If said protein or a partial peptide thereof is obtained in a free form by these methods, the product can be converted into a suitable salt by a known method, or if the product

is obtained in a salt form, it can be converted into a free form by a known method.

As the DNA encoding the protein of the present invention, any DNAs may be used as long as they comprise a nucleotide sequence encoding the protein of the present invention described above. Further, the DNA may be any of a genomic DNA, a genomic DNA library, a cDNA derived from the aforementioned cells or tissues, a cDNA isolated by the per se known method from a cDNA library derived from the aforementioned cells or tissues, and a synthetic DNA.

A vector used for a library may be any of bacteriophage, plasmid, cosmid and phagemide. In addition, amplification can be performed directly by Reverse Transcriptase Polymerase Chain Reaction (abbreviated hereinafter to RT-PCR method) using total RNA or mRNA fraction prepared from the aforementioned cells or tissues.

As the DNA encoding the protein of the present invention, for example, (1) a DNA comprising a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8 or (2) a DNA comprising a nucleotide sequence hybridizing under high stringent conditions with a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8 and encoding a protein having substantially homogenous physiological activities (for example activities such as receptor affinity and signal transmission ability, and specificity of distribution of

expression thereof in organs) with those of a protein comprising an amino acid sequence represented by SEQ ID NO: 5 or NO: 6 may be used.

As the DNA which can hybridize, under high stringent
5 conditions, with a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8, for example, a DNA comprising a nucleotide sequence having about 70 % or more, preferably about 80 % or more, more preferably about 90 % or more and most preferably about 95 % or more homology with a
10 nucleotide sequence represented by SEQ ID NO: 7 or NO: 8, etc. are used.

Hybridization can be carried out according to a per se known method or its analogous method, for example a method described in Molecular Cloning, 2nd ed. (J. Sambrook et al.,
15 Cold Spring Harbor Lab. Press, 1989). If a commercial library is used, hybridization can be carried out according to the manufacture's instructions. Preferably, hybridization can be conducted under high stringent conditions.

20 The high stringent conditions refer to those conditions under which the concentration of sodium is about 19 to 40 mM, preferably about 19 to 20 mM, and the temperature is about 50 to 70 °C, preferably about 60 to 65 °C. In particular, the case where the concentration of
25 sodium is about 19 mM and the temperature is about 65 °C is

the most preferable.

More specifically, as a DNA encoding a protein comprising an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, a DNA comprising a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8, etc. are used.

As a DNA encoding the partial protein of the present invention, any DNAs may be used as long as they comprise a nucleotide sequence encoding the partial protein of the present invention described above. Further, said DNA may be any of a genomic DNA, a genomic DNA library, a cDNA derived from the aforementioned cells or tissues, a cDNA isolated by the per se known method from a cDNA library derived from the aforementioned cells or tissues, and a synthetic DNA.

As a DNA encoding the partial peptide of the present invention, for example, (1) a DNA comprising a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8 or (2) a DNA comprising a nucleotide sequence hybridizing under high stringent conditions with a nucleotide sequence represented by SEQ ID NO: 7 or NO: 8 and encoding a protein having substantially homogenous physiological activities with those of a protein comprising an amino acid sequence represented by SEQ ID NO: 5 or NO: 6, etc. are used.

As the hybridization method and high stringent conditions, those described above are used.

As a means of cloning the DNA encoding the protein of the present invention or a partial peptide thereof (hereinafter referred to collectively as the protein of the present invention), it is possible to use amplification by the PCR method using synthetic DNA primers having a nucleotide sequence encoding a partial sequence of the protein of the present invention as well as isolation by hybridizing, with a labeled DNA fragment or synthetic DNA encoding a part or the whole of the protein of the present invention, a DNA integrated in a suitable vector or a genomic DNA, a genomic DNA library, a cDNA derived from the aforementioned cells or tissues, a cDNA isolated by the per se known method from a cDNA library derived from the aforementioned cells or tissues. Hybridization can be carried out according to a method described in, for example, Molecular Cloning, 2nd ed. (J. Sambrook et al., Cold Spring Harbor Lab. Press, 1989). If a commercial library is used, hybridization can be carried out according to the manufacture's instructions.

Conversion of DNA into a nucleotide sequence can be carried out by a per se known method, for example by the Gapped duplex method or Kunkel method or its analogous method by using a known kit such as MutantTM-G (Takara Shuzo Co., Ltd.) or MutantTM-K (Takara Shuzo Co., Ltd.).

A DNA encoding a cloned protein can be used as it is

depending on the object, or if desired, by digesting it with a restriction enzyme or adding a linker thereto. The DNA may have ATG as a translation initiation codon at the 5'-terminal thereof and TAA, TGA or TAG as a translation
5 termination codon at the 3'-terminal thereof. These translation initiation and termination codons can also be added to the DNA via a suitable synthetic DNA adaptor.

An expression vector for the protein of the present invention can be produced for example by (A) cutting the
10 desired DNA fragment off from the DNA encoding the protein of the present invention and then (B) ligating said DNA fragment to a region downstream from a promoter in a suitable expression vector.

The vector used includes E. coli-derived plasmid (e.g.,
15 pBR322, pBR325, pUC12, pUC13), Bacillus subtilis-derived plasmid (e.g., pUB110, pTP5, pC194), yeast-derived plasmid (e.g., pSH19, pSH15), bacteriophage such as λ -phage, and animal viruses such as retrovirus, vaccinia virus and Baculovirus, as well as pA1-11, pXT1, pRc/CMV, pRc/RSV,
20 pcDNAI/Neo, etc.

The promoter used in the present invention may be any suitable promoter compatible with a host used for expression of the gene. For example, when animal cells are used as the host, mention is made of SV40-derived promoter,
25 retrovirus promoter, metallothionein promoter, heat shock

promoter, cytomegalovirus promoter, SR α promoter, etc. Among these, cytomegalovirus promoter, SR α promoter, etc. are preferably used. It is preferable to use trp promoter, lac promoter, recA promoter, λ P_L promoter, lpp promoter, etc. for microorganisms of the genus Escherichia as the host, SP01 promoter, SP02 promoter, penP promoter, etc. for microorganisms of the genus Bacillus as the host, and PH05 promoter, PGK promoter, GAP promoter, ADH promoter, etc. for yeasts as the host. When insect cells are used as the host, polyhedron promoter, P10 promoter, etc. are preferable.

The expression vector may contain an enhancer, a splicing signal, a poly A-added signal, a selective marker, an SV40 origin of replication (also referred to hereinafter as SV40 ori), etc. in addition to the promoter described above. The selective marker includes, for example, dihydrofolate reductase (also referred to hereinafter as dhfr) gene [methotrexate (MTX) resistance], ampicillin resistance gene (also referred to hereinafter as Amp^r) and neomycin resistance gene (G418 resistance, also referred to hereinafter as Neo^r). In particular, if the dhfr gene is used as a selective marker for CHO (dhfr⁻) cells, a transformant carrying the desired gene can also be selected in a thymidine-free medium.

A signal sequence compatible with the host is added as

necessary to a nucleotide sequence for the N-terminal of the protein of the present invention. An alkali phosphatase signal sequence, Omp A signal sequence, etc. can be utilized for microorganisms of the genus Escherichia used as the host; an α -amylase signal sequence, subtilisin signal sequence, etc. for microorganisms of the genus Bacillus as the host; a mating factor α signal sequence, invertase signal sequence, etc. for yeasts as the host; and an insulin signal sequence, α -interferon signal sequence, antibody molecule signal sequence, etc. for animal cells as the host.

The thus constructed vector containing the DNA encoding the protein of the present invention can be used to produce transformants.

The microorganisms of the genus Escherichia, the microorganisms of the genus Bacillus, yeasts, insect cells, insects, animal cells, etc. are used as the host.

The microorganisms of the genus Escherichia used include, for example, Escherichia coli K12 DH1 (Proc. Natl. Acad. Sci. USA, 60, 160 (1968)), JM103 (Nucleic Acids Research, 9, 309 (1981)), JA221 (Journal of Molecular Biology, 120, 517 (1978)), HB101 (Journal of Molecular Biology, 41, 459 (1969)), C600 (Genetics, 39, 440 (1954)), etc.

The microorganisms of the genus Bacillus used include,

for example, Bacillus subtilis MI114 (Gene, 24, 255 (1983)),
207-221 (Journal of Biochemistry, 95, 87 (1984)), etc.

The yeasts used include, for example, Saccharomyces
cerevisiae AH22, AH22R⁻, NA87-11A, DKD-5D, 20B-12,
5 Schizosaccharomyces pombe NCYC1913, NCYC2036, Saccharomyces
picjia pastoris, etc.

The insect cells used when the virus is AcNPV include,
for example, cells (Spodoptera frugiperda cells; Sf cells)
from an established cell line derived from caterpillars of
10 Spodoptera frugiperda, MG1 cells derived from the midgut in
Trichoplusia ni, High FiveTM cells derived from eggs of
Trichoplusia ni, cells derived from Mamestra brassicae and
cells derived from Estigmena acrea. When the virus is
BmNPV, cells (Bombyx mori N cells; BmN cells) from an
15 established cell line derived from silkworms, etc., are
used. The Sf cells used include, for example, Sf9 cells
(ATCC CRL1711), Sf21 cells (Vaughn, J. L. et al., In Vivo,
13, 213-217 (1977)), etc.

The insects used include, for example, silkworm
20 caterpillars (Maeda et al., Nature, 315, 592 (1985)).

The animals cells used include, for example, simian
cell COS-7, Vero, Chinese hamster cell CHO, DHFR gene-
defect Chinese hamster cell CHO (dhfr⁻ CHO cells), mouse L
cells, mouse AtT-20, mouse myeloma cells, rat GH3, human FL
25 cells, etc.

The microorganisms of the genus Escherichia can be transformed according to a method described in e.g. Proc. Natl. Acad. Sci. USA, 69, 2110 (1972) or Gene, 17, 107 (1982).

5 The microorganisms of the genus Bacillus can be transformed according to a method described in e.g. Molecular & General Genetics, 168, 111 (1979), etc.

10 The yeasts can be transformed according to a method described in e.g. Methods in Enzymology, 194, 182-187 (1991), Proc. Natl. Acad. Sci. USA, 75, 1929 (1978), etc.

 The insect cells or insects can be transformed according to a method described in e.g. Bio/Technology, 6, 47-55 (1988), etc.

15 The animal cells can be transformed according to a method described in e.g. "Saibo Kogaku Bessatsu 8, Shin-Saibo Kogaku Jikken Protocol (Cell Technology, Extra Number 8, New Experimental Protocol in Cell Technology)", 263-267 (1995) (published by Shujunsha), Virology, 52, 456 (1973), etc.

20 Thereby, a transformant transformed with an expression vector comprising a DNA encoding the protein can be obtained.

25 When transformants derived from the microorganisms of the genus Escherichia or Bacillus as the host are cultured, the medium used for their culture is preferably a liquid

medium containing a carbon source, a nitrogen source, inorganic matter, etc. necessary for growth of the transformants. The carbon source includes, for example, glucose, dextrin, soluble starch, sucrose, etc.; the
5 nitrogen source includes, for example, inorganic or organic materials such as ammonium salts, nitrates, corn steep liquor, peptone, casein, meat extract, soybean cake and potato extract; and the inorganic matter includes, for example, calcium chloride, sodium dihydrogen phosphate,
10 magnesium chloride, etc. In addition, yeast extracts, vitamins, growth promoters, etc. may be added. The pH value of the medium is desirably about 5 to 8.

For example, the medium for culturing the microorganisms of the genus Escherichia is preferably M9
15 medium containing glucose and casamino acid (Miller, Journal of Experiments in Molecular Genetics, 431-433, Cold Spring Harbor Laboratory, New York, 1972). To permit the promoter to work efficiently, a chemical such as 3 β -indolylacrylic acid can be added as necessary.

20 The transformants from the microorganisms of the genus Escherichia as the host are cultured usually at about 15 to 43 °C for about 3 to 24 hours during which the medium may be aerated or stirred as necessary.

The transformants from the microorganisms of the genus
25 Bacillus as the host are cultured usually at about 30 to 40

°C for about 6 to 24 hours during which the medium may be aerated or stirred as necessary.

The medium used for culturing the transformants from yeasts as the host includes, for example, Burkholder
5 minimum medium (Bostian, K. L. et al., Proc. Natl. Acad. Sci. USA, 77, 4505 (1980)) and SD medium containing 0.5 % casamino acid (Bitter, G. A. et al., Proc. Natl. Acad. Sci. USA, 81, 5330 (1984)). The pH value of the medium is adjusted preferably to about 5 to 8. The transformants are
10 cultured usually at about 20 to 35 °C for about 24 to 72 hours during which the medium may be aerated or stirred as necessary.

The medium used for culturing the transformants from insect cells or insects as the host includes, for example,
15 a medium prepared by adding inactivated additives such as 10 % bovine serum as necessary to Grace's insect medium (Grace, T. C. C., Nature, 195, 788 (1962)). The pH value of the medium is adjusted preferably to about 6.2 to 6.4. The transformants are cultured usually at about 27 °C for
20 about 3 to 5 days during which the medium may be aerated or stirred as necessary.

The medium used for culturing the transformants from animal cells as the host includes, for example, MEM medium containing about 5 to 20 % fetal bovine serum (Science, 122,
25 501 (1952)), DMEM medium (Virology, 8, 396 (1959)), RPMI

1640 medium (Journal of the American Medical Association,
199, 519 (1967)), 199 medium (Proceeding of the Society for
the Biological Medicine, 73, 1 (1950)), etc. The pH value
is preferably about 6 to 8. The transformants are cultured
5 usually at about 30 to 40 °C for about 15 to 60 hours
during which the medium may be aerated or stirred as
necessary.

As described above, the protein of the present
invention can be formed in the culture medium or in the
10 transformants.

Separation and purification of the protein of the
present invention from the above culture can be performed
for example by the following method.

To extract the protein of the present invention from
15 the cultured microorganisms or cells, the cultured
microorganisms or cells are collected in a usual manner,
suspended in a suitable buffer, disrupted by sonication,
lysozyme and/or freezing and thawing, and centrifuged or
filtered to give a crude extract of the protein. The
20 buffer may contain protein denaturants such as urea and
guanidine hydrochloride and surfactants such as Triton X-
100TM. When the protein is secreted into the culture
liquid, the culture supernatant is collected by separating
the supernatant from the cultured microorganisms or cells
25 by a per se known method.

The culture supernatant thus obtained, or the protein contained in the extract, can be purified by a suitable combination of separation and purification techniques known per se. These known separation and purification techniques
5 make use of a method of utilizing solubility, such as salting-out and solvent precipitation, a method of mainly utilizing a difference in molecular weight, such as dialysis, ultrafiltration, gel filtration, and SDS-polyacrylamide gel electrophoresis, a method of utilizing a
10 difference in electric charge, such as ion-exchange chromatography, a method of utilizing specific affinity, such as affinity chromatography, a method of utilizing a difference in hydrophobicity, such as reverse-phase high performance liquid chromatography, a method of utilizing a
15 difference in isoelectric point, such as isoelectric focusing.

If the protein thus obtained is in a free form, it can be converted into a salt by a per se known method or its analogous method, while if the resulting protein is
20 obtained in the form of a salt, it can be converted into a free protein or another salt by a per se known method or its analogous method.

Before or after purification, the protein produced by the transformants can be arbitrarily modified or its
25 partial polypeptide can be removed by allowing a suitable

protein-modifying enzyme to act on the protein. For example, trypsin, chymotrypsin, arginyl endopeptidase, protein kinase or glycosidase is used as the protein-modifying enzyme.

5 The thus formed protein of the present invention or a salt thereof can be measured for its activity by an experiment of binding to a labeled ligand, enzyme immunoassays using specific antibody, etc.

10 The antibody against the protein of the present invention, a partial peptide thereof or a salt thereof may be a polyclonal or monoclonal antibody capable of recognizing the protein of the present invention, a partial peptide thereof or a salt thereof.

15 The antibody against the protein of the present invention, a partial peptide thereof or a salt thereof (referred to collectively as the protein of the present invention) can be produced by a known process for producing antibody or antiserum by using the protein of the present invention as the antigen.

20 [Preparation of a monoclonal antibody]

(a) Preparation of monoclonal antibody-producing cells

25 The protein of the present invention is administered alone or together with a carrier and a diluent into mammals at a site where the antibody can be produced by administration. To enhance the ability of the animals upon

administration to produce the antibody, complete Freund's adjuvant or incomplete Freund's adjuvant may be administered. Administration is conducted usually once every 2 to 6 weeks and about 2 to 10 times in total. The mammals used include, for example, monkey, rabbit, dog, guinea pig, mouse, rat, sheep, goat and chicken, among which mouse and rat are preferably used.

For production of the monoclonal antibody-producing cells, those animals having antibody titer are selected from the warm-blooded animals (e.g. mice) immunized with the antigen, and on the second to fifth day after the final immunization, their spleens or lymph nodes are collected, and the antibody-producing cells contained therein are fused with myeloma cells, whereby monoclonal antibody-producing hybridoma can be produced. The antibody titer in antiserum can be measured for example by reacting the antiserum with the labeled protein described below and then measuring the activity of the labeling agent bound to the antibody. Fusion can be carried out by a known method such as the method of Keller and Millstein (Nature, 256, 495 (1975)). The fusion promoter includes polyethylene glycol (PEG) and Sendai virus, and PEG is preferably used.

The myeloma cells include, for example, NS-1, P3U1, SP2/0 and AP-1 among which P3U1 is preferably used. The ratio of the antibody-producing cells (spleen cells) to the

myeloma cells used is from about 1 : 1 to about 20 : 1, and cell fusion can be effected efficiently by incubating the cells for about 1 to 10 minutes at 20 to 40 °C, preferably 30 to 37 °C, in the presence of PEG (preferably PEG 1000 to
5 PEG 6000) at a concentration of about 10 to 80 %.

The anti-protein antibody-producing hybridoma can be screened by various methods, for example by adding a culture supernatant of the hybridoma to a solid phase (e.g., a microplate) having the antibody to the protein, etc.,
10 adsorbed thereon directly or along with a carrier and then adding a radioactive substance- or enzyme-labeled anti-immunoglobulin antibody (which is e.g. an anti-mouse immunoglobulin antibody when mouse cells are subjected to cell fusion) or protein A to detect the monoclonal antibody
15 bound to the solid phase or by adding a culture supernatant of the hybridoma to a solid phase having an anti-immunoglobulin antibody or protein A adsorbed thereon and then adding the protein labeled with a radioactive substance or with an enzyme to detect the anti-protein
20 monoclonal antibody bound to the solid phase.

The anti-protein monoclonal antibody can be screened according to a per se known method or its analogous method. Screening can be carried out usually in an animal cell culture medium to which HAT (hypoxanthine, aminopterin,
25 thymidine) was added. The screening and breeding medium

may be any medium in which the hybridoma can grow. Examples of such medium include PRMI 1640 medium containing 1 to 20 % (preferably 10 to 20 %) FBS, GIT medium containing 1 to 10 % FBS (Wako Pure Chemical Industries, Ltd.), and a serum-free medium for hybridoma culture (SFM-101, Nissui Pharmaceutical, Co., Ltd.). The culture temperature is usually 20 to 40 °C, preferably about 37 °C. The culture time is usually 5 days to 3 weeks, preferably 1 to 2 weeks. Culture can be conducted usually in 5 % CO₂ gas. The antibody titer in a culture supernatant of the hybridoma can be measured in the same manner as in the measurement of the anti-protein antibody titer in antiserum as described above.

(b) Purification of the monoclonal antibody

Separation and purification of the anti-protein monoclonal antibody can be performed according to the per se known method, for example, a method of separating and purifying an immunoglobulin, such as a salting-out method, an alcohol precipitation method, an isoelectric precipitation method, an electrophoresis method, an absorbing-desorbing method with an ion-exchanger (e.g. DEAE), an ultracentrifugation method, a gel filtration method], a specific purifying method for obtaining an antibody by collecting it exclusively on antigen-bound solid phase or on an active adsorbent such as protein A or

protein G and then dissociating a bonding thereof to obtain the antibody.

[Preparation of a polyclonal antibody]

The polyclonal antibody of the present invention can be produced by the per se known method or its analogous method. For example, the polyclonal antibody can be produced by making an immune antigen (protein antigen)-carrier protein conjugate, then immunizing a mammal therewith in the same manner as in the method of producing the above monoclonal antibody, collecting a material comprising an antibody to the protein of the present invention from the immunized animal, and separating and purifying the antibody.

For the immune antigen-carrier protein conjugate used for immunizing mammals, the type of the carrier protein and the mixing ratio of the carrier to the hapten are not particularly limited insofar as the desired antibody can be efficiently produced by immunization with the antigen crosslinked via the hapten with the carrier, and for example, the conjugate is produced by coupling the hapten with bovine serum albumin, bovine cytoglobulin, hemocyanin or the like in a ratio of from 1 to about 0.1 to 20, preferably from 1 to about 1 to 5.

The hapten can be coupled with the carrier by use of various condensation agents such as glutaraldehyde,

carbodiimide, maleimide-activated ester, and activated ester reagents containing thiol group and dithiopyridyl group.

5 The resulting condensation product is administered alone or together with a carrier and a diluent into warm-blooded animals at a site where the antibody can be produced. To enhance the ability of the animals upon administration to produce the antibody, complete Freund's adjuvant or incomplete Freund's adjuvant may be
10 administered. Administration is conducted usually once every 2 to 6 weeks and about 3 to 10 times in total.

The polyclonal antibody can be collected from blood, ascites, etc. preferably blood in the warm-blooded animals immunized by the method described above.

15 The polyclonal antibody titer in antiserum can be measured in the same manner as in the measurement of the antibody titer in antiserum as described above. Separation and purification of the polyclonal antibody can be carried out by a method of separating and purifying immunoglobulin,
20 which is similar to the separation and purification of the monoclonal antibody as described above.

As the antisense DNA having a nucleotide sequence substantially complementary to a DNA or mRNA encoding the protein of the present invention or a partial peptide
25 thereof, any antisense DNAs may be used as long as they are

oligonucleotides or derivatives thereof comprising a nucleotide sequence substantially complementary to the whole or a part of the nucleotide sequence of a DNA or mRNA encoding the protein of the present invention or a partial peptide thereof and having an ability to suppress expression of said protein or a partial peptide thereof.

As the nucleotide sequence substantially complementary to the DNA or mRNA, for example, a nucleotide sequence having about 70 % or more, preferably about 80 % or more, more preferably about 90 % or more, and most preferably about 95 % or more homology with the whole or a part of a nucleotide sequence (that is, a complementary chain of said DNA or mRNA) complementary to said DNA or mRNA, etc., are exemplified. Particularly, antisense DNAs having about 70 % or more, preferably about 80 % or more, more preferably about 90 % or more, and further preferably about 95 % or more homology with the partial nucleotide sequence which codes, in the entire nucleotide sequence of the complementary chain of the DNA or mRNA of the present invention, for an N-terminal region of the protein of the present invention (e.g., a nucleotide sequence in the vicinity of the initiation codon) are preferable. These antisense DNAs can be produced using e.g. a known DNA synthesizer.

The protein of the present invention, a partial

peptide thereof or a salt thereof (also referred to hereinafter as the protein of the present invention), the DNA encoding the protein of the present invention or a partial peptide thereof (also referred to hereinafter as the DNA of the present invention), the antibody against the protein of the present invention (also referred to hereinafter as the antibody of the present invention) and the antisense DNA can be used in (1) a method of determining a binding protein to the protein of the present invention, (2) construction of a system for expressing a recombinant protein, (3) development of an assay system using the two-hybrid method and screening of candidates for pharmaceutical compounds, (4) practice of a drug design based on comparison with a structurally analogous ligand receptor and as (5) a reagent in formation of probes, PCR primers in genetic diagnosis etc., and further can be used as (6) a chemical for gene therapy etc.

In particular, by acquisition of a binding protein to the protein of the present invention by using the two-hybrid method and further by a binding assay system using the protein of the present invention and an activin receptor, acquired other receptors or an activin intracellular information transmission molecule, a promoter or an inhibitor in an information transmission system specific to warm-blooded animals such as human beings can

be screened, and the promoter or the inhibitor can be used as an agent for preventing or treating various diseases, as described in detail below.

(1) A method of determining a binding protein to the
5 protein of the present invention

The protein of the present invention is useful as a reagent for screening or determining a binding protein to the protein of the present invention.

That is, the present invention provides a method of
10 determining a binding protein to the protein of the present invention, which comprises screening a binding protein interacting with the protein of the present invention.

Specifically, the method of determining a binding protein according to the present invention is a method of
15 determining a protein or a salt thereof interacting with the protein of the present invention, which comprises introducing (1) a vector having the protein of the present invention fused with a DNA binding region of a transcriptional factor on a host cell expression vector and
20 (2) a fusion library of a test protein with a transcription-activating region, into a host cell having a reporter gene maintaining said binding region of the transcriptional factor on a promoter, and measuring a change in the amount of the expressed reporter gene which
25 is increased by binding of the two kinds of proteins.

The method of determining a binding protein according to the present invention comprises use of the two-hybrid method of detecting the binding protein by converting the interaction of the protein of the present invention with a test protein into the expression of a specific reporter gene.

The method of determining a binding protein according to the present invention is described below in more detail.

As the DNA encoding the protein of the present invention used in the method of determining a binding protein, any DNAs can be used insofar as they comprise a nucleotide sequence encoding a protein containing an amino acid sequence substantially identical to an amino acid sequence represented by SEQ ID NO: 5 or NO: 6 in the present invention or a nucleotide sequence encoding a partial peptide thereof. Further, the DNA may be a genomic DNA from warm-blooded animals (e.g., humans), a genomic DNA library from warm blooded animals (e.g., humans), a cDNA derived from tissues and cells in warm blooded animals (e.g., humans), a DNA isolated by a per se known method from a cDNA library derived from tissues and cells in warm-blooded animals (e.g., humans), a synthetic or semi-synthetic DNA. The vector used in the library may be any of bacteriophage, plasmid, cosmid and phagemide. Alternatively, the DNA can be isolated directly by the RT-

PCR method using an mRNA fraction prepared from tissues and cells, or can also be produced by chemically producing partial nucleotide sequences respectively and ligating them.

As the test protein library screened, commercial cDNA
5 libraries (e.g. MATCHMAKER cDNA produced by Clontech)
derived from various animals, etc. are used.

As the vector expressing a fusion protein of the DNA-binding region with a test protein, pAS2-1, pGBT9, pKAD-09, pSD09, pEG202, pBTM116, etc. are used.

10 As the vector expressing a fusion protein of an
activating region of a transcriptional factor with the
present protein, pGAD424, pACT2, pKT10Gal-VP, pJG4-5, pVP16,
etc. are used.

As the transcriptional factor, GAL4, LexA, SRF, etc.
15 are used.

As the reporter gene, b-galactosidase gene (LacZ),
histidine gene (HIS3), luciferase gene, chloramphenicol
acetyl transferase gene, etc. are used.

As the host cells, yeast (Saccharomyces cerevisiae),
20 Escherichia coli, etc. are used, among which CG-1945, Y190,
Y187, HF7c, SFY526, L40, EGY48, HIS/L1, 62L yeast strains,
etc. are used.

Specifically, in the method of determining a binding
protein to the present protein, first a cDNA library
25 expressed in the form of a fusion protein by ligating a

plasmid bound to the same frame as that of a DNA fragment containing a nucleotide sequence encoding the present protein and a DNA fragment containing a nucleotide sequence encoding a DNA-binding region on a plasmid (e.g., pAS2-1),
5 to a DNA fragment containing a nucleotide sequence encoding a transcription-activating region of a transcriptional factor (e.g., GAL4) on a plasmid (e.g., pACT2) can be formed according to a method described in, for example, Molecular Cloning, 2nd ed. (J. Sambrook et al., Cold Spring
10 Harbor Lab. Press, 1989).

To introduce the two plasmids described above into host cells (e.g. Saccharomyces cerevisiae Y190), both the plasmids may be transformed simultaneously into the cells, or one plasmid may be first introduced followed by
15 introducing the other plasmid, and for example, transformation can be carried out in a method described in, for example, Methods in Enzymology, 194, 182-187 (1991) or Proc. Natl. Acad. Sci. USA, 75, 1929 (1978).

Expression of a reporter gene in the transformant can
20 be detected by conversion into the amount, emission, etc. of a coloring or fluorescent material formed by the reporter enzyme. More specifically, when the host cells are yeasts, expression of the reporter gene (e.g., β -galactosidase activity) can be detected by blue/white-
25 coloring screening by use of the replica plate method or

the filter method, preferably the filter method.

The colonies colored (e.g. blue) by the filter method are treated according to a method described in, for example, A Practical Approach (Bartel, P. L. et al., Oxford University Press, Oxford; 153-179, 1993a), whereby the
5 positive clone can be separated as a single colony.

Recovery of the plasmid DNA from the single transformant thus separated is performed according to a method described in, for example, Gene, 57, 267 (1987), Bio
10 Techniques, 14, 552 (1993), etc., and by determining the nucleotide sequence of said DNA, a receptor for the protein of the present invention can be determined.

Further, the binding protein to the protein of the present invention can also be determined by using a
15 commercial kit, for example MATCHMAKER GAL4 Two-Hybrid Systems (Clontech) according to the manufacture's instructions.

(2) Agent for preventing or treating deficiency of the protein of the present invention

20 If the binding protein to the protein of the present invention is revealed in the method described in (1) above, the expression site of said binding protein and the action possessed by the protein of the present invention via said binding protein, etc., could be revealed. On the basis of
25 these findings, the DNA encoding the protein of the present

invention can be used as an agent for preventing or treating diseases occurring via said binding protein. Further, the protein of the present invention exhibits the activity of binding to activin receptors or activin intracellular information transmission molecules, so it can be used as an agent for preventing or treating diseases occurring via the activin receptors or activin intracellular information transmission molecules.

For example, the gene for the protein of the present invention is expressed specifically in the brain, and therefore, when there are those patients with nerve cell abnormalities or cerebral diseases correlated with said binding protein, activin receptors or activin intracellular information transmission molecules, the action of the protein of the present invention in cerebral cells in said patients can be sufficiently demonstrated by (A) administering the DNA encoding the protein of the present invention into the patients, or (B) inserting the DNA encoding the protein of the present invention into cerebral cells etc. to express the present protein, followed by transplanting said cells in the patients. Accordingly, the DNA encoding the protein of the present invention can be used as a safe and low-toxic preventive or therapeutic agent for diseases occurring via the binding protein to the protein of the present invention or via activin receptors

or activin intracellular information transmission molecules

To use the DNA of the present invention as the above therapeutic agent, said DNA can be administered in a usual manner into a human being or a mammal directly or after

5 insertion thereof into a suitable vector such as retrovirus vector, adenovirus vector and adenovirus-associated virus vector. For example, the DNA of the present invention can

be used orally as tablets coated with sugar as necessary, as capsules, elixirs and microcapsules, or parenterally as

10 an aseptic solution with water or with other pharmaceutically acceptable solutions or as an injection such as suspension. For example, the DNA of the present

invention can be formed by blending with physiologically acceptable carriers, flavors, excipients, vehicles,

15 preservatives, stabilizers and binders, into a unit dosage form required for the generally recognized preparation implementation. An amount of an active ingredient in these

pharmaceutical preparations is such that a suitable volume in an indicated range can be obtained.

20 The additives which can be admixed with the tablets, capsules, etc. include, for example, binders such as gelatin, corn starch, tragacanth, gum arabic, excipients

such as crystalline cellulose, swelling agents such as corn starch, gelatin and alginic acid, lubricants such as

25 magnesium stearate, sweeteners such as sucrose, lactose and

saccharine, and flavors such as peppermint, akamono oil and cherry. When one capsule is in a unit form, liquid carriers such as fats and oils can be contained in the materials described above. The aseptic composition for injection can be formulated according to conventional pharmaceutical manufacturing by dissolving or suspending the active material and naturally occurring vegetable oils such as sesame oil and coconut oil in vehicles such as injection water. The aqueous solution for injection includes physiological saline or an isotonic solution containing glucose and other supplementary agents (e.g., D-sorbitol, D-mannitol, sodium chloride, etc.), and may be used in combination with suitable solubilizers such as alcohols (e.g., ethanol etc.), polyalcohols (e.g., propylene glycol, polyethylene glycol, etc.) and nonionic surfactants (e.g., Polysorbate 80TM, HCO-50, etc.). The oily solution includes sesame oil, soybean oil, etc., and may be used in combination with solubilizer such as benzyl benzoate, benzyl alcohol, etc.

Further, the above preventing or treating agent may be blended, for example, with the buffer (e.g., phosphate buffer, sodium acetate buffer, etc.), soothing agents (e.g., benzalkonium chloride, procaine hydrochloride, etc.), stabilizers (e.g., human serum albumin, polyethylene glycol, etc.), preservatives (e.g., benzyl alcohol, phenol, etc.),

antioxidants, etc. Usually, the prepared injection is introduced into suitable ampoules. Since the thus obtained preparation is safe and low toxic, it can be administered, for example, to a mammal (e.g., rat, rabbit, sheep, pig, cow, cat, dog, monkey, human being, etc.). A dose of the DNA is different depending on symptom etc., and in the case of oral administration, the dose is usually about 0.1 to 100 mg, preferably about 1.0 to 50 mg and more preferably about 1.0 to 20 mg per day in an adult (60 kg). When parenterally administered, one time dose is different depending upon an administration subject, a subject organ, symptom and an administration method and, for example, in the form of an injection, it is advantageous to administer about 0.01 to 30 mg, preferably about 0.1 to 20 mg and more preferably about 0.1 to 10 mg per day through intravenous injection in an adult (60 kg). In the case of other animals, an amount calculated per 60 kg can be administered.

(3) A method of screening a compound inhibiting or promoting a binding of the protein of the present invention to its binding protein

By use of the protein of the present invention or a salt thereof, or by use of a protein competitive binding assay system using an expression system constructed for a recombinant binding protein, a compound (e.g. peptides, proteins, non-peptide compounds, synthetic compounds,

fermentation products, etc.) or a salt thereof inhibiting or promoting a binding of the protein of the present invention to its binding protein can be screened. Further, by use of an expression system (two-hybrid method) for a reporter gene by interaction of the two kinds of proteins in a transformant into which a DNA encoding the protein of the present invention was introduced, a compound (e.g. peptides, proteins, non-peptide compounds, synthetic compounds, fermentation products, etc.) or a salt thereof for inhibiting or promoting a binding of the protein of the present invention to its binding protein can also be screened.

Such compounds include a compound (so-called an agonist) having a cell-stimulating activity (e.g., the activity of promoting or inhibiting growth promotion and phosphorylation of intracellular proteins) via the binding protein and a compound (so-called an antagonist) not having said cell-stimulating activity.

As the protein of the present invention, the protein determined in (1) above, activin receptors and the aforementioned activin intracellular information transmission molecules (particularly, Smad such as Smad3), etc., are used. Hereinafter, these are also referred to as a binding protein to the protein of the present invention.

That is, the present invention provides:

A method of screening a compound or a salt thereof inhibiting or promoting a binding of the protein of the present invention to a binding protein to the protein of the present invention, which comprises comparing:

- 5 1) (i) the case where the protein of the present invention or a salt thereof is contacted with a binding protein to the protein of the present invention with (ii) the case where the protein of the present invention or a salt thereof and a test compound are contacted with a binding
10 protein to the protein of the present invention, or
2) (i) a transformant into which a DNA encoding the protein of the present invention and a binding protein to the protein of the present invention with (ii) said transformant brought into contact with a test compound.

15 The screening method of the present invention comprises comparing:

- 1) (i) the case where the protein of the present invention or a salt thereof is contacted with a binding protein to the protein of the present invention with (ii) the case
20 where the protein of the present invention or a salt thereof and a test compound are contacted with a binding protein to the protein of the present invention, for example by measuring the amount of the protein of the present invention or a salt thereof bound to a binding
25 protein to the protein of the present invention, the cell-

stimulating activity thereof, etc., or

2) (i) a transformant into which a DNA encoding the protein
of the present invention and a binding protein to the
protein of the present invention with (ii) said
5 transformant brought into contact with a test compound, for
example by measuring the degree of coloration indicative of
the degree of expression of a reporter gene.

More specifically, the present invention provides:

(1) a method of screening a compound or a salt thereof
10 inhibiting or promoting a binding of the protein of the
present invention or a salt thereof to a binding protein to
the protein of the present invention, which comprises
comparing the case where the labeled protein of the present
invention or a salt thereof is contacted with a binding
15 protein to the present invention with the case where the
labeled protein of the present invention or a salt thereof
and a test compound are contacted with a binding protein to
the protein of the present invention, by measuring the
amount of the labeled protein or a salt thereof bound to
20 the binding protein in both the cases,

(2) a method of screening a compound or a salt thereof
inhibiting or promoting a binding of the protein of the
present invention or a salt thereof to a binding protein to
the protein of the present invention in the case where the
25 binding protein to the present invention is a membrane-

bound protein (e.g., membrane penetration receptor, channel, etc.), which comprises comparing the case where the labeled protein of the present invention or a salt thereof is contacted with cells containing the binding protein to the protein of the present invention or with a membrane fraction of said cells, with the case where the labeled protein of the present invention or a salt thereof and a test compound are contacted with cells containing the binding protein to the protein of the present invention or with a membrane fraction of said cells, by measuring the amount of the labeled protein of the present invention or a salt thereof bound to said cells or said membrane fraction in both the cases,

(3) a method of screening a compound or a salt thereof inhibiting or promoting a binding of the protein of the present invention or a salt thereof to a binding protein to the protein of the present invention, which comprises comparing the case where the labeled protein of the present invention or a salt thereof is contacted with the binding protein expressed in cells by culturing a transformant containing a DNA encoding the binding protein to the protein of the present invention, with the case where the labeled protein of the present invention or a salt thereof and a test compound are contacted with the binding protein expressed in cells by culturing a transformant containing a

DNA encoding the binding protein to the protein of the present invention, by measuring the amount of the labeled protein of the present invention or a salt thereof bound to the binding protein in both the cases, and

5 (4) a method of screening a compound or a salt thereof inhibiting or promoting a binding of the protein of the present invention or a salt thereof to a binding protein to the protein of the present invention, which comprises comparing host cells into which a plasmid expressing a gene
10 for the protein of the present invention and a plasmid expressing a gene for a binding protein to the protein of the present invention were introduced, with said host cells brought into contact with a test compound, by measuring the expression of a gene group whose expression is regulated by
15 interaction between the protein of the present invention and the binding protein to the protein of the present invention, or physiological reaction (e.g., secretion of FSH, etc., when said binding protein is an activin receptor) in both the host cells.

20 Hereinafter, the screening method of the present invention is specifically described.

 First, as the binding protein used in the screening method of the present invention, any proteins may be used as long as they contain said binding protein or a salt
25 thereof, but extracts from organs in mammals are preferable.

However, since organs derived particularly from human beings are hardly obtainable, the binding protein or a salt thereof expressed in a large amount by genetic recombination technology is suitable for screening.

5 In order to prepare the binding protein, the aforementioned method is used, but preparation can be performed by expressing a DNA encoding said protein in a mammal cell or an insect cell. As a DNA fragment encoding a protein part of interest, a complementary DNA is usually
10 used but is not necessarily limited to it. For example, a gene fragment or a synthetic DNA may be used. In order to introduce a DNA fragment encoding the binding protein into a host animal cell and express it effectively, it is preferable that the DNA fragment is integrated in
15 downstream of polyhedron promoter from nuclear polyhedrosis virus (NPV) belonging to Baculovirus whose host is an insect, SV40-derived promoter, promoter of retrovirus, metallothionein promoter, human heat shock promoter, cytomegalovirus promoter and SR α promoter. Investigation
20 of an amount and quality of the expressed receptor can be conducted by the per se known method. The investigation can be conducted by, for example, a method described in a reference [Nambi, P. et al., J. Biol. Chem., vol. 267, pp. 19555-19559, 1992].

25 Therefore, in the screening method of the present

invention, a material comprising the binding protein or a salt thereof or a salt thereof may be the binding protein or a salt thereof purified according to the per se known method, or a cell comprising said protein may be used, or
5 when said protein is a membrane-bound protein, a cell membrane fraction containing the same may be used.

In the screening method of the present invention, when a cell comprising the binding protein is used, the cell may be fixed with glutaraldehyde or formalin. The fixing
10 method can be conducted according to the per se known method.

A cell comprising the binding protein refers to a host cell which expressed the binding protein and, as the host cell, Escherichia coli, Bacillus subtilis, yeast, insect
15 cell and animal cell are exemplified.

A cell membrane fraction refers to a fraction in which a cell membrane obtained by the per se known method after disruption of a cell is contained in a large amount. As a method for disrupting a cell, there are a method of
20 squeezing a cell with a Potter-Elvehjem type homogenizer, disruption with a Waring blender or Polytron (manufactured by Kinematica), disruption by sonication, and disruption by jetting a cell through a thin nozzle while pressurizing with a French press. For fractionating a cell membrane, a
25 fractionating method by centrifugation force such as a

fractionating centrifugation method and density gradient centrifugation method is mainly used. For example, a disrupted cell solution is centrifuged at a low speed (500 to 3000 rpm) in a short time (usually about 1 minute to 10 minutes), the supernatant is centrifuged at a higher speed (15000 to 30000 rpm) usually for 30 minutes to 2 hours, and the resulting precipitate is used as a membrane fraction. An expressed binding protein and membrane components such as cell-derived phospholipid and membrane protein are contained in a large amount in the membrane fraction.

An amount of a binding protein in a cell comprising the binding protein or its membrane fraction is preferably 10^2 to 10^8 molecules, suitably 10^5 to 10^7 molecules per cell. In addition, as an expressed amount is larger, the activity of binding to the protein of the present invention per a cellular extract or a membrane fraction (specific activity) is raised, and not only high sensitive screening system can be constructed but also a large amount of samples can be measured at the same lot.

In order to conduct the above methods (1) to (3) for screening a compound inhibiting binding of the protein of the present invention to a binding protein to the protein of the present invention, a suitable binding protein fraction and the labeled protein of the present invention are necessary. As the binding protein fraction, a natural

binding protein fraction or a recombinant binding protein fraction having the equal activity thereto are desirable. Here, equal activity denotes an equal activity of binding to the protein of the present invention, etc.

5 As the labeled protein of the present invention, the labeled protein of the present invention, a labeled compound analogous to the protein of the present invention, etc. are used. For example, the protein of the present invention labeled with [³H], [¹²⁵I], [¹⁴C], [¹³⁵S] or the like
10 can be utilized.

Specifically, in order to conduct screening of a compound inhibiting binding of the protein of the present invention to a binding protein to the protein of the present invention, first, an authentic binding protein is
15 prepared by suspending a cell comprising the binding protein or a membrane fraction of the cell in a buffer suitable for screening. Any buffers which do not inhibit binding of the protein of the present invention to a binding protein to the protein of the present invention,
20 for example phosphate buffer and Tris-HCl buffer, pH 4 to 10 (desirably pH 6 to 8, may be used. In addition, in order to decrease non-specific binding, surfactants such as CHAPS, Tween-80TM (Kao-Atlas Company), digitonin, and deoxycholate may be added to the buffer. Further, in order
25 to suppress degradation of the binding protein and the

protein of the present invention by a protease, a protease inhibitor such as PMSF, leupeptin, E-64 (manufactured by Peptide Laboratory) and pepstain may be added. The protein of the present invention labeled with a predetermined amount of (5000 to 500000 cpm) of a radioactive substance is added to 0.01 ml to 10 ml of the binding protein solution, and simultaneously 10^{-4} to 10^{-10} M test compound is allowed to be coexist. In order to know an amount of non-specific binding (NSB), a reaction tube to which the unlabeled protein of the present invention has been added in large excess is also prepared. The reaction is conducted at about 0 to 50 °C, preferably about 4 to 37 °C, for about 20 minutes to 24 hours, desirably about 30 minutes to 3 hours. After the reaction, the reaction solution is filtered with a glass fiber filtering paper and washed with a suitable amount of the same buffer, and the radioactivity remaining in the glass fiber filtering paper is measured by a scintillation counter or a γ -counter. Assuming that the count (B0 - NSB) obtained by subtracting the amount of non-specific binding (NSB) from the count (B0) in the absence of the antagonist is 100 %, a test compound by which the amount of specific binding (B - NSB) becomes e.g. 50 % or less can be selected as a candidate for a substance having an antagonistic inhibitory ability, while a test compound by which the amount of specific

binding (B - NSB) becomes e.g. 150 % or more can be selected as a candidate for a compound having a binding-promoting ability.

To practice the method in (4) above for screening a
5 compound inhibiting or promoting a binding of the protein to a binding protein to the protein of the present invention, expression of a gene group whose expression is regulated by interaction between the protein of the present invention and a binding protein to the protein of the
10 present invention, or physiological reactions (e.g., secretion of FSH, etc. when the binding protein is an activin receptor) in host cells to which a plasmid expressing a gene for the protein of the present invention and a plasmid expressing a gene for a binding protein to
15 the protein of the present invention were introduced can be measured by a known method. Specifically, when the host cell is yeast, a transformant transformed by introducing, to a yeast cell, a plasmid expressing a fusion protein between the protein of the present invention and a DNA-
20 binding region of a transcription-regulating factor and a plasmid expressing a fusion protein between a binding protein to the protein of the present invention and an activating region of a transcription-regulating factor is created by the same method as described above. For
25 screening, this transformant is cultured at 30 °C for 2 to

4 days on an agar medium containing 10^{-4} to 10^{-10} M test compound. As the agar medium, a tryptophan/leucine/histidine-deficient SD medium, a tryptophan/leucine-deficient SD medium, etc. are used.

5 Thereafter, the filter method or the like is used in order to detect expression of the reporter gene as coloration of colonies by β -galactosidase activity. For this detection, a Whatman #5 filter or VWR grade 410 filter moistened with buffer Z/X-gal (Clontech) is placed on a filter to which
10 the transformant adheres, and incubated at 30 °C for 30 minutes to 8 hours, and then the coloration of the colonies on the filter is compared with the coloration of a control transformant not containing a test compound. A test compound added to the culture medium of colonies showing
15 darker coloration than that of the control can be selected as a candidate for a compound having an ability to promote binding, while a test compound added to the culture medium of colonies showing lighter coloration than that of the control can be selected as a candidate as a compound having
20 an ability to inhibit binding. Further, expression of the reporter gene can also be quantified as β -galactosidase activity by measuring the amount of o-nitrophenol formed by decomposition of the substrate. First, the transformant is cultured under shaking at 30 °C for 8 to 24 hours in a
25 liquid medium containing 10^{-4} to 10^{-10} M test compound. As

the liquid medium, a tryptophan/leucine/histidine
-deficient SD medium, a tryptophan/leucine-deficient SD
medium, etc. are used. Preferably, after overnight culture,
an aliquot of the culture liquid is inoculated into YPD
5 medium and cultured under shaking at 30 °C for 3 to 5 hours
during which the OD₆₀₀ of 0.5 is increased to 1.0. An
aliquot of this culture liquid is centrifuged, and the
resulting precipitate is added to a mixture of buffer Z/ β -
mercaptoethanol and then reacted by adding an o-nitrophenyl
10 galactoside solution (Sigma). The reaction is carried out
at 0 to 50 °C for 3 minutes to 24 hours, desirably at 30 °C
for 30 minutes to 15 hours, and the absorbance (OD₄₂₀) of
the thus yellowed supernatant at 420 nm is measured. From
the absorbance (OD₄₂₀) thus obtained, the β -galactosidase
15 activity is calculated as Miller unit by use of the
equation below.

In the case where upon adding a test compound to the
medium, the β -galactosidase activity is increased by about
10 % or more, preferably about 20 % or more, more
20 preferably about 30 % or more and most preferably about
50 % or more, said test compound can be selected as a
candidate for a compound having an ability to promote
binding between the protein of the present invention and a
binding protein to the present invention. On the other
25 hand, in the case where upon adding a test compound to the

medium, the β -galactosidase activity is decreased by about 10 % or more, preferably about 20 % or more, more preferably 30 % or more and most preferably about 50 % or more, said test compound can be selected as a candidate for
5 a compound having an ability to inhibit binding between the protein of the present invention and a binding protein to the present invention.

To perform screening by measuring the expression activity of the reporter gene, a DNA encoding the binding
10 protein to the protein of the present invention is necessary. As the DNA encoding the binding protein to the protein of the present invention, a DNA comprising said DNA, a DNA hybridizing therewith under high stringent conditions, etc. are desirable.

15 As the test compound, for example, peptides, proteins, non-peptide compounds, synthetic compounds, fermentation products, cellular extracts, plant extracts, animal tissue extracts, etc. are exemplified, and these compounds may be novel compounds or known compounds.

20 A kit for screening a compound or a salt thereof inhibiting or promoting a binding of the protein of the present invention to a binding protein to the protein of the present invention comprises the protein of the present invention or a salt thereof, a partial peptide thereof or a
25 salt thereof, cells containing the binding protein to the

protein of the present invention, a fraction extracted from cells containing the binding protein to the protein of the present invention, or a transformant into which a plasmid expressing a fusion protein between the protein of the present invention and a DNA-binding region of a transcription-regulating factor and a plasmid expressing a fusion protein between the binding protein to the protein of the present invention and an activating region of a transcription-regulating factor were introduced.

As the screening kit of the present invention, for example the following is exemplified.

1. Reagents for screening

(1) Transformant

Yeast Y190 strain transformed by introducing a plasmid expressing a fusion protein between the protein of the present invention and a DNA-binding region of a transcription-regulating factor and a plasmid expressing a fusion protein between an activin IIA-N receptor protein and an activating region of a transcription-regulating factor.

(2) Liquid culture medium

Tryptophan/leucine/histidine-deficient SD medium: An aqueous solution of a yeast nitrogen source (Difco) is sterilized in an autoclave, and then a drop-out solution consisting of L-isoleucine, L-valine, L-adenine hemisulfate,

L-arginine hydrochloride, L-lysine hydrochloride, L-methionine, L-phenyl alanine, L-threonine, L-tyrosine and L-uracil, which was sterilized in an autoclave and stored at 4 °C, is added thereto. Further, 40 % dextrose/stock
5 solution (Sigma) sterilized by filtration with a filter is added thereto at a concentration of 2 %, which may be stored at 4 °C or prepared when the medium is used.

YPD medium: Difco peptone and an aqueous solution of a yeast extract are sterilized in an autoclave, and 40 %
10 dextrose sterilized by filtration with a filter is added thereto at a final concentration of 2 %, which may be stored at 4 °C or prepared when the medium is used.

(3) Buffer

Z buffer: A buffer containing $\text{Na}_2\text{HPO}_4 \cdot 7\text{H}_2\text{O}$, $\text{NaH}_2\text{PO}_4 \cdot \text{H}_2\text{O}$,
15 KCl , $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ is adjusted to pH 7 or thereabout and sterilized in an autoclave, which may be stored at 4 °C or prepared when the buffer is used.

Buffer Z/ β -mercaptoethanol: A mixture of 100 parts of buffer Z and 0.27 part of β -mercaptoethanol.

20 (4) Substrate for β -mercaptoethanol

An o-nitrophenyl galactoside solution (Sigma) is dissolved in buffer Z and adjusted to a concentration of 4 mg/ml, and this substrate is prepared when it is used.

(5) Reaction termination solution

25 1 M sodium carbonate solution stored at 4 °C is used,

and this solution may be prepared when it is used.

2. Measurement method

(1) A yeast transformant is cultured under shaking at 30 °C overnight in 1 ml tryptophan/leucine-deficient SD medium
5 containing 5 μ l of 10^{-3} to 10^{-10} M test compound solution.

(2) 0.4 ml of the culture liquid is added to 1.6 ml YPD medium and cultured under shaking at 30 °C for 3 to 5 hours during which the OD₆₀₀ of 0.5 is increased to 1.0.

(3) 0.3 ml of the culture liquid is centrifuged at 14000
10 rpm for 30 seconds, and the resulting precipitates are suspended in 0.3 ml buffer Z and centrifuged again, and the resulting precipitates are suspended in 0.1 ml buffer Z.

(4) After the suspension is frozen once in liquid nitrogen, it is melted at 37 °C for 30 seconds to 1 minute. 0.7 ml
15 mixture of buffer Z/ β -mercaptoethanol and 0.16 ml o-nitrophenyl galactoside solution are added thereto, and the mixture is incubated at 30 °C until the solution turns yellow, and thereafter, 0.4 ml of 1 M sodium carbonate solution is added thereto.

(5) The mixture is centrifuged at 14000 rpm for 10 minutes, and the absorbance of its supernatant at 420 nm is measured, and the β -galactosidase activity is calculated as Miller unit by use of Equation 1 below.

[Equation 1]

25 β -Galactosidase activity = $1000 \times \text{OD}_{420} / (t \times V \times \text{OD}_{600})$

OD₄₂₀: Absorbance at 420 nm

t: Reaction time (min)

V: The volume of a suspension of the transformant in buffer Z, used in the reaction × the degree of dilution

5 A compound or a salt thereof obtained by use of the screening method or the screening kit according to the present invention is either a compound inhibiting binding between the present protein (particularly the protein which has 5 PDZ domains and/or 2 WW domains, is expressed
10 particularly in the brain, and binds to activin receptors and/or activin intracellular information transmission molecules) and a binding protein to the protein of the present invention, or a compound promoting said binding (hereinafter, a promoter).

15 The compound inhibiting binding of the protein of the present invention and a binding protein to the protein of the present invention includes (1) a compound or a salt thereof (so-called an agonist) which binds to a binding protein to the protein of the present invention thereby
20 inhibiting binding between the protein of the present invention and the binding protein to the protein of the present invention and which in itself has a cell-stimulating activity via the binding protein, (2) a compound or a salt thereof (so-called an antagonist) which
25 binds to a binding protein to the protein of the present

invention thereby inhibiting binding between the protein of the present invention and the binding protein to the protein of the present invention but which in itself does not have a cell-stimulating activity via the binding protein, and (3) a compound (referred to hereinafter as an inhibitor) or a salt thereof which inhibits binding between the protein of the present invention and a binding protein to the present invention without binding to the binding protein to the protein of the present invention.

Whether the binding between the compound or a salt thereof obtained by using the screening method or the screening kit according to the present invention and the binding protein to the protein of the present invention is present or not can be confirmed according to the above method of measuring the binding activity.

The cell-stimulating activity via the binding protein can be measured according to the per se known method or its analogous method.

As the compound obtained by using the screening method or the screening kit according to the present invention, peptides, proteins, non-peptide compounds, synthetic compounds, fermentation products, etc. are exemplified, and these compounds may be novel compounds or known compounds.

The agonist and the promoter have the same action as the physiological activity possessed by the protein of the

present invention or have the action of enhancing its physiological activity so that depending on the activity of said protein, they are useful in a safe and low-toxic pharmaceutical composition, particularly as an agent for preventing or treating nerve cell abnormalities or cerebral diseases (for example, Alzheimer's disease, Parkinson's disease, epilepsy, Huntington's chorea, etc.) correlated with said binding protein or activin receptors.

On the other hand, since the antagonist or the inhibitor can inhibit the physiological activity possessed by the protein of the present invention, it can be used in a safe and low-toxic pharmaceutical composition inhibiting the activity of said protein, particularly as an agent for preventing or treating nerve cell abnormalities or cerebral diseases (for example, Alzheimer's disease, Parkinson's disease, epilepsy, Huntington's chorea, etc.) correlated with said binding protein or activin receptors.

When the compound or a salt thereof obtained by using the screening method or the screening kit according to the present invention is used as the aforementioned pharmaceutical composition, a conventional means can be used. For example, said compound or a salt thereof can be used orally as tablets coated with sugar as necessary, as capsules, elixirs and microcapsules, or parenterally as an aseptic solution with water or with other pharmaceutically

acceptable solutions or as an injection such as suspension. For example, said compound or a salt thereof can be formed by blending with physiologically acceptable carriers, flavors, excipients, vehicles, preservatives, stabilizers and binders, into a unit dosage form required for the generally recognized preparation implementation. An amount of the active ingredient in these pharmaceutical preparations is such that a suitable volume in an indicated range can be obtained.

The additives which can be admixed with the tablets, capsules, etc. include, for example, binders such as gelatin, corn starch, tragacanth, gum arabic, excipients such as crystalline cellulose, swelling agents such as corn starch, gelatin and alginic acid, lubricants such as magnesium stearate, sweeteners such as sucrose, lactose and saccharine, and flavors such as peppermint, akamono oil and cherry. When one capsule is in a unit form, liquid carriers such as fats and oils can be contained in the materials described above. The aseptic composition for injection can be formulated according to conventional pharmaceutical manufacturing by dissolving or suspending the active material and naturally occurring vegetable oils such as sesame oil and coconut oil in vehicles such as injection water.

The aqueous solution for injection includes, for

example, physiological saline or an isotonic solution containing glucose and other supplementary agents (e.g., D-sorbitol, D-mannitol, sodium chloride, etc.), and may be used in combination with suitable solubilizers such as
5 alcohols (e.g., ethanol etc.), polyalcohols (e.g., propylene glycol, polyethylene glycol, etc.) and nonionic surfactants (e.g., Polysorbate 80TM, HCO-50, etc.).

The oily solution includes, for example, sesame oil, soybean oil, etc., and may be used in combination with
10 solubilizer such as benzyl benzoate, benzyl alcohol, etc. Further, the above preventing or treating agent may be blended, for example, with the buffer (e.g., phosphate buffer, sodium acetate buffer, etc.), soothing agents (e.g., benzalkonium chloride, procaine hydrochloride, etc.),
15 stabilizers (e.g., human serum albumin, polyethylene glycol, etc.), preservatives (e.g., benzyl alcohol, phenol, etc.), antioxidants, etc. Usually, the prepared injection is introduced into suitable ampoules.

Since the thus obtained preparation is safe and low
20 toxic, it can be administered, for example, to a mammal (e.g., rat, rabbit, sheep, pig, cow, cat, dog, monkey, human being, etc.).

A dose of said compound or a salt thereof is different depending on symptom etc., and in the case of oral
25 administration, the dose is usually about 0.1 to 100 mg,

preferably about 1.0 to 50 mg and more preferably about 1.0 to 20 mg per day in an adult (60 kg). When parenterally administered, one time dose is different depending upon an administration subject, a subject organ, symptom and an administration method and, for example, in the form of an injection, it is advantageous to administer about 0.01 to 30 mg, preferably about 0.1 to 20 mg and more preferably about 0.1 to 10 mg per day through intravenous injection in an adult (60 kg). In the case of animals other than human beings, an amount calculated per 60 kg can be administered.

(4) Quantification of the protein of the present invention or a partial peptide or a salt thereof

The antibody against the protein of the present invention can specifically recognize the protein etc. of the present invention so that it can be used for example in quantification of the protein etc. of the present invention in a test sample, particularly in quantification thereof by sandwich immunoassays.

That is, the present invention provides:

(i) a method of quantifying the protein etc. of the present invention in a test solution, which comprises allowing a test solution and the labeled protein etc. of the present invention to react competitively with an antibody reacting with the protein etc. of the present invention and determining the ratio of the labeled protein

etc. of the present invention bound to said antibody; and

(ii) a method of quantifying the protein etc. of the present invention in a test solution, which comprises allowing a test solution to react with the antibody of the present invention insolubilized on a carrier and the labeled antibody of the present invention simultaneously or successively and then measuring the activity of the labeling agent on the insolubilizing carrier, wherein one of the two antibodies is an antibody recognizing an N-terminal region or a C-terminal region of the protein etc. of the present invention, and the other antibody is an antibody reacting with an amino acid sequence of SEQ ID NO: 5 or NO: 6.

The monoclonal antibody recognizing the protein etc. of the present invention (hereinafter, also referred to as the anti-protein antibody) can be used not only for quantifying the protein etc. of the present invention but also for detection thereof by tissue staining etc. For these purposes, the antibody molecule itself may be used, or an $F(ab')_2$, Fab' or Fab fraction of the antibody molecule may be used.

The method of quantifying the protein etc. of the present invention by means of the antibody of the present invention is not particularly limited as long as the method comprises detecting the amount of the antibody, the antigen

or an antigen-antibody conjugate, corresponding to the amount of the antigen (e.g., the amount of the protein of the present invention) in a test solution, by chemical or physical means and calculating it on the basis of a
5 standard curve prepared using standard solutions containing known amounts of the antigen. For example, nephelometry, the competitive method, the immunometric method and the sandwich method are preferably used, but the sandwich method described later is particularly preferably used in
10 respect of sensitivity and specificity.

The labeling agent used in the measurement method using a labeling substance includes, for example, a radioisotope, enzyme, fluorescent material and luminescent material. The radioisotope includes, for example, [¹²⁵I],
15 [¹³¹I], [³H], [¹⁴C], etc.; the enzyme described above is preferably a stable enzyme with high specific activity, which includes, for example, β -galactosidase, β -glucosidase, alkali phosphatase, peroxidase and malate dehydrogenase; the fluorescent material includes, for example,
20 fluorescamine and fluorescein isocyanate; the luminescent material includes luminol, luminol derivatives, luciferin and lucigenin. Further, a biotin-avidin system can also be used for binding the labeling agent to the antibody or antigen.

25 The antigen or antibody may be insolubilized by

physical adsorption or via chemical bonding used conventionally for insolubilization or immobilization of proteins, enzymes, etc. The carrier includes insoluble polysaccharides such as agarose, dextran and cellulose, 5 synthetic resin such as polystyrene, polyacrylamide and silicon, and glass, etc.

In the sandwich method, a test sample is allowed to react with the insolubilized anti-protein antibody (primary reaction), then the labeled anti-protein antibody is 10 allowed to react therewith (secondary reaction), and the activity of the labeling agent on the insolubilizing carrier is measured, whereby the protein of the present invention in the test sample can be quantitatively determined. The primary and secondary reactions may be 15 carried out in the reverse order, simultaneously, or separately after a predetermined time. The labeling agent and the method of insolubilization may be in accordance with those described above. The antibody used as the solid-phase antibody or as the labeling antibody in 20 immunoassays by the sandwich method may not necessarily be one kind of antibody and may use a mixture of two or more kinds of antibodies for the purpose of improving measurement sensitivity etc.

In the method of measuring the protein etc. of the 25 present invention by the sandwich method according to the

present invention, the anti-protein antibodies used in the primary and secondary reactions are preferably those antibodies to which the protein etc. of the present invention are bound at different sites. That is, when the antibody used in, for example, the secondary reaction recognizes a C-terminal region of the protein etc. of the present invention, the other antibody used in the primary reaction recognizes preferably an amino acid sequence represented by SEQ ID NO: 5 or NO: 6.

The antibody against the protein etc. of the present invention can be used not only in the sandwich method but also in other measurement systems such as competitive method, immunometric method and nephelometry.

In the competitive method, an antigen in a test sample and its labeled antigen are allowed to react competitively with the antibody, then the unreacted labeled antigen (F) and the labeled antibody (B) bound to the antibody are separated from each other (B/F separation), and the amount of either labeled B or F is determined to quantify the amount of the antigen in the sample. This reaction method employs either a liquid-phase method in which a soluble antibody is used as the antibody and polyethylene glycol and a second antibody against the above antibody are used in B/F separation, or a solid-phase method in which a solid-phase (i.e. immobilized) antibody is used as the

first antibody, or a soluble antibody is used as the first antibody while a solid-phase antibody is used as the second antibody.

5 In the immunometric method, the antigen in a test solution and the solid-phase antigen are allowed to react competitively with a predetermined amount of the labeled antibody, followed by separating the solid phase from the liquid phase, or alternatively the antigen in a test solution is allowed to react with an excess of the labeled antibody, and then the solid-phase antigen is added thereto 10 to permit the unreacted labeled antibody to be bound to the solid phase, followed by separating the solid phase from the liquid phase. Then, the amount of the label in either phase is measured to quantify the amount of the antigen in 15 the test solution.

Further, in nephelometry, the amount of insoluble precipitates in gel or solution, formed by antigen-antibody reaction, is measured. Laser nephelometry using laser scattering can be applied preferably to the case where 20 precipitates are obtained in a small amount because of a very small amount of antigen in a test solution.

For application of these immunoassays to the quantification method of the present invention, it is not necessary to establish special conditions, procedures etc. 25 A measurement system for the protein etc. of the present

invention can be established in an ordinary manner by those skilled in the art by modifying the conventional conditions and procedures. The details of these general technical means can be referred to in general remarks, books, etc.

5 [for example, reference can be made of "Radioimmunoassays" edited by Hirhoshi Irie (published by Kodansha in 1974), "Radioimmunoassays" (2nd edition) edited by Hiroshi Irie (published by Kodansha in 1979), "Enzyme Immunoassays" edited by Eiji Ishikawa et al. (published by Igakushoin in 10 1978), "Enzyme Immunoassays" (2nd edition) edited by Eiji Ishikawa et al. (published by Igakushoin in 1982), "Enzyme Immunoassays" (3rd edition) edited by Eiji Ishikawa et al. (published by Igakushoin in 1987), Methods in Enzymology, Vol. 70 (Immunochemical Techniques (Part A), Methods in 15 Enzymology, Vol. 73 (Immunochemical Techniques (Part B)), Methods in Enzymology, Vol. 74 (Immunochemical Techniques (Part C)), Methods in Enzymology, Vol. 84 (Immunochemical Techniques (Selected Immunoassays (Part D)), and Methods in Enzymology, Vol. 92 (Immunochemical Techniques (Monoclonal 20 Antibodies and General Immunoassay Methods (Part E)), Methods in Enzymology, Vol. 121 (Immunochemical Techniques (Hybridoma Technology and Monoclonal Antibodies (Part I)) (which are published by Academic Press Ltd.).

By using the antibody of the present invention in the 25 manner as described above, the protein etc. of the present

invention can be quantified with good sensitivity.

Further, by quantifying the concentration of the protein etc. of the present invention by means of the antibody against the protein etc. of the present invention, for example, diagnosis of those diseases correlated with the protein etc. of the present invention can be performed.

Further, the antibody of the present invention can be used to detect the protein etc. of the present invention present in humor and tissues. In addition, the antibody of the present invention can be used for preparation of an antibody column used in purification of the protein etc. of the present invention, for detection of the protein etc. of the present invention in each fraction during purification, and for analysis of the behavior of the protein etc. of the present invention in cells examined.

(5) Genetic diagnostic agent

Since the DNA of the present invention can be used for example as a probe to detect abnormalities in a gene encoding the protein of the present invention or a partial peptide thereof in a human being or a mammal (e.g., rat, mouse, guinea pig, rabbit, bird, sheep, pig, cow, horse, cat, dog, monkey, chimpanzee, etc.), the DNA of the present invention is useful as a genetic diagnostic agent for e.g. mutation of said DNA or abnormal accumulation or abnormal reduction of the mRNA, etc.

This genetic diagnosis using the DNA of the present invention can be carried out by techniques known per se, for example Northern hybridization and the PCR-SSCP method (Genomics, Vol. 5, pp. 874-879 (1989) and Proceedings of
5 the National Academy of Sciences of the United States of America, Vol. 86, pp. 2766-2770 (1989)).

(6) DNA containing an antisense DNA

An antisense DNA capable of binding complementarily to the DNA or mRNA encoding the protein etc. of the present
10 invention to suppress transcription or translation of the DNA or mRNA can suppress abnormal expression of a gene encoding the protein etc. of the present invention. Therefore, the antisense DNA can be used as e.g. a therapeutic or preventive agent for diseases attributable
15 to abnormal expression of a gene encoding the protein etc. of the present invention.

When the antisense DNA is used as the above therapeutic or preventive agent, it can be produced in the same manner as for the aforementioned pharmaceutical
20 preparation containing the DNA of the present invention. For example, said antisense DNA can be administered into a human being or a mammal in a usual manner directly or after insertion thereof into a suitable vector such as retrovirus vector, adenovirus vector and adenovirus-associated virus
25 vector. The antisense DNA is formed alone or along with

physiologically acceptable carriers such as a supplementary agent for promoting ingestion, into a pharmaceutical preparation which can be administered via a gene gun or a catheter such as hydrogel catheter.

5 Further, said antisense DNA can also be used as a diagnostic oligonucleotide probe for examining the presence or expression of the DNA of the present invention in tissues or cells.

(7) Creation of DNA-transferred animals

10 The present invention provides non-human mammals having an extraneous DNA encoding the protein etc. of the present invention (referred to hereinafter as the extraneous DNA of the present invention) or a mutated DNA thereof (also referred to hereinafter as the extraneous
15 mutated DNA of the present invention).

That is, the present invention provides:

(i) non-human mammals having the extraneous DNA of the present invention or a mutated DNA thereof;

(ii) the non-human mammals according to item (i),
20 wherein the non-human mammals are rodents;

(iii) the non-human mammals according to item (ii),
wherein the rodents are mice;

(iv) the non-human mammals according to item (ii),
wherein the rodents are rats; and

25 (v) a recombinant vector comprising the extraneous DNA

of the present invention or a mutated DNA thereof and capable of expression in mammals.

The non-human mammals having the extraneous DNA of the present invention or its mutated DNA (hereinafter, referred to as the DNA-transferred animals of the present invention) can be produced by transferring the intended DNA to embryonic cells such as unfertilized cells, fertilized cells, sperms and their initial cells, preferably at the stage of embryonic development in the development of non-human mammals (more preferably at the stage of single cells or fertilized egg cells and generally before the 8-cell stage) by the calcium phosphate method, electric pulse method, lipofection method, aggregation method, microinjection method, particle gun method or DEAE-dextran method. By the method of transferring the DNA, the intended extraneous DNA of the present invention can be transferred to somatic cells, organs in a living body, tissues and cells, and utilized in cell culture and tissue culture, and these cells can also be fused with embryonic cells by the cell fusion per se known method in order to produce the DNA-transferred animals of the present invention.

The non-human mammals used include, for example, rat, mouse, guinea pig, hamster, rabbit, sheep, pig, cow, cat, dog, etc. In particular, rodents, particularly mice (for

example, C57BL/6 strain and DBA2 strain as pure strain and B6C3F₁ strain, BDF₁ strain, B6D2F₁ strain, BALB/c strain and ICR strain as crossed strain) and rats (for example, Wister, SD, etc.) are preferable in formation of a morbid animal
5 model system because of their easy breeding with a relatively short period of development and biological cycle.

With respect to the recombinant vector capable of expression in mammals, the "mammals" include human beings in addition to the non-human mammals described above.

10 The extraneous DNA of the present invention is not the DNA of the present invention inherent in non-human mammals but the DNA encoding the protein of the present invention, which was once isolated and extracted from mammals.

The mutated DNA of the present invention includes a
15 DNA derived from the original DNA of the present invention by mutating the nucleotide sequence thereof, specifically by adding or deleting some bases or by substituting some bases with other bases, and the mutated DNA also includes an abnormal DNA.

20 The abnormal DNA refers to a gene expressing the abnormal protein of the present invention, and for example, a DNA expressing a protein suppressing the functions of the normal protein of the present invention is used.

The extraneous DNA of the present invention may be
25 derived from animals of either the same species as or a

different species from that of the intended animals. For transferring the DNA of the present invention to the intended animals, the DNA is generally advantageously used as a DNA construct having the DNA bound to a region downstream from a promoter capable of expressing the DNA in animal cells. For example, in the case where the DNA of the present invention is to be transferred, a DNA construct (e.g. a vector) having the DNA of the present invention bound to a region downstream from a promoter permitting expression of a DNA having high homology with the DNA of the present invention and derived from various mammals (e.g., rabbit, dog, cat, guinea pig, hamster, rat, mouse, etc.) carrying the DNA of the present invention is microinjected into fertilized eggs of the intended animals, such as mouse fertilized eggs, whereby DNA-transferred animals highly expressing the DNA of the present invention can be produced.

The expression vector carrying said construct includes plasmids derived from E. coli, Bacillus subtilis or yeast, bacteriophage such as λ -phage, retrovirus such as MALONEY [phonetic transcription] leukemia virus, and animal viruses such as vaccinia virus and Baculovirus. In particular, the plasmids derived from E. coli or yeast are preferably used.

The promoter for regulating expression of the DNA includes, for example, promoters derived from viruses (e.g.,

simian virus, cytomegalovirus, MALONEY [phonetic] leukemia virus, JC virus, breast cancer virus, poliovirus, etc.), promoters derived from various mammals (human being, rabbit, dog, cat, guinea pig, hamster, rat, mouse, etc.), for
5 example, promoters for albumin, insulin II, UROPLAKIN (phonetic) II, elastase, erythropoietin, endoserine, muscular creatine kinase, glia fibrous acidic protein, glutathione S-transferase, platelet-derived growth factor β , keratin K1, K10 and K14, collagen I and II type, cyclic
10 AMP-dependent kinase β I subunit, dystrophin, tartaric acid-resistant alkali phosphatase, atrial natriuretic factor, endothelial receptor tyrosine kinase (abbreviated generally to Tie 2), sodium potassium adenosine triphosphatase (Na,K-ATPase), neurofilament light chain, metallothionein I and
15 IIA, metalloproteinase 1 tissue inhibitor, MHC class I antigen (H-2L), H-ras, renin, dopamine β -hydroxylase, thyroid peroxidase, polypeptide chain-extending factor 1α (EF- 1α), β -actin, α - and β -myosin heavy chains, myosin light chains 1 and 2, myelin basic protein, thyroglobulin,
20 Thy-1, immunoglobulin, H chain variable region (VNP), serum amyloid P component, myoglobin, troponin, smooth muscular α -actin, preproenkephalin A and vasopressin, among which promoters such as cytomegalovirus promoter, human polypeptide-extending factor 1α (EF- 1α) promoter, human
25 and chicken β actin promoters can be preferably used

because of their ability to express the DNA in the whole body.

Preferably, the vector has a sequence (generally called a terminator) for terminating transcription of the intended mRNA in the DNA-transferred mammals, and for
5 example the sequence of each DNA derived from viruses or mammals, preferably SV40 terminator from simian virus, can be used.

Further, a splicing signal for each DNA, an enhancer
10 region, a part of introns from eucaryotic DNA, etc. can also be linked depending on the object to a 5'-upstream region from the promoter, to a region between the promoter and the translating region, or to a 3'-downstream region from the translating region.

15 The translating region for the normal protein of the present invention can be obtained in the per se known method as the whole or a part of genomic DNA derived from the liver, kidney, thyroid cells, and fibroblasts derived from various mammals (e.g., rabbit, dog, cat, guinea pig,
20 hamster, rat, mouse, human being) and genomic DNA from a variety of commercial genomic DNA libraries or by use of a complementary DNA as the starting material prepared in the per se known method from RNA derived from the liver, kidney, thyroid cells, and fibroblasts. Further, the extraneous
25 abnormal DNA can be obtained from the above cells or

tissues with those diseases attributable to a mutation in the protein of the present invention. In addition, a mutated translating region can be produced by a point mutagenesis method of mutating the translating region for the normal protein obtained from the above cells or tissues.

The translating region can be constructed as a DNA construct capable of expression in the DNA-transferred animals by conventional DNA engineering means for linking it to a downstream region from the promoter or if desired to an upstream region from the transcription termination site.

The extraneous DNA of the present invention is transferred to cells at the stage of fertilized egg cells so as to secure the presence of the DNA in all embryonic cells and somatic cells of the intended mammals. The presence of the extraneous DNA of the present invention in the embryonic cells in the produced animals after transfer of the DNA means that the extraneous DNA of the present invention is maintained in all embryonic cells and somatic cells in the offspring of the produced animals. The offspring of the produced animals, which has inherited the extraneous DNA of the present invention, has the extraneous DNA of the present invention in all embryonic cells and somatic cells thereof.

After it is confirmed that the non-human mammals

having the extraneous normal DNA of the present invention transferred to them maintain the DNA stably through crossbreeding, they can be bred generation after generation as animals carrying said DNA under normal breeding environments.

The extraneous DNA of the present invention is transferred to cells at the stage of fertilized egg cells so as to secure the presence of the DNA in excess in all embryonic cells and somatic cells in the intended mammals.

The presence of the extraneous DNA of the present invention in excess in the embryonic cells in the produced animals after transfer of the DNA means that the extraneous DNA of the present invention is maintained in excess in all embryonic cells and somatic cells in the offspring of the produced animals. The offspring of the produced animals, which has inherited the extraneous DNA of the present invention, has the extraneous DNA of the present invention in excess in all embryonic cells and somatic cells thereof. Homozygote animals having the introduced DNA in both homogenous chromosomes are obtained, and these male and female animals are crossbred to produce the offspring having said DNA in excess and capable of breeding generation after generation.

The non-human mammals having the normal DNA of the present invention can be used as a morbid animal model

since the normal DNA of the present invention is highly expressed in these animals to promote the functions of endogenous normal DNA, resulting in the onset of functional exasperation by the protein of the present invention. For
5 example, the normal DNA-transferred animals of the present invention can be used to elucidate the mechanism of the morbid functional exasperation caused by the protein of the present invention or those diseases correlated with the protein of the present invention or to examine the method
10 of treating these diseases.

Further, the mammals having the extraneous normal DNA of the present invention transferred to them can also be used in a test for screening a therapeutic agent against those diseases correlated with the protein of the present
15 invention because these animals have the symptom of an increase in the protein of the present invention.

On the other hand, the non-human mammals having the extraneous abnormal DNA of the present invention transferred to them are confirmed to maintain the
20 extraneous DNA stably through crossbreeding, and they can be then bred generation after generation as animals carrying said DNA under normal breeding environments. Further, the desired extraneous DNA can be used as a starting material by integrating it in the plasmid
25 described above. The DNA construct having the promoter can

be produced in usual DNA engineering means. The abnormal DNA of the present invention is transferred to cells at the stage of fertilized egg cells so as to secure the presence of the DNA in all embryonic cells and somatic cells of the intended mammals. The presence of the abnormal DNA of the present invention in the embryonic cells in the produced animals after transfer of the DNA means that the abnormal DNA of the present invention is maintained in all embryonic cells and somatic cells in the offspring of the produced animals. The offspring of the produced animals, which has inherited the DNA, has the abnormal DNA of the present invention in all embryonic cells and somatic cells thereof. Homozygote animals having the introduced DNA in both homogenous chromosomes are obtained, and these male and female animals can be crossbred to produce the offspring having said DNA and capable of breeding generation after generation.

The non-human mammals having the abnormal DNA of the present invention can be used as a morbid animal model since the abnormal DNA of the present invention is highly expressed in these animals to inhibit the functions of endogenous normal DNA, resulting in the onset of functional inactivated refractory conditions [literal translation] of the protein of the present invention. For example, the abnormal DNA-transferred animals of the present invention

can be used to elucidate the mechanism of the morbid functional inactivated refractory conditions of the protein of the present invention or to examine the method of treating this disease.

5 For specific applicability, the abnormal DNA high-expression animals of the present invention serve as a model for elucidation of the functional inhibition (dominant negative action) of the normal protein by the abnormal protein of the present invention in the functional
10 inactivated refractory conditions of the protein of the present invention. Further, the mammals having the extraneous abnormal DNA of the present invention transferred to them can also be used in a test for screening a therapeutic agent against the functional
15 inactivated refractory conditions of the protein of the present invention since these animals have the symptom of an increase in the protein of the present invention.

Other applicability of the two DNA-transfected animals of the present invention includes:

- 20 1. Use thereof as a cellular source for tissue culture;
2. Analysis of the relationship with proteins specifically expressed or activated by the protein of the present invention by direct analysis of mRNA in tissues in the DNA-transferred animals of the present invention or by analysis
25 of tissues where the protein is expressed in the DNA-

transferred animals;

3. Screening of chemicals promoting or inhibiting the functions of cells by using the cells described according to item 1; and

5 4. Isolation and purification of the mutated protein of the present invention and preparation of an antibody against said protein.

Further, the DNA-transferred animals of the present invention can be used to examine the clinical symptoms of
10 those diseases correlated with the protein of the present invention, including the functional inactivated refractory conditions of the protein of the present invention, or to obtain further detailed pathological findings in each organ in a model with those diseases correlated with the protein
15 of the present invention, and to contribute to development of new therapeutic methods as well as to the study and treatment of secondary diseases resulting from said diseases.

Further, each organ is removed from the DNA-
20 transferred animals of the present invention, cut into thin pieces, and digested with proteinase such as trypsin, whereby the free DNA-transferred cells can be obtained, and these cells can be cultured and these cultured cells can be established as a cell line. By identifying those cells
25 producing the protein of the present invention and by

examining their relationship with differentiation or multiplication or the signal transfer mechanism therein, their abnormalities can be examined so that these cells can be used as an useful material for study on the protein of the present invention and in elucidating their action.

The DNA-transferred animals of the present invention can also be used to provide an effective and rapid screening method for therapeutic agents for those diseases correlated with the protein of the present invention by use of the examination method and quantification method described above, so that the therapeutic agents against these diseases including the functional inactive type refractory conditions of the protein of the present invention can be developed. Further, the DNA-transferred animals of the present invention or the extraneous DNA expression vector of the present invention can be used for examination and development of the gene therapeutic method to treat those diseases correlated with the protein of the present invention.

When bases or amino acids are expressed in abbreviations in the present specification and drawings, the following abbreviations in accordance with IUPAC-IUB Commission on Biochemical Nomenclature or based on customary abbreviations in this field are used. If amino acids can occur as optical isomers, L-isomers are referred

to unless otherwise specified.

DNA: deoxyribonucleic acid

cDNA: complementary deoxyribonucleic acid

A: adenine

5 T: thymine

G: guanine

C: cytosine

RNA: ribonucleic acid

mRNA: messenger ribonucleic acid

10 dATP: deoxyadenosine triphosphate

dTTP: deoxythymidine triphosphate

dGTP: deoxyguanosine triphosphate

dCTP: deoxycytidine triphosphate

ATP: adenosine triphosphate

15 EDTA: ethylene diamine tetraacetate

SDS: sodium dodecyl sulfate

EIA: enzyme immunoassay

Gly: glycine

Ala: alanine

20 Val: valine

Leu: leucine

Ile: isoleucine

Ser: serine

Thr: threonine

25 Cys: cysteine

Met: methionine

Glu: glutamic acid

Asp: aspartic acid

Lys: lysine

5 Arg: arginine

His: histidine

Phe: phenyl alanine

Tyr: tyrosine

Trp: tryptophan

10 Pro: proline

Asn: asparagine

Gln: glutamine

pGl: pyroglutamic acid

Me: methyl group

15 Et: ethyl group

Bu: butyl group

Ph: phenyl group

TC: thiazolidine-4 (R)-carboxamide group

20 The substituent groups, protecting groups and reagents
appearing frequently in the present specification are
expressed in the following symbols.

Tos: p-toluene sulfonyl

CHO: formyl

Bzl: benzyl

25 Cl₂-Bzl: 2,6-dichlorobenzyl

Bom: benzyloxymethyl

Z: benzyloxycarbonyl

Cl-Z: 2-chlorobenzyloxycarbonyl

Br-Z: 2-bromobenzyloxycarbonyl

5 Boc: t-butoxycarbonyl

DNP: dinitrophenol

Trt: trityl

Bom: t-butoxymethyl

Fmoc: N-9-fluorenyl methoxycarbonyl

10 HOBt: 1-hydroxybenzotriazole

HOObt: 3,4-dihydro-3-hydroxy-4-oxo-1,2,3-benzotriazine

HONB: 1-hydroxy-5-norbornane-2,3-dicarbodiimide

DCC: N,N'-dicyclohexyl carbodiimide

15 The sequence numbers in the Sequence Listing in the present specification show the following sequences:

SEQ ID NO: 1 shows the nucleotide sequence of a DNA fragment encoding an intracellular domain of activin IIA-N receptor protein as bait plasmid used in the two-hybrid method.

20 SEQ ID NO: 2 shows the nucleotide sequence of a DNA fragment as a probe used in Northern hybridization.

SEQ ID NO: 3 shows the nucleotide sequence of a DNA fragment as a probe used in Northern hybridization.

25 SEQ ID NO: 4 shows the nucleotide sequence of a DNA fragment as a probe used in Northern hybridization.

SEQ ID NO: 5 shows the amino acid sequence of the protein of the present invention.

SEQ ID NO: 6 shows the amino acid sequence of the protein of the present invention.

5 SEQ ID NO: 7 shows the nucleotide sequence of a cDNA encoding the protein of the present invention having an amino acid sequence represented by SEQ ID NO: 5.

10 SEQ ID NO: 8 shows the nucleotide sequence of a cDNA encoding the protein of the present invention having an amino acid sequence represented by SEQ ID NO: 6.

The transformant Escherichia coli DH5 α /pBSYN3-6 obtained in Example 1 below has been deposited under FERM BP-6592 from December 2, 1998 with the National Institute of Bioscience and Human-Technology (NIBH), the Agency of Industrial Science of Technology, the Ministry of International Trade and Industry, and under IFO 16221 with Institution Fermentation Organization (IFO) from November 18, 1998.

20 Hereinafter, the present invention is described in more detail by reference to the Reference Example and the Examples, but is not limited thereto. The method of operating the two-hybrid method using yeasts was conducted according to instructions attached to the commercial kit, and the genetic manipulation using E. coli was conducted
25 according to methods described in Molecular Cloning.

Example 1 Cloning of a cDNA encoding the protein of the present invention

(1) Screening of a protein interacting with activin IIA-N receptor protein by the two-hybrid method

5 For the following screening of a protein interacting with activin IIA-N receptor protein from a cDNA library, MATCHMAKER™ Two-Hybrid System 2 (Cat. No. K1604-1, Clontech) was used as a kit.

10 According to the method described above, an Eco RI site and Bam HI site in vector pAS2-1 for DNA ligation were cleaved with suitable restriction enzymes, treated with alkali phosphatase and then purified. Thereafter, a DNA fragment encoding an intracellular domain in the activin IIA-N receptor protein set forth in SEQ ID NO: 1 was
15 ligated thereto, to give a plasmid (pAS-IIA-N) expressing the desired GAL4 DNA-binding domain and activin IIA-N receptor intracellular domain as a fusion protein. As the GAL4 transcription-activating region-fused library plasmid, a commercial mouse brain MATCHMAKER cDNA library (Clontech)
20 was used.

A single colony (diameter of 2 to 3 mm) of yeast strain Y190 was inoculated into 20 ml tryptophan-deficient SD medium and cultured under shaking for 18 hours at 30 °C. 10 ml of this culture was inoculated into 300 ml YPD medium
25 such that its OD₆₀₀ of 0.2 became 0.3, and the yeast was

cultured under shaking at 30 °C for 3 hours. The supernatant was transferred to a sterilized centrifuge tube and centrifuged at 1000×g for 5 minutes at room temperature. The supernatant was discarded, and the cells were suspended
5 in 25 ml sterilized water and centrifuged again at 1000×g for 5 minutes at room temperature. The supernatant was discarded, and the cells were suspended in a mixed solution of 1.5 ml TE buffer (0.01 M Tris-HCl, 1 mM EDTA, pH 7.5)/0.1 M lithium acetate (pH 7.5) and used for the
10 subsequent transformation.

1 ml of the cell suspension prepared above was added to 10 µg of the previously prepared plasmid (pAS-IIA-N) and 2 mg herring spermary carrier DNA (Clontech), and well mixed. Further, 6 ml mixed solution of PEG (40 % PEG
15 4000)/TE buffer/0.1 M lithium acetate was added thereto and mixed in a vortex mixer. After the mixture was shaken at 30 °C at 200 rpm for 30 minutes, 700 µl DMSO (final concentration, 10 %) was added thereto, and the mixture was stirred gently. Thereafter, the mixture was heated at 42
20 °C for 15 minutes while being occasionally shaken, and after the container was cooled on ice, the mixture was centrifuged at 1000×g for 5 minutes. The supernatant was discarded, and the cells were suspended in 0.5 ml TE buffer. 100 µl of the resulting cell suspension was spread onto a
25 tryptophan-deficient SD medium and cultured at 30 °C for 4

days to give a strain maintaining said plasmid stably. Subsequently, the GAL4 transcription-activating region-fused library plasmid was also introduced in the same manner into said yeast strain. 0.2 ml of the resulting transformant was plated on a tryptophan/leucine/histidine-deficient SD medium and cultured at 30 °C for 8 days, and His⁺ colonies were streaked on a tryptophan/leucine/histidine-deficient agar medium.

5 ml mixture of buffer Z (Na₂HPO₄·7H₂O, NaH₂PO₄·H₂O, KCl, MgSO₄·7H₂O, pH 7)/X-gal solution (2 % 5-bromo-4-chloro-3-indolyl-β-D-galactoside solution in DMF)/β-mercaptoethanol was placed in a Petri dish, to moisten a sterilized Whatman #5 filter therein. Another filter was placed on the agar plate containing said transformant colonies, and then this filter was removed and frozen in liquid nitrogen with the colonies facing upward. The filter was removed from liquid nitrogen and then thawed at room temperature, and with the colonies facing upward, the filter was placed on the previously moistened filter. A lid was placed on the Petri dish, and the colonies were incubated at 30 °C for 1 hour, and about 100 blue positive colonies were separated therefrom. The resulting respective positive clones were inoculated into 3 ml leucine-deficient SD liquid medium and cultured for 2 days, and the culture liquid was diluted 10000-fold, plated on a leucine-deficient SD plate, and

incubated at 30 °C for 3 days. 20 to 30 colonies were picked up with a sterilized picker and replicated onto a tryptophan/leucine-deficient SD plate and a leucine-deficient SD plate. Then, those colonies requiring a tryptophan nutrient were selected from the previous positive colonies, and their β -galactosidase activity was assayed to select 2 colonies having no activity.

The resulting true positive colonies were inoculated into 2 ml YPD liquid medium and cultured at 30 °C overnight. The culture was centrifuged for 5 minutes at room temperature, and after the supernatant was discarded, 0.2 ml yeast-lyzing solution (2 % Triton X-100, 1 % SDS, 100 mM NaCl, 10 mM Tris-HCl (pH 8.0), 1 mM EDTA) was added to suspend the microorganisms. 0.2 ml phenol/chloroform/isoamyl alcohol (25 : 24 : 1) and glass beads washed with an acid were added thereto, and the mixture was stirred for 2 minutes with a vortex mixer. After centrifugation at 14000 rpm for 5 minutes at room temperature, the separated supernatant was mixed with a 1/10 volume of 3 M sodium acetate solution (pH 5.2) and a 2.5-fold excess volume of ethanol. The resulting precipitates were washed with 70 % ethanol and dissolved in 20 μ l sterilized water. 1 μ l of this solution was introduced by electroporation into *E. coli* HB101, and the plasmid DNA was purified by the usual mini-prep method to

give the desired cDNA (YN3).

To obtain the full-length cDNA containing a whole coding region of the nucleotide sequence of YN3, the full-length insert in the resulting YN3 was labeled with ^{32}P by the random prime method. Using the labeled YN3 was used as the probe, a mouse brain Lambda cDNA library (Lambda ZAPII vector, Stratagene) was screened by conventional plaque hybridization. The full-length insert in the resulting clones was further used as the probe, and the same screening was carried out repeatedly to give those considered as having the maximum full-length insert containing the whole coding region for the protein of the present invention. One of the resulting positive phages was converted into phagimide ((pBluescript SK) (-) vector) according to the method of Stratagene, and its nucleotide sequence was determined. The full-length cDNA (YN3-6) for the protein of the present invention was 5156 bp encoding a polypeptide [Figs. 1-11] consisting of 1161 amino acids represented by SEQ ID NO: 5 or a polypeptide [Figs. 12-22] consisting of 1112 amino acids represented by SEQ ID NO: 6.

Plasmid pBSYN3-6 containing the full-length cDNA (YN3-6) encoding a polypeptide consisting of amino acids represented by SEQ ID NO: 5 or NO: 6 was transformed into *E. coli* DH5 α , to give a transformant (*E. coli* DH5 α /pBSYN3-6).

Example 2 Detection of expression by Northern

hybridization using poly(A) ⁺RNA derived from a variety of mouse organs

In the insert of YN3-6, the DNA fragments represented by SEQ ID NOS: 2, 3 and 4 were labeled with digoxigenin by use of a DIG-PCR probe synthesis kit (Boehringer) and used as probes.

Further, the brain, liver, spleen, embryo, kidney, hearts, testicles, ovary and skeletal muscles were excised from Balb/c mice, and total RNA was extracted therefrom with a TRIzol reagent (GIBCO BRL). Thereafter, poly(A) ⁺RNAs were purified by use of PolyAT tract mRNA Isolation System (Promega).

1 µg each of poly(A) ⁺RNAs was subjected to 1 % agarose gel electrophoresis in a formalin-gel method and blotted onto Hybond N (Amersham-Pharmacia) by a vacuum blotting method using a blotting unit (Amersham-Pharmacia). The blotted sample was incubated at 65 °C overnight in a hybridization buffer (5×SSC, 0.1 % N-lauroylsarcosine, 0.02 % SDS, 0.5 % blocking reagent (Boehringer), 100 µg/ml salmon testicle DNA) containing 5 ng/ml of each of the previously prepared probe DNAs. Thereafter, the sample was washed 3 times with 0.1×SSC and 0.1 % SDS at 65 °C for 20 minutes. Further, the sample was maintained in a solution (0.1 M Tris-HCl pH 7.5, 0.15 M NaCl) containing 150 mU/ml alkali phosphatase-labeled anti-digoxigenin antibody

(Boehringer), and then washed 3 times with 0.1 M Tris-HCl (pH 7.5), 0.15 M NaCl and 0.1 % Tween 20 for 15 minutes. Finally, the sample was subjected to chemiluminescence using Lumi-Phos 530 (Wako Pure Chemical Industries, Ltd.)
5 as the substrate and detected by exposure onto an X-ray film, and as a result, it was confirmed that a majority of the proteins of the present invention are expressed particularly in the brain [Fig. 23].

Example 3 Interaction between ARIP1 and Smad3

10 In CHO cells transduced with the full-length Smad3 DNA fused with the GAL4 DNA-binding domain and the protein of the present invention (ARIP1) fused with the VP16-activating domain, the specific interaction of ARIP1 with Smad3 was examined in the mammalian two-hybrid system.

15 A DNA construct for the mammalian two-hybrid screening was prepared as follows. That is, for expression of a fusion protein between the GAL4 DNA-binding domain and Smad3, plasmid pBIND was used, and the full-length cDNA encoding human Smad3 was ligated to the plasmid pBIND to
20 construct pBIND-Smad3. Further, for expression of a fusion protein between the VP16-activating domain and ARIP1, plasmid pACT was used. A cDNA fragment having a partial nucleotide sequence (from the 923- to 4446-positions) of the full-length cDNA encoding ARIP1 constructed by ligating
25 the fragment obtained by PCR to a fragment prepared from

plasmid pBS-ARIP1-short having a partial nucleotide sequence (from the 1187- to 4446-positions) of the full-length cDNA encoding ARIP1 was subcloned into pACT, to construct pACT-ARIP1.

5 For the mammalian two-hybrid assay, Check Mate Mammalian Two-Hybrid System (Promega) was used as the kit, and the assay was carried out according to its protocol. The plasmids pBIND-Smad3 and pACT-ARIP1 obtained above, cytomegalovirus promoter-derived β gal (CMV- β gal), and
10 reporter plasmid pG5luc inducing a luciferase gene under the control of a GAL4 response promoter were introduced into CHO cells. The strength of the interaction between Smad3 and ARIP1 in the CHO cells was evaluated where the activity of luciferase expressed by the luciferase gene was
15 used as the indicator. The luciferase activity was measured by the same per se known method (Endocrinology, 136, 5493-5503 (1995)) as in measurement of β -galactosidase activity, and as a result, the protein of the present invention (ARIP1), similar to the activin receptor IIA,
20 showed direct interaction with Smad3 in the mammalian cells [Fig. 24].

Industrial Applicability

25 The protein of the present invention, a partial peptide thereof or a salt thereof (also referred to

hereinafter as the protein of the present invention), the DNA encoding the protein of the present invention or a partial peptide thereof (also referred to hereinafter as the DNA of the present invention), the antibody against the protein of the present invention and the antisense DNA can be used as reagents in (1) a method of determining a binding protein to the protein of the present invention, (2) construction of a system for expressing a recombinant protein, (3) development of a binding assay system using said expression system and an assay system using the two-hybrid method and screening of candidates for pharmaceutical compounds, (4) practice of a drug design based on comparison with structurally analogous ligand receptors and (5) formation of probes and PCR primers in gene diagnosis, etc., and further can be used as chemicals in (6) gene therapy, etc. In particular, elucidation of the structure and properties of the protein of the present invention leads to development of unique pharmaceutical preparations acting on these systems.

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SEQUENCE LISTING

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 25 Ser Ile Glu Pro Pro Glu Glu Glu Glu Glu Glu Arg Pro Val Val Asn
 115 120 125
 Gly Asn Gly Val Val Ile Thr Pro Glu Ser Ser Glu His Glu Asp Lys
 130 135 140
 Ser Ala Gly Ala Ser Gly Glu Thr Pro Ser Gln Pro Tyr Pro Ala Pro

6/19

	145		150		155		160									
	Val	Tyr	Ser	Gln	Pro	Glu	Glu	Leu	Lys	Asp	Gln	Met	Asp	Asp	Thr	Lys
				165				170					175			
	Pro	Thr	Lys	Pro	Glu	Glu	Asn	Glu	Asp	Ser	Asp	Pro	Leu	Pro	Asp	Asn
5			180					185					190			
	Trp	Glu	Met	Ala	Tyr	Thr	Glu	Lys	Gly	Glu	Val	Tyr	Phe	Ile	Asp	His
			195					200					205			
	Asn	Thr	Lys	Thr	Thr	Ser	Trp	Leu	Asp	Pro	Arg	Leu	Ala	Lys	Lys	Ala
		210					215					220				
10	Lys	Pro	Pro	Glu	Glu	Cys	Lys	Glu	Asn	Glu	Leu	Pro	Tyr	Gly	Trp	Glu
	225					230				235				240		
	Lys	Ile	Asp	Asp	Pro	Ile	Tyr	Gly	Thr	Tyr	Tyr	Val	Asp	His	Ile	Asn
				245						250				255		
	Arg	Arg	Thr	Gln	Phe	Glu	Asn	Pro	Val	Leu	Glu	Ala	Lys	arg	Lys	Leu
15			260					265					270			
	Gln	Gln	His	Asn	Met	Pro	His	Thr	Glu	Leu	Gly	Ala	Lys	Pro	Leu	Gln
			275					280					285			
	Ala	Pro	Gly	Phe	Arg	Glu	Lys	Pro	Leu	Phe	Thr	Arg	Asp	Ala	Ser	Gln
		290					295					300				
20	Leu	Lys	Gly	Thr	Phe	Leu	Ser	Thr	Thr	Leu	Lys	Lys	Ser	Asn	Met	Gly
	305					310				315				320		
	Phe	Gly	Phe	Thr	Ile	Ile	Gly	Gly	Asp	Glu	Pro	Asp	Glu	Phe	Leu	Gln
				325						330				335		
	Val	Lys	Ser	Val	Ile	Pro	Asp	Gly	Pro	Ala	Ala	Gln	Asp	Gly	Lys	Met
25			340					345					350			
	Glu	Thr	Gly	Asp	Val	Ile	Val	Tyr	Ile	Asn	Glu	Val	Cys	Val	Leu	Gly
			355					360					365			
	His	Thr	His	Ala	Asp	Val	Val	Lys	Leu	Phe	Gln	Ser	Val	Pro	Ile	Gly
		370					375					380				

7/19

Gln Ser Val Asn Leu Val Leu Cys Arg Gly Tyr Pro Leu Pro Phe Asp
 385 390 395 400
 Pro Glu Asp Pro Ala Asn Ser Met Val Pro Pro Leu Ala Ile Met Glu
 405 410 415
 5 Arg Pro Pro Pro Val Met Val Asn Gly Arg His Asn Tyr Glu Thr Tyr
 420 425 430
 Leu Glu Tyr Ile Ser Arg Thr Ser Gln Ser Val Pro Asp Ile Thr Asp
 435 440 445
 Arg Pro Pro His Ser Leu his Ser Met Pro Ala Asp Gly Gln Leu Asp
 10 450 455 460
 Gly Thr Tyr Pro Pro Pro Val His Asp Asp Asn Val Ser Met Ala Ser
 465 470 475 480
 Ser Gly Ala Thr Gln Ala Glu Leu Met Thr Leu Thr Ile Val Lys Gly
 485 490 495
 15 Ala Gln Gly Phe Gly Phe Thr Ile Ala Asp Ser Pro Thr Gly Gln Arg
 500 505 510
 Val Lys Gln Ile Leu Asp Ile Gln Gly Cys Pro Gly Leu Cys Glu Gly
 515 520 525
 Asp Leu Ile Val Glu Ile Asn Gln Gln Asn Val Gln Asn Leu Ser His
 20 530 535 540
 Thr Glu Val Val Asp Ile Leu Lys Asp Cys Pro Val Gly Ser Glu Thr
 545 550 555 560
 Ser Leu Ile Ile His Arg Gly Gly Phe Phe Ser Pro Trp Lys Thr Pro
 565 570 575
 25 Lys Pro Met Met Asp Arg Trp Glu Asn Gln Gly Ser Pro Gln Thr Ser
 580 585 590
 Leu Ser Ala Pro Ala Val Pro Gln Asn Leu Pro Phe Pro Pro Ala Leu
 595 600 605
 His Arg Ser Ser Phe Pro Asp Ser Thr Glu Ala Phe Asp Pro Arg Lys

8/19

	610		615		620														
	Pro	Asp	Pro	Tyr	Glu	Leu	Tyr	Glu	Lys	Ser	Arg	Ala	Ile	Tyr	Glu	Ser			
	625					630					635					640			
	Arg	Gln	Gln	Val	Pro	Pro	Arg	Thr	Ser	Phe	Arg	Met	Asp	Ser	Ser	Gly			
5					645					650						655			
	Pro	Asp	Tyr	Lys	Glu	Leu	Asp	Val	His	Leu	Arg	Arg	Met	Glu	Ser	Gly			
					660					665						670			
	Phe	Gly	Phe	Arg	Ile	Leu	Gly	Gly	Asp	Glu	Pro	Gly	Gln	Pro	Ile	Leu			
					675					680						685			
10	Ile	Gly	Ala	Val	Ile	Ala	Met	Gly	Ser	Ala	Asp	Arg	Asp	Gly	Arg	Leu			
					690					695						700			
	His	Pro	Gly	Asp	Glu	Leu	Val	Tyr	Val	Asp	Gly	Ile	Pro	Val	Ala	Gly			
	705					710					715					720			
	Lys	Thr	His	Arg	Tyr	Val	Ile	Asp	Leu	Met	His	His	Ala	Ala	Arg	Asn			
15					725					730						735			
	Gly	Gln	Val	Asn	Leu	Thr	Val	Arg	Arg	Lys	Val	Leu	Cys	Gly	Gly	Glu			
					740					745						750			
	Pro	Cys	Pro	Glu	Asn	Gly	Arg	Ser	Pro	Gly	Ser	Val	Ser	Thr	His	His			
					755					760						765			
20	Ser	Ser	Pro	Arg	Ser	Asp	Tyr	Ala	Thr	Tyr	Ser	Asn	Ser	Asn	His	Ala			
					770					775						780			
	Ala	Pro	Ser	Ser	Asn	Ala	Ser	Pro	Pro	Glu	Gly	Phe	Ala	Ser	His	Ser			
	785					790					795					800			
	Leu	Gln	Thr	Ser	Asp	Val	Val	Ile	His	Arg	Lys	Glu	Asn	Glu	Gly	Phe			
25					805					810						815			
	Gly	Phe	Val	Ile	Ile	Ser	Ser	Leu	Asn	Arg	Pro	Glu	Ser	Gly	Ala	Thr			
					820					825						830			
	Ile	Thr	Val	Pro	His	Lys	Ile	Gly	Arg	Ile	Ile	Asp	Gly	Ser	Pro	Ala			
					835					840						845			

9/19

	Asp	Arg	Cys	Ala	Lys	Leu	Lys	Val	Gly	Asp	Arg	Ile	Leu	Ala	Val	Asn
	850						855							860		
	Gly	Gln	Ser	Ile	Ile	Asn	Met	Pro	His	Ala	Asp	Ile	Val	Lys	Leu	Ile
	865					870					875				880	
5	Lys	Asp	Ala	Gly	Leu	Ser	Val	Thr	Leu	Arg	Ile	Ile	Pro	Gln	Glu	Glu
						885					890				895	
	Leu	Asn	Ser	Pro	Thr	Ser	Ala	Pro	Ser	Ser	Glu	Lys	Gln	Ser	Pro	Met
					900					905					910	
	Ala	Gln	Gln	His	Ser	Pro	Leu	Ala	Gln	Gln	Ser	Pro	Leu	Ala	Gln	Pro
10			915						920						925	
	Ser	Pro	Ala	Thr	Pro	Asn	Ser	Pro	Val	Ala	Gln	Pro	Ala	Pro	Pro	Gln
			930						935					940		
	Pro	Leu	Gln	Leu	Gln	Gly	His	Glu	Asn	Ser	Tyr	Arg	Ser	Glu	Val	Lys
	945					950					955				960	
15	Ala	Arg	Gln	Asp	Val	Lys	Pro	Asp	Ile	Arg	Gln	Pro	Pro	Phe	Thr	Asp
						965					970				975	
	Tyr	Arg	Gln	Pro	Pro	Leu	Asp	Tyr	Arg	Gln	Pro	Pro	Gly	Gly	Asp	Tyr
					980					985				990		
	Ser	Gln	Pro	Pro	Pro	Leu	Asp	Tyr	Arg	Gln	His	Ser	Pro	Asp	Tyr	Arg
20			995						1000					1005		
	Gln	Tyr	Pro	Leu	Ser	Asp	Tyr	Arg	Gln	Pro	Gln	Asp	Phe	Asp	Tyr	Phe
			1010						1015					1020		
	Thr	Val	Asp	Met	Glu	Lys	Gly	Ala	Lys	Gly	Phe	Gly	Phe	Ser	Ile	Arg
	1025					1030					1035				1040	
25	Gly	Gly	Arg	Glu	Tyr	Lys	Met	Asp	Leu	Tyr	Val	Leu	Arg	Leu	Ala	Glu
						1045					1050				1055	
	Asp	Gly	Pro	Ala	Ile	Arg	Asn	Gly	Arg	Met	Arg	Val	Gly	Asp	Gln	Ile
					1060					1065					1070	
	Ile	Glu	Ile	Asn	Gly	Glu	Ser	Thr	Arg	Asp	Met	Thr	His	Ala	Arg	Ala

10/19

	1075	1080	1085
	Ile Glu Leu Ile Lys Ser Gly Gly Arg Arg Val Arg Leu Leu Leu Lys		
	1090	1095	1100
	Arg Gly Thr Gly Gln Val Pro Glu Tyr Gly Met Val Pro Ser Ser Leu		
5	1105	1110	1115 1120
	Ser Met Cys Met Lys Ser Asp Lys His Gly Ser Pro Tyr Phe Tyr Leu		
	1125	1130	1135
	Leu Gly His Pro Lys Asp Thr Thr Asn Pro Thr Pro Gly Val Leu Pro		
	1140	1145	1150
10	Leu Pro Pro Pro Gln Ala Cys Arg Lys		
	1155	1160 1161	
	<210>6		
	<211>1112		
	<212>PRT		
15	<213>Mouse		
	<400>6		
	Met Glu Leu Glu Lys Ser Gly Ala Leu Leu Glu Ser Gly Thr Tyr Glu		
	5	10	15
	Asp Asn Tyr Tyr Gly Thr Pro lys Pro Pro Ala Glu Pro Ala Pro Leu		
20	20	25	30
	Leu Asn Val Thr Asp Gln Ile Leu Pro Gly Ala Thr Pro Ser Ala Glu		
	35	40	45
	Gly Lys Arg Lys Arg Asn Lys Ser Val Thr Asn Met Glu Lys Ala Ser		
	50	55	60
25	Ile Glu Pro Pro Glu Glu Glu Glu Glu Glu Arg Pro Val Val Asn Gly		
	65	70	75 80
	Asn Gly Val Val Ile Thr Pro Glu Ser Ser Glu His Glu Asp Lys Ser		
	85	90	95
	Ala Gly Ala Ser Gly Glu Thr Pro Ser Gln Pro Tyr Pro Ala Pro Val		

11/19

	100	105	110
	Tyr Ser Gln Pro Glu Glu Leu Lys Asp Gln Met Asp Asp Thr Lys Pro		
	115	120	125
	Thr Lys Pro Glu Glu Asn Glu Asp Ser Asp Pro Leu Pro Asp Asn Trp		
5	130	135	140
	Glu Met Ala Tyr Thr Glu Lys Gly Glu Val Tyr Phe Ile Asp His Asn		
	145	150	155
	Thr Lys Thr Thr Ser Trp Leu Asp Pro Arg Leu Ala Lys Lys Ala Lys		
	165	170	175
10	Pro Pro Glu Glu Cys Lys Glu Asn Glu Leu Pro Tyr Gly Trp Glu Lys		
	180	185	190
	Ile Asp Asp Pro Ile Tyr Gly Thr Tyr Tyr Val Asp His Ile Asn Arg		
	195	200	205
	Arg Thr Gln Phe Glu Asn Pro Val Leu Glu Ala Lys arg Lys Leu Gln		
15	210	215	220
	Gln His Asn Met Pro His Thr Glu Leu Gly Ala Lys Pro Leu Gln Ala		
	225	230	235
	Pro Gly Phe Arg Glu Lys Pro Leu Phe Thr Arg Asp Ala Ser Gln Leu		
	245	250	255
20	Lys Gly Thr Phe Leu Ser Thr Thr Leu Lys Lys Ser Asn Met Gly Phe		
	260	265	270
	Gly Phe Thr Ile Ile Gly Gly Asp Glu Pro Asp Glu Phe Leu Gln Val		
	275	280	285
	Lys Ser Val Ile Pro Asp Gly Pro Ala Ala Gln Asp Gly Lys Met Glu		
25	290	295	300
	Thr Gly Asp Val Ile Val Tyr Ile Asn Glu Val Cys Val Leu Gly His		
	305	310	315
	Thr His Ala Asp Val Val Lys Leu Phe Gln Ser Val Pro Ile Gly Gln		
	325	330	335

12/19

Ser Val Asn Leu Val Leu Cys Arg Gly Tyr Pro Leu Pro Phe Asp Pro
 340 345 350
 Glu Asp Pro Ala Asn Ser Met Val Pro Pro Leu Ala Ile Met Glu Arg
 355 360 365
 5 Pro Pro Pro Val Met Val Asn Gly Arg His Asn Tyr Glu Thr Tyr Leu
 370 375 380
 Glu Tyr Ile Ser Arg Thr Ser Gln Ser Val Pro Asp Ile Thr Asp Arg
 385 390 395 400
 Pro Pro His Ser Leu his Ser Met Pro Ala Asp Gly Gln Leu Asp Gly
 10 405 410 415
 Thr Tyr Pro Pro Pro Val His Asp Asp Asn Val Ser Met Ala Ser Ser
 420 425 430
 Gly Ala Thr Gln Ala Glu Leu Met Thr Leu Thr Ile Val Lys Gly Ala
 435 440 445
 15 Gln Gly Phe Gly Phe Thr Ile Ala Asp Ser Pro Thr Gly Gln Arg Val
 450 455 460
 Lys Gln Ile Leu Asp Ile Gln Gly Cys Pro Gly Leu Cys Glu Gly Asp
 465 470 475 480
 Leu Ile Val Glu Ile Asn Gln Gln Asn Val Gln Asn Leu Ser His Thr
 20 485 490 495
 Glu Val Val Asp Ile Leu Lys Asp Cys Pro Val Gly Ser Glu Thr Ser
 500 505 510
 Leu Ile Ile His Arg Gly Gly Phe Phe Ser Pro Trp Lys Thr Pro Lys
 515 520 525
 25 Pro Met Met Asp Arg Trp Glu Asn Gln Gly Ser Pro Gln Thr Ser Leu
 530 535 540
 Ser Ala Pro Ala Val Pro Gln Asn Leu Pro Phe Pro Pro Ala Leu His
 545 550 555 560
 Arg Ser Ser Phe Pro Asp Ser Thr Glu Ala Phe Asp Pro Arg Lys Pro

13/19

	565	570	575
	Asp Pro Tyr Glu Leu Tyr Glu Lys Ser Arg Ala Ile Tyr Glu Ser Arg		
	580	585	590
	Gln Gln Val Pro Pro Arg Thr Ser Phe Arg Met Asp Ser Ser Gly Pro		
5	595	600	605
	Asp Tyr Lys Glu Leu Asp Val His Leu Arg Arg Met Glu Ser Gly Phe		
	610	615	620
	Gly Phe Arg Ile Leu Gly Gly Asp Glu Pro Gly Gln Pro Ile Leu Ile		
	625	630	635
10	Gly Ala Val Ile Ala Met Gly Ser Ala Asp Arg Asp Gly Arg Leu His		
	645	650	655
	Pro Gly Asp Glu Leu Val Tyr Val Asp Gly Ile Pro Val Ala Gly Lys		
	660	665	670
	Thr His Arg Tyr Val Ile Asp Leu Met His His Ala Ala Arg Asn Gly		
15	675	680	685
	Gln Val Asn Leu Thr Val Arg Arg Lys Val Leu Cys Gly Gly Glu Pro		
	690	695	700
	Cys Pro Glu Asn Gly Arg Ser Pro Gly Ser Val Ser Thr His His Ser		
	705	710	715
20	Ser Pro Arg Ser Asp Tyr Ala Thr Tyr Ser Asn Ser Asn His Ala Ala		
	725	730	735
	Pro Ser Ser Asn Ala Ser Pro Pro Glu Gly Phe Ala Ser His Ser Leu		
	740	745	750
	Gln Thr Ser Asp Val Val Ile His Arg Lys Glu Asn Glu Gly Phe Gly		
25	755	760	765
	Phe Val Ile Ile Ser Ser Leu Asn Arg Pro Glu Ser Gly Ala Thr Ile		
	770	775	780
	Thr Val Pro His Lys Ile Gly Arg Ile Ile Asp Gly Ser Pro Ala Asp		
	785	790	795
			800

14/19

	Arg	Cys	Ala	Lys	Leu	Lys	Val	Gly	Asp	Arg	Ile	Leu	Ala	Val	Asn	Gly
						805				810					815	
	Gln	Ser	Ile	Ile	Asn	Met	Pro	His	Ala	Asp	Ile	Val	Lys	Leu	Ile	Lys
					820				825					830		
5	Asp	Ala	Gly	Leu	Ser	Val	Thr	Leu	Arg	Ile	Ile	Pro	Gln	Glu	Glu	Leu
					835				840					845		
	Asn	Ser	Pro	Thr	Ser	Ala	Pro	Ser	Ser	Glu	Lys	Gln	Ser	Pro	Met	Ala
					850			855				860				
	Gln	Gln	His	Ser	Pro	Leu	Ala	Gln	Gln	Ser	Pro	Leu	Ala	Gln	Pro	Ser
10	865					870				875					880	
	Pro	Ala	Thr	Pro	Asn	Ser	Pro	Val	Ala	Gln	Pro	Ala	Pro	Pro	Gln	Pro
					885					890				895		
	Leu	Gln	Leu	Gln	Gly	His	Glu	Asn	Ser	Tyr	Arg	Ser	Glu	Val	Lys	Ala
					900				905					910		
15	Arg	Gln	Asp	Val	Lys	Pro	Asp	Ile	Arg	Gln	Pro	Pro	Phe	Thr	Asp	Tyr
					915			920					925			
	Arg	Gln	Pro	Pro	Leu	Asp	Tyr	Arg	Gln	Pro	Pro	Gly	Gly	Asp	Tyr	Ser
					930			935				940				
	Gln	Pro	Pro	Pro	Leu	Asp	Tyr	Arg	Gln	His	Ser	Pro	Asp	Tyr	Arg	Gln
20	945					950				955					960	
	Tyr	Pro	Leu	Ser	Asp	Tyr	Arg	Gln	Pro	Gln	Asp	Phe	Asp	Tyr	Phe	Thr
					965					970				975		
	Val	Asp	Met	Glu	Lys	Gly	Ala	Lys	Gly	Phe	Gly	Phe	Ser	Ile	Arg	Gly
					980				985				990			
25	Gly	Arg	Glu	Tyr	Lys	Met	Asp	Leu	Tyr	Val	Leu	Arg	Leu	Ala	Glu	Asp
					995			1000				1005				
	Gly	Pro	Ala	Ile	Arg	Asn	Gly	Arg	Met	Arg	Val	Gly	Asp	Gln	Ile	Ile
					1010			1015				1020				
	Glu	Ile	Asn	Gly	Glu	Ser	Thr	Arg	Asp	Met	Thr	His	Ala	Arg	Ala	Ile

15/19

	1025		1030		1035		1040									
	Glu	Leu	Ile	Lys	Ser	Gly	Gly	Arg	Arg	Val	Arg	Leu	Leu	Leu	Lys	Arg
				1045				1050							1055	
	Gly	Thr	Gly	Gln	Val	Pro	Glu	Tyr	Gly	Met	Val	Pro	Ser	Ser	Leu	Ser
5				1060				1065							1070	
	Met	Cys	Met	Lys	Ser	Asp	Lys	His	Gly	Ser	Pro	Tyr	Phe	Tyr	Leu	Leu
				1075				1080							1085	
	Gly	His	Pro	Lys	Asp	Thr	Thr	Asn	Pro	Thr	Pro	Gly	Val	Leu	Pro	Leu
				1090				1095							1100	
10	Pro	Pro	Pro	Gln	Ala	Cys	Arg	Lys								
	1105							1110			1112					
	<210>7															
	<211>3483															
	<212>DNA															
15	<213>Mouse															
	<400>7															
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	CTGAGCTCAG	CTCCAGGCAC	CACAAGGCCA	CATAAGGAGG	GTGAGGTCCC	TGGAGTGGAC	120									
	TACATTTTCA	TAACCGTTGA	GGAGTTTATG	GAATTGGAGA	AAAGTGGTGC	TCTCCTAGAA	180									
20	AGCGGGACCT	ATGAAGACAA	CTACTACGGT	ACCCCGAAGC	CTCCAGCTGA	ACCAGCACCA	240									
	TTATTAAATG	TAACAGACCA	GATACTTCCG	GGAGCTACTC	CAAGTGCTGA	GGGGAAGCGG	300									
	AAAAGAAATA	AGTCAGTGAC	CAACATGGAG	AAAGCAAGTA	TAGAGCCTCC	AGAGGAGGAA	360									
	GAAGAAGAAA	GGCCTGTAGT	CAATGGAAAC	GGCGTGGTCA	TAACCCCA	ATCCAGTGAA	420									
	CATGAAGACA	AAAGTGCAGG	TGCCTCAGGG	GAGACACCCT	CCCAGCCTTA	CCCTGCACCC	480									
25	GTGTACAGCC	AGCCCGAAGA	GCTCAAGGAC	CAGATGGACG	ATACAAAGCC	AACAAAGCCT	540									
	GAGGAGAACG	AGGACTCTGA	TCCATTGCCT	GATAACTGGG	AAATGGCCTA	CACAGAGAAG	600									
	GGGGAAGTCT	ACTTCATTGA	CCATAACACA	AAGACAACAT	CATGGCTGGA	TCCGCGACTT	660									
	GCGAAAAGG	CTAAACCTCC	AGAAGAGTGC	AAAGAAAATG	AGCTTCCATA	TGGCTGGGAA	720									
	AAAATCGATG	ATCCTATATA	TGGCACTTAC	TATGTTGACC	ACATAAATAG	AAGAACACAG	780									

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 GATGCATCCC AGTTGAAGGG AACGTTCTC AGCACCACCC TCAAAAAGAG CAACATGGGC 960
 TTTGGGTTTA CCATAATTGG TGGAGACGAG CCGGATGAGT TTCTACAGGT GAAAAGTGTG 1020
 5 ATCCCGGATG GGCCTGCCGC ACAGGATGGG AAAATGGAGA CAGGTGATGT CATTGTCTAT 1080
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17/19

CGAATCATTG ATGGGAGCCC TGCAGATCGC TGTGCCAAAC TCAAAGTGGG CGACCGTATC 2580
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 AAGGACGCCG GTCTCAGTGT CACCCTTCGC ATCATTCCTC AGGAGGAGCT CAACAGCCCA 2700
 ACATCAGCAC CCAGTTCAGA GAAACAGAGC CCCATGGCCC AGCAGCACAG CCCTCTGGCC 2760
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20 <212>DNA

<213>Mouse

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 25 CCGGGAGCTA CTCCAAGTGC TGAGGGGAAG CGGAAAAGAA ATAAGTCAGT GACCAACATG 180
 GAGAAAGCAA GTATAGAGCC TCCAGAGGAG GAAGAAGAAG AAAGGCCTGT AGTCAATGGA 240
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 TGCAAAGAAA ATGAGCTTCC ATATGGCTGG GAAAAAATCG ATGATCCTAT ATATGGCACT 600
 TACTATGTTG ACCACATAAA TAGAAGAACA CAGTTTGAAA ACCCTGTCCT GGAAGCAAAA 660
 5 AGGAAGCTAC AGCAACATAA CATGCCCCAC ACAGAACTTG GAGCAAAGCC CCTGCAGGCC 720
 CCAGGTTTCC GAGAAAAGCC ACTCTTCACC CGGGATGCAT CCCAGTTGAA GGGAACGTTC 780
 CTCAGCACCA CCCTCAAAAA GAGCAACATG GGCTTTGGGT TTACCATAAT TGGTGGAGAC 840
 GAGCCGGATG AGTTTCTACA GGTGAAAAGT GTGATCCCGG ATGGGCCTGC CGCACAGGAT 900
 GGGAAAATGG AGACAGGTGA TGTCATTGTC TATATTAATG AAGTTTGTGT CCTTGGACAC 960
 10 ACTCATGCAG ATGTTGTCAA ACTTTTCCAG TCTGTTCCCTA TTGGTCAGAG TGTCAACTTG 1020
 GTGTTGTGTC GTGGCTACCC TTTGCCCTTT GACCCTGAAG ATCCTGCTAA CAGCATGGTG 1080
 CCACCCCTTG CAATAATGGA GAGGCCACCT CCGGTGATGG TCAATGGAAG ACATAACTAT 1140
 GAAACATACT TGGAATACAT TTCTCGGACC TCACAGTCGG TCCCAGATAT TACAGACCGG 1200
 CCACCTCATT CTTTGCCTC CATGCCAGCT GACGGCCAGC TAGATGGCAC GTATCCACCA 1260
 15 CCCGTCCATG ACGACAATGT GTCTATGGCT TCGTCTGGAG CCACTCAAGC TGAAGTTATG 1320
 ACCTTAACCA TTGTGAAAGG TGCCCAGGGA TTTGGCTTTA CTATTGCCGA CAGTCCCACG 1380
 GGACAGCGGG TGAAACAAAT CCTTGACATT CAGGGATGCC CTGGGCTGTG TGAAGGAGAC 1440
 CTCATTGTTG AGATCAACCA ACAGAATGTA CAGAACCTGA GCCATACAGA AGTAGTGGAT 1500
 AACTTAAGG ACTGCCCCGT TGGAAGTGAG ACTTCTTTAA TCATCCATCG AGGAGGTTTC 1560
 20 TTTTCTCCAT GGAAAACTCC AAAGCCTATG ATGGACCGAT GGGAGAACCA AGGCAGTCCA 1620
 CAAACAAGTT TATCTGCTCC GGCCGTCCCA CAGAACCTGC CCTTCCCACC TGCCCTTCAC 1680
 AGGAGCTCCT TTCCTGATTC AACAGAGGCC TTTGACCCAC GGAAGCCTGA CCCATATGAG 1740
 CTCTACGAGA AATCGAGAGC CATTTATGAA AGTAGGCAAC AAGTGCCACC CAGGACCAGT 1800
 TTTCGAATGG ATTCCTCTGG TCCAGATTAT AAGGAACTGG ATGTTACCT TCGGAGGATG 1860
 25 GAGTCTGGAT TTGGCTTTAG AATCCTTGGG GGAGATGAAC CTGGACAGCC TATTTTGATC 1920
 GGAGCCGTCA TTGCCATGGG CTCAGCTGAC AGAGACGGCC GTCTACACCC AGGAGATGAG 1980
 CTTGTCTATG TCGATGGGAT CCCAGTGGCT GGCAAGACCC ACCGCTATGT CATCGACCTC 2040
 ATGCACCACG CGGCCCCGAA TGGGCAGGTT AACCTCACTG TGAGAAGAAA GGTGCTATGT 2100
 GGAGGGGAGC CCTGCCCAGA GAATGGGAGG AGTCCAGGCT CTGTATCAAC TCACCACAGC 2160

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TCTCCGCGCA GTGACTATGC CACCTACTCC AACAGCAACC ACGCCGCCCC CAGCAGCAAT 2220
GCCTCACCTC CTGAAGGCTT TGCCTCACAC AGCTTGCAGA CCAGTGATGT GGTCATTAC 2280
CGCAAAGAAA ACGAAGGGTT TGGCTTCGTC ATCATCAGCT CTCTGAACAG GCCTGAGTCT 2340
GGAGCCACCA TAACTGTGCC CCATAAAATT GGACGAATCA TTGATGGGA GCCCTGCAGAT 2400
5 CGCTGTGCCA AACTCAAAGT GGGCGACCGT ATCTTAGCAG TCAACGGCCA GTCTATCATC 2460
AACATGCCTC ACGCTGACAT TGTGAAGCTC ATCAAGGACG CCGGTCTCAG TGTACCCTT 2520
CGCATCATTC CTCAGGAGGA GCTCAACAGC CCAACATCAG CACCCAGTTC AGAGAAACAG 2580
AGCCCCATGG CCCAGCAGCA CAGCCCTCTG GCCCAGCAGA GTCCTCTGGC CCAGCCAAGC 2640
CCCGCCACCC CCAACAGCCC AGTCGCACAG CCAGCTCCTC CCCAACCTCT CCAGCTGCAA 2700
10 GGACACGAAA ATAGTTACAG GTCAGAAGTT AAAGCGAGGC AAGATGTGAA GCCAGACATC 2760
CGGCAGCCTC CCTTCACAGA CTACAGGCAG CCCCCGCTGG ACTACAGGCA GCCCCCGGGA 2820
GGAGACTACT CACAGCCCCC ACCCTTGGAC TACAGGCAGC ACTCTCCAGA CACCAGGCAG 2880
TACCCTCTGT CAGACTACAG GCAGCCACAG GATTTTGATT ATTTCACTGT GGACATGGAG 2940
AAAGGAGCCA AAGGATTTGG ATTCAGCATT CGTGGAGGAA GGAATACAA GATGGATCTG 3000
15 TATGTGTTGA GATTGGCAGA GGATGGGCCA GCCATAAGGA ACGGCAGGAT GAGGGTAGGA 3060
GATCAGATCA TTGAAATAAA TGGGGAAAGC ACACGAGACA TGACCCACGC CAGAGCAATA 3120
GAACTCATCA AGTCTGGAGG AAGAAGAGTG CGGCTGCTGC TGAAGAGAGG CACGGGGCAG 3180
GTCCCGGAGT ATGGAATGGT ACCTTCCAGC CTCTCCATGT GCATGAAAAG TGACAAGCAT 3240
GGGTCCCAT ATTTCTACTT ACTGGGCCAC CCTAAAGACA CGACGAACCC CACGCCTGGA 3300
20 GTGCTGCCGC TGCCGCCGCC CCAGGCCTGC CGGAAG 3336

CLAIMS

1. A protein which has an amino acid sequence identical to or substantially identical to an amino acid sequence represented by SEQ ID NO: 5, or a salt thereof.
- 5 2. A protein which has an amino acid sequence identical to or substantially identical to an amino acid sequence represented by SEQ ID NO: 6, or a salt thereof.
3. The protein according to claim 1 or 2, which has PDZ domains and WW domains and is expressed specifically in the
10 brain and has an ability to bind to activin receptors and/or activin intracellular information transmission molecules.
4. The protein according to claim 3, wherein the activin intracellular information transmission molecule is Smad3.
- 15 5. The protein according to claim 1 or 2, which has 5 PDZ domains and 2 WW domains and is expressed specifically in the brain and has an ability to bind to activin receptors and Smad3.
6. A partial peptide of the protein according to claim 1, a
20 partial peptide of the protein according to claim 2, or a salt thereof.
7. A recombinant DNA which comprises a DNA having a nucleotide sequence encoding the protein according to claim 1 or the protein according to claim 2.
- 25 8. The DNA according to claim 7, which has a nucleotide

sequence represented by SEQ ID NO: 7, a nucleotide sequence represented by SEQ ID NO: 8 or a nucleotide sequence hybridizing therewith under high stringent conditions.

5 9. A recombinant DNA which comprises a DNA having a nucleotide sequence encoding the partial peptide according to claim 6.

10. A recombinant vector which comprises the DNA according to claim 7.

10 11. A transformant comprising the recombinant vector according to claim 10.

12. A method for producing the protein according to claim 1, the protein according to claim 2 or a salt thereof, which comprises culturing the transformant according to claim 11 to produce and accumulate the protein according to claim 1 or the protein according to claim 2 is formed and accumulated, and then recovering the product.

13. An antibody against the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof.

20 14. A method for quantifying the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof, which comprises allowing a test solution containing the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof and the

25

labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof to react competitively with the antibody according to claim 13.

5 15. A method for determining a binding protein to the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof, which comprises using the protein according to claim 1, the protein according to claim 2, the partial
10 peptide according to claim 6 or a salt thereof.

16. The method according to claim 15, which comprises introducing (1) an expression vector which fuses the protein according to claim 1, the protein according to claim 2 or the partial peptide according to claim 6 with a
15 DNA-binding region of a transcriptional factor and (2) a fusion library between a gene encoding a test protein and a transcription-activating region, into a host cell having a reporter gene maintaining a region binding to the transcriptional factor on a promoter, and measuring a
20 change in the amount of the expressed reporter gene which is increased by a binding of the protein according to claim 1, the protein according to claim 2 or the partial peptide according to claim 6 to the test compound.

17. A protein or a salt thereof which binds to the protein
25 according to claim 1, the protein according to claim 2, the

partial peptide according to claim 6 or a salt thereof,
which is obtained by the method according to claim 15.

18. A method for screening a compound or a salt thereof
inhibiting or promoting a binding of the protein according
5 to claim 1, the protein according to claim 2, the partial
peptide according to claim 6 or a salt thereof to the
protein according to claim 17 or a salt thereof, activin
receptors or activin intracellular information transmission
molecules, which comprises using the protein according to
10 claim 1, the protein according to claim 2, the partial
peptide according to claim 6 or a salt thereof.

19. A method for screening a compound or a salt thereof
inhibiting or promoting a binding of the protein according
to claim 1, the protein according to claim 2, the partial
15 peptide according to claim 6 or a salt thereof to activin
receptors or activin intracellular information transmission
molecules, which comprises comparing the case where the
labeled protein according to claim 1, the labeled protein
according to claim 2, the labeled partial peptide according
20 to claim 6 or a labeled salt thereof is contacted with
activin receptors or activin intracellular information
transmission molecules, with the case where the labeled
protein according to claim 1, the labeled protein according
to claim 2, the labeled partial peptide according to claim
25 6 or a labeled salt thereof and a test compound are

contacted with activin receptors or activin intracellular information transmission molecules, by measuring the amount of the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof bound to the activin receptors or activin intracellular information transmission molecules in both the cases.

20. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof to the protein according to claim 17 or a salt thereof, which comprises comparing the case where the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof is contacted with the protein according to claim 17 or a salt thereof, with the case where the labeled protein according to claim 1, the labeled protein according to claim 2 or a labeled salt thereof and a test compound are contacted with the protein according to claim 17 or a salt thereof, by measuring the amount of the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof bound to the protein according to claim 17 or a salt thereof in both the cases.

21. A method for screening a compound or a salt thereof inhibiting or promoting a binding of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof to the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which comprises comparing the case where the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof is introduced into cells expressing the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, with the case where the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof and a test compound are introduced into cells expressing the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the amount of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof bound to the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules in the cells in both the cases.

22. A method for screening a compound or a salt thereof

inhibiting or promoting a binding of the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof to the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which comprises comparing the case where the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof is contacted with a membrane fraction of cells expressing the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, with the case where the labeled protein according to claim 1, the labeled protein according to claim 2, the labeled partial peptide according to claim 6 or a labeled salt thereof and a test compound are contacted with a membrane fraction of cells expressing the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the amount of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof bound to the membrane fraction of the cells in both the cases.

23. A method for screening a compound or a salt thereof

inhibiting or promoting a binding of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof to the protein according to claim 17 or a salt thereof, activin
5 receptors or activin intracellular information transmission molecules, which comprises comparing the case where the protein according to claim 1, the protein according to claim 2, or a salt thereof is introduced into cells expressing the protein according to claim 17 or a salt
10 thereof, activin receptors or activin intracellular information transmission molecules, with the case where the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof and a test compound are introduced into cells
15 expressing the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, by measuring the cell-stimulating activity via the protein according to claim 17 or a salt thereof, activin receptors or activin
20 intracellular information transmission molecules in both the cases.

24. The method for determining a protein according to claim 16 or the screening method according to any one of claims 18 to 23, which comprises using the two-hybrid method.

25. A kit for screening a compound or a salt thereof

inhibiting or promoting a binding of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof to the protein according to claim 17 or a salt thereof, activin
5 receptors or activin intracellular information transmission molecules, which comprises the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof.

26. A compound or a salt thereof which inhibits or promotes
10 a binding of the protein according to claim 1, the protein according to claim 2, the partial peptide according to claim 6 or a salt thereof to the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which is
15 obtained by the screening method according to any one of claims 18 to 23 or by the screening kit according to claim 25.

27. A pharmaceutical composition comprising the protein according to claim 17, the compound according to claim 26
20 or a salt thereof.

28. The pharmaceutical composition according to claim 27, which is an agent for preventing or treating Alzheimer's disease, Parkinson's disease, epilepsy or Huntington's chorea.

25 29. A method for preventing and treating abnormalities in

nerve cells or cerebral diseases correlated with the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules, which administering an effective amount of the protein according to claim 17, the compound according to claim 26 or a salt thereof to mammals.

30. Use of the protein according to claim 17, the compound according to claim 26 or a salt thereof for preparing an agent for preventing and treating abnormalities in nerve cells or cerebral diseases correlated with the protein according to claim 17 or a salt thereof, activin receptors or activin intracellular information transmission molecules.

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Patent Agents

Fig. 1

TCGCCGCCACGACGGCCAGCACCTCCGAGCGACTGACCGACCTCCACGGCGTCCCGA	60
ACACACTGCCACCGCCGCCGCCGGCGCTCGCGCCGCACTCCCTCGCAGCTCACC	120
ACGTGGCTGCCGCCAACGCCCTCCCGGCCGCTTCCGGCTCTGATGCCCTGAGCGAATCACA	180
GGCGAGCTCCCGGAAGATCCCGCTCTGAGGCTCCGCCCCCGACAGGCCCCGCCACC	240
TCATAGCTCTTTTCCCTCAGCCGCCCTCCTTCCCTTCTCGGCTCAACTAGGTCAGCGCAA	300
GGTGATCCCGGAGAGCGGGGGGGGACCGCTCCTCTTACTTATCGAGCGCGCGC	360
TCCCTCCCGAGCCTCACACCCCTCGCTTCGCCCTTTTTCCTGTCAGGAACTGGTT	420
CCCTCCTTCCCTCTCCACCTGCCCTACCTTCTCCAGAGATCCGACGTGGCGATTAGAGTT	480
CTCAGCGTCACACTGACTTCTAGGCAACTAGCCTAGACTGGAGCTGCCGTGTGTGGGAAC	540
CCCGCGCAGTAGTTGAGCATCAGGCTCTTACCTTGGAGGTGGAGGGGTGAGAAGAAATAG	600
AGGAAGAAGGATAAGTCAGAGGAGGGCCCTGAACAACCTAGCCCCCTCTATTGGCCCTGCTT	660
GGGTGAGCATTCAGTGAGTGTGTTTAAAAAAGGAGGGGAAACAAAGACCTCAG	720
GAGCAGTTTGTGTGTGTCTGGCTTCAAGAAGAAAATTCTAGACATTTATGCCCGC	780
AAGACCAAGCTCAGCTAAGACTACTTCTCCCAAGAAGATAATTGTATCAGAGGATGGGT	840
TGGATCAGTACAGGTGGTTGA GGA GAC GCT GAC AGA GGA CCA TGG AAA GGT GGG	895
Gly Asp Ala Asp Arg Gly Pro Trp Lys Gly Gly	11

Fig. 2

AGA GGA CGC GCG GCT CCT GGG CTT CCT CTG AGC TCA GCT CCA GGC ACC ACA	946
Arg Gly Arg Ala Ala Pro Gly Leu Pro Leu Ser Ser Ala Pro Gly Thr Thr	28
AGG CCA CAT AAG GAG GGT GAG GTC CCT GGA GTG GAC TAC ATT TTC ATA ACC	997
Arg Pro His Lys Glu Gly Glu Val Pro Gly Val Asp Tyr Ile Phe Ile Thr	45
GTT GAG GAG TTT ATG GAA TTG GAG AAA AGT GGT GCT CTC CTA GAA AGC GGG	1048
Val Glu Glu Phe Met Glu Leu Glu Lys Ser Gly Ala Leu Leu Glu Ser Gly	62
ACC TAT GAA GAC AAC TAC TAC GGT ACC CCG AAG CCT CCA GCT GAA CCA GCA	1099
Thr Tyr Glu Asp Asn Tyr Tyr Gly Thr Pro lys Pro Pro Ala Glu Pro Ala	79
CCA TTA TTA AAT GTA ACA GAC CAG ATA CTT CCG GGA GCT ACT CCA AGT GCT	1150
Pro Leu Leu Asn Val Thr Asp Gln Ile Leu Pro Gly Ala Thr Pro Ser Ala	96
GAG GGG AAG CGG AAA AGA AAT AAG TCA GTG ACC AAC ATG GAG AAA GCA AGT	1201
Glu Gly Lys Arg Lys Arg Asn Lys Ser Val Thr Asn Met Glu Lys Ala Ser	113
ATA GAG CCT CCA GAG GAG GAA GAA GAA AGG CCT GTA GTC AAT GGA AAC	1252
Ile Glu Pro Pro Glu Glu Glu Glu Glu Arg Pro Val Val Asn Gly Asn	130
GGC GTG GTC ATA ACC CCA GAA TCC AGT GAA CAT GAA GAC AAA AGT GCA GGT	1303
Gly Val Val Ile Thr Pro Glu Ser Ser Glu His Glu Asp Lys Ser Ala Gly	147

Fig. 3

GCC TCA GGG GAG ACA CCC TCC CAG CCT TAC CCT GCA CCC GTG TAC AGC CAG 1354
 Ala Ser Gly Glu Thr Pro Ser Gln Pro Tyr Pro Ala Pro Val Tyr Ser Gln 164
 CCC GAA GAG CTC AAG GAC CAG ATG GAC GAT ACA AAG CCA ACA AAG CCT GAG 1405
 Pro Glu Glu Leu Lys Asp Gln Met Asp Asp Thr Lys Pro Thr Lys Pro Glu 181
 GAG AAC GAG GAC TCT GAT CCA TTG CCT GAT AAC TGG GAA ATG GCC TAC ACA 1456
 Glu Asn Glu Asp Ser Asp Pro Leu Pro Asp Asn Trp Glu Met Ala Tyr Thr 198
 GAG AAG GGG GAA GTC TAC TTC ATT GAC CAT AAC ACA AAG ACA ACA TCA TGG 1507
 Glu Lys Gly Glu Val Tyr Phe Ile Asp His Asn Thr Lys Thr Thr Ser Trp 215
 CTG GAT CCG CGA CTT GCG AAA AAG GCT AAA CCT CCA GAA GAG TGC AAA GAA 1558
 Leu Asp Pro Arg Leu Ala Lys Lys Ala Lys Pro Pro Glu Glu Cys Lys Glu 232
 AAT GAG CTT CCA TAT GGC TGG GAA AAA ATC GAT GAT CCT ATA TAT GGC ACT 1609
 Asn Glu Leu Pro Tyr Gly Trp Glu Lys Ile Asp Asp Pro Ile Tyr Gly Thr 249
 TAC TAT GTT GAC CAC ATA AAT AGA AGA ACA CAG TTT GAA AAC CCT GTC CTG 1660
 Tyr Tyr Val Asp His Ile Asn Arg Arg Thr Gln Phe Glu Asn Pro Val Leu 266
 GAA GCA AAA AGG AAG CTA CAG CAA CAT AAC ATG CCC CAC ACA GAA CTT GGA 1711
 Glu Ala Lys arg Lys Leu Gln His Asn Met Pro His Thr Glu Leu Gly 283

Fig. 4

GCA AAG CCC CTG CAG GCC CCA GGT TTC CGA GAA AAG CCA CTC TTC ACC CGG	1762
Ala Lys Pro Leu Gln Ala Pro Gly Phe Arg Glu Lys Pro Leu Phe Thr Arg	300
GAT GCA TCC CAG TTG AAG GGA ACG TTC CTC AGC ACC ACC CTC AAA AAG AGC	1813
Asp Ala Ser Gln Leu Lys Gly Thr Phe Leu Ser Thr Thr Leu Lys Lys Ser	317
AAC ATG GGC TTT GGG TTT ACC ATA ATT GGT GGA GAC GAG CCG GAT GAG TTT	1864
Asn Met Gly Phe Gly Phe Thr Ile Ile Gly Gly Asp Glu Pro Asp Glu Phe	334
CTA CAG GTG AAA AGT GTG ATC CCG GAT GGG CCT GCC GCA CAG GAT GGG AAA	1915
Leu Gln Val Lys Ser Val Ile Pro Asp Gly Pro Ala Ala Gln Asp Gly Lys	351
ATG GAG ACA GGT GAT GTC ATT GTC TAT ATT AAT GAA GTT TGT GTC CTT GGA	1966
Met Glu Thr Gly Asp Val Ile Val Tyr Ile Asn Glu Val Cys Val Leu Gly	368
CAC ACT CAT GCA GAT GTT GTC AAA CTT TTC CAG TCT GTT CCT ATT GGT CAG	2017
His Thr His Ala Asp Val Val Lys Leu Phe Gln Ser Val Pro Ile Gly Gln	385
AGT GTC AAC TTG GTG TTG TGT CGT GGC TAC CCT TTT GAC CCT GAA	2068
Ser Val Asn Leu Val Leu Cys Arg Gly Tyr Pro Leu Pro Phe Asp Pro Glu	402
GAT CCT GCT AAC AGC ATG GTG CCA CCC CTT GCA ATA ATG GAG AGG CCA CCT	2119
Asp Pro Ala Asn Ser Met Val Pro Pro Leu Ala Ile Met Glu Arg Pro Pro	419

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Fig. 5

CCG GTG ATG GTC AAT GGA AGA CAT AAC TAT GAA ACA TAC TTG GAA TAC ATT 2170
 Pro Val Met Val Asn Gly Arg His Asn Tyr Glu Thr Tyr Leu Glu Tyr Ile 436
 TCT CGG ACC TCA CAG TCG GTC CCA GAT ATT ACA GAC CGG CCA CCT CAT TCT 2221
 Ser Arg Thr Ser Gln Ser Val Pro Asp Ile Thr Asp Arg Pro Pro His Ser 453
 TTG CAC TCC ATG CCA GCT GAC GGC CAG CTA GAT GGC ACG TAT CCA CCA CCC 2272
 Leu his Ser Met Pro Ala Asp Gly Gln Leu Asp Gly Thr Tyr Pro Pro Pro 470
 GTC CAT GAC GAC AAT GTG TCT ATG GCT TCG TCT GGA GCC ACT CAA GCT GAA 2323
 Val His Asp Asp Asn Val Ser Met Ala Ser Ser Gly Ala Thr Gln Ala Glu 487
 CTT ATG ACC TTA ACC ATT GTG AAA GGT GCC CAG GGA TTT GGC TTT ACT ATT 2374
 Leu Met Thr Leu Thr Ile Val Lys Gly Ala Gln Gly Phe Gly Phe Thr Ile 504
 GCC GAC AGT CCC ACG GGA CAG CGG GTG AAA CAA ATC CTT GAC ATT CAG GGA 2425
 Ala Asp Ser Pro Thr Gly Gln Arg Val Lys Gln Ile Leu Asp Ile Gln Gly 521
 TGC CCT GGG CTG TGT GAA GGA GAC CTC ATT GTT GAG ATC AAC CAA CAG AAT 2476
 Cys Pro Gly Leu Cys Glu Gly Asp Leu Ile Val Glu Ile Asn Gln Gln Asn 538
 GTA CAG AAC CTG AGC CAT ACA GAA GTA GTG GAT ATA CTT AAG GAC TGC CCC 2527
 Val Gln Asn Leu Ser His Thr Glu Val Val Asp Ile Leu Lys Asp Cys Pro 555

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Fig. 6

GTT GGA AGT GAG ACT TCT TTA ATC ATC CAT CGA GGA GGT TTC TTT TCT CCA 2578
 Val Gly Ser Glu Thr Ser Leu Ile Ile His Arg Gly Gly Phe Phe Ser Pro 572
 TGG AAA ACT CCA AAG CCT ATG ATG GAC CGA TGG GAG AAC CAA GGC AGT CCA 2629
 Trp Lys Thr Pro Lys Pro Met Met Asp Arg Trp Glu Asn Gln Gly Ser Pro 589
 CAA ACA AGT TTA TCT GCT CCG GCC GTC CCA CAG AAC CTG CCC TTC CCA CCT 2680
 Gln Thr Ser Leu Ser Ala Pro Ala Val Pro Gln Asn Leu Pro Phe Pro Pro 606
 GCC CTT CAC AGG AGC TCC TTT CCT GAT TCA ACA GAG GCC TTT GAC CCA CGG 2731
 Ala Leu His Arg Ser Ser Phe Pro Asp Ser Thr Glu Ala Phe Asp Pro Arg 623
 AAG CCT GAC CCA TAT GAG CTC TAC GAG AAA TCG AGA GCC ATT TAT GAA AGT 2782
 Lys Pro Asp Pro Tyr Glu Leu Tyr Glu Lys Ser Arg Ala Ile Tyr Glu Ser 640
 AGG CAA CAA GTG CCA CCC AGG ACC AGT TTT CGA ATG GAT TCC TCT GGT CCA 2833
 Arg Gln Gln Val Pro Pro Arg Thr Ser Phe Arg Met Asp Ser Ser Gly Pro 657
 GAT TAT AAG GAA CTG GAT GTT CAC CTT CGG AGG ATG GAG TCT GGA TTT GGC 2884
 Asp Tyr Lys Glu Leu Asp Val His Leu Arg Arg Met Glu Ser Gly Phe Gly 674
 TTT AGA ATC CTT GGG GGA GAT GAA CCT GGA CAG CCT ATT TTG ATC GGA GCC 2935
 Phe Arg Ile Leu Gly Gly Asp Glu Pro Gly Gln Pro Ile Leu Ile Gly Ala 691

Fig. 7

GTC ATT GCC ATG GGC TCA GCT GAC AGA GAC GGC CGT CTA CAC CCA GGA GAT 2986
 Val Ile Ala Met Gly Ser Ala Asp Arg Asp Gly Arg Leu His Pro Gly Asp 708
 GAG CTT GTC TAT GTC GAT GGG ATC CCA GTG GCT GGC AAG ACC CAC CGC TAT 3037
 Glu Leu Val Tyr Val Asp Gly Ile Pro Val Ala Gly Lys Thr His Arg Tyr 725
 GTC ATC GAC CTC ATG CAC CAC GCG GCC CGC AAT GGG CAG GTT AAC CTC ACT 3088
 Val Ile Asp Leu Met His His Ala Ala Arg Asn Gly Gln Val Asn Leu Thr 742
 GTG AGA AGA AAG GTG CTA TGT GGA GGG GAG CCC TGC CCA GAG AAT GGG AGG 3139
 Val Arg Arg Lys Val Leu Cys Gly Gly Glu Pro Cys Pro Glu Asn Gly Arg 759
 AGT CCA GGC TCT GTA TCA ACT CAC CAC AGC TCT CCG CGC AGT GAC TAT GCC 3190
 Ser Pro Gly Ser Val Ser Thr His His Ser Ser Pro Arg Ser Asp Tyr Ala 776
 ACC TAC TCC AAC AGC AAC CAC GCC GCC CCC AGC AGC AAT GCC TCA CCT CCT 3241
 Thr Tyr Ser Asn Ser Asn His Ala Ala Pro Ser Ser Asn Ala Ser Pro Pro 793
 GAA GGC TTT GCC TCA CAC AGC TTG CAG ACC AGT GAT GTG GTC ATT CAC CGC 3292
 Glu Gly Phe Ala Ser His Ser Leu Gln Thr Ser Asp Val Val Ile His Arg 810
 AAA GAA AAC GAA GGG TTT GGC TTC GTC ATC ATC AGC TCT CTG AAC AGG CCT 3343
 Lys Glu Asn Glu Gly Phe Gly Phe Val Ile Ser Ser Leu Asn Arg Pro 827

Fig. 8

GAG TCT GGA GCC ACC ATA ACT GTG CCC CAT AAA ATT GGA CGA ATC ATT GAT 3394
 Glu Ser Gly Ala Thr Ile Thr Val Pro His Lys Ile Gly Arg Ile Ile Asp 844
 GGG AGC CCT GCA GAT CGC TGT GCC AAA CTC AAA GTG GGC GAC CGT ATC TTA 3445
 Gly Ser Pro Ala Asp Arg Cys Ala Lys Leu Lys Val Gly Asp Arg Ile Leu 861
 GCA GTC AAC GGC CAG TCT ATC ATC AAC ATG CCT CAC GCT GAC ATT GTG AAG 3496
 Ala Val Asn Gly Gln Ser Ile Ile Asn Met Pro His Ala Asp Ile Val Lys 878
 CTC ATC AAG GAC GCC GGT CTC AGT GTC ACC CTT CGC ATC ATT CCT CAG GAG 3547
 Leu Ile Lys Asp Ala Gly Leu Ser Val Thr Leu Arg Ile Ile Pro Gln Glu 895
 GAG CTC AAC AGC CCA ACA TCA GCA CCC AGT TCA GAG AAA CAG AGC CCC ATG 3598
 Glu Leu Asn Ser Pro Thr Ser Ala Pro Ser Ser Ser Glu Lys Gln Ser Pro Met 912
 GCC CAG CAG CAC AGC CCT CTG GCC CAG CAG AGT CCT CTG GCC CAG CCA AGC 3649
 Ala Gln Gln His Ser Pro Leu Ala Gln Gln Ser Pro Leu Ala Gln Pro Ser 929
 CCC GCC ACC CCC AAC AGC CCA GTC GCA CAG CCA GCT CCT CCC CAA CCT CTC 3700
 Pro Ala Thr Pro Asn Ser Pro Val Ala Gln Pro Ala Pro Pro Gln Pro Leu 946
 CAG CTG CAA GGA CAC GAA AAT AGT TAC AGG TCA GAA GTT AAA GCG AGG CAA 3751
 Gln Leu Gln Gly His Glu Asn Ser Tyr Arg Ser Glu Val Lys Ala Arg Gln 963

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Fig. 9

GAT GTG AAG CCA GAC ATC CGG CAG CCT CCC TTC ACA GAC TAC AGG CAG CCC 3802
Asp Val Lys Pro Asp Ile Arg Gln Pro Pro Phe Thr Asp Tyr Arg Gln Pro 980
CCG CTG GAC TAC AGG CAG CCC CCG GGA GAC TAC TCA CAG CCC CCA CCC 3853
Pro Leu Asp Tyr Arg Gln Pro Pro Gly Gly Asp Tyr Ser Gln Pro Pro Pro 997
TTG GAC TAC AGG CAG CAC TCT CCA GAC ACC AGG CAG TAC CCT CTG TCA GAC 3904
Leu Asp Tyr Arg Gln His Ser Pro Asp Tyr Arg Gln Tyr Pro Leu Ser Asp 1014
TAC AGG CAG CCA CAG GAT TTT GAT TAT TTC ACT GTG GAC ATG GAG AAA GGA 3955
Tyr Arg Gln Pro Gln Asp Phe Asp Tyr Phe Thr Val Asp Met Glu Lys Gly 1031
GCC AAA GGA TTT GGA TTC AGC ATT CGT GGA GGA AGG GAA TAC AAG ATG GAT 4006
Ala Lys Gly Phe Gly Phe Ser Ile Arg Gly Gly Arg Glu Tyr Lys Met Asp 1048
CTG TAT GTG TTG AGA TTG GCA GAG GAT GGG CCA GCC ATA AGG AAC GGC AGG 4057
Leu Tyr Val Leu Arg Leu Ala Glu Asp Gly Pro Ala Ile Arg Asn Gly Arg 1065
ATG AGG GTA GGA GAT CAG ATC ATT GAA ATA AAT GGG GAA AGC ACA CGA GAC 4108
Met Arg Val Gly Asp Gln Ile Ile Glu Ile Asn Gly Glu Ser Thr Arg Asp 1082
ATG ACC CAC GCC AGA GCA ATA GAA CTC ATC AAG TCT GGA GGA AGA AGA GTG 4159
Met Thr His Ala Arg Ala Ile Glu Leu Ile Lys Ser Gly Gly Arg Arg Val 1099

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Fig. 10

CGG CTG CTG AAG AGA GGC ACG GGG CAG GTC CCG GAG TAT GGA ATG GTA 4210
Arg Leu Leu Leu Lys Arg Gly Thr Gly Gln Val Pro Glu Tyr Gly Met Val 1116
CCT TCC AGC CTC TCC ATG TGC ATG AAA AGT GAC AAG CAT GGG TCC CCA TAT 4261
Pro Ser Ser Leu Ser Met Cys Met Lys Ser Asp Lys His Gly Ser Pro Tyr 1133
TTC TAC TTA CTG GGC CAC CCT AAA GAC ACG ACG AAC CCC ACG CCT GGA GTG 4312
Phe Tyr Leu Leu Gly His Pro Lys Asp Thr Thr Asn Pro Thr Pro Gly Val 1150
CTG CCG CTG CCG CCC CAG GCC TGC CGG AAG TAGGCGTCTCCCTCGAAGACATC 4368
Leu Pro Leu Pro Pro Pro Gln Ala Cys Arg Lys 1161
CTCTCTCCATTCTCTCCATCACATCCAGCCCCACCCCTCCGACCCTTCCCACCATAGGC 4428
CCAGACCCAACTTGGGATATCCAAAGGGAACACGACGTTAGGAAACCAAGGAGCTTTCG 4488
GCCGGGGCCAGAAGACAGCGCCTGGGGAGCAGAGGAGCGCTCGCGGAGCCCGCAG 4548
CGCAGTGGCGGCCACGGCTGGAGGAGGTGCCCGGGCCAGGGCGGCCGAGGCCGGC 4608
AGGCCCGCCTCGGAGCGGCCGACGGGAAGGAGCGCTGCCGGCCGCGGTGAGGGCCTC 4668
GGGGCCGGGGCGCGGAGGCCGAGCCAAGGTGGTGTGCGCTCGGGGCCCCGACCC 4728
GCAGCGGGCCACGGGGGGGGCCAGCGCGCAAGGCGACGATGGCGCCGGGCCCTGG 4788
AAGTGCCGGGCTCCGACAAGCTGCCGGGGGCCCTGCAGCCTGGCGCCTCGGCCGGGGC 4848

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Fig. 11

AGATGAGCCCCAAGCGAGGGCCCCCGCCCTCCACGAGGCCGATCTTCTGGGTT 4908
CCGTCTACGGCGTTTAATTATTTCCACTGTCACACGCATAGATCTATACGAGCGCC 4968
GAAGCCGGGAGCGCGCGTGCGACGCGCGTAGCGGCACGGTGTCGCGCAGG 5028
CAGACCTAAACTGATCCTAAAGCCCCGGTTCCATGGTGGAGCTTTGGCAGCTACGGAA 5088
GAACCAAAATCACGCAAAACATCACAGAGAGACAGTGCAGTGTAGCTTAGATTCAAAAAA 5148
AAAAAAA 5156

Fig. 12

TCGCCGCACGACGCGCCAGCACCTCCGAGCGACTGACCGACCTCCACGCGCGTCCCGA 60
ACACACTGCCACCGCGCCGCCGCGCGGCTCGCGCCGCACTCCCTCGCACGTCACC 120
ACGTGGCTGCCGCCAACGCCCTCCCGCGCGCTTCGGCTCTGATGCCGTAGCGAATCACA 180
GGGAGCTCCCGGAAGATCCCGCTCTGAGGCTCCGCCCCCGGACAGGCCCCGCCACC 240
TCATAGCTCTTTCCCTCAGCCGCCCTCCCTTCCTTCGGCTCAACTAGGTCAGCGCAA 300
GGTGATCCCGAGAGCGGGGCGGGGACCGCTCCTCCTGTTACTTATCGAGCGCGCGC 360
TCCCTCCGAGCCTCACACCCCTCGCTTCGCCCTTTTTCACACTGTCCAGGAAGTGGTT 420
CCCTCCTTCCTCTTCCACCTGCCCTACCTTCTCCAGAGATCCGACGTGGCGATTAGAGTT 480
CTCAGCGTCACACTGACTTCTAGGCAACTAGCCTAGACTGGAGCTGCCGTGTGTGGGAAC 540
CCCGCGCAGTAGTTGAGCATCAGGCTCTTACCTTGGAGGTGGAGGGTGAGAAAGATAG 600
AGGAAGAAGGATAAGTCAGAGGAGGCCCTGAACAACCTAGCCCCCTCTATTGGCCCTGCTT 660
GGTGAGCATTCAGTGAGTGTTTAAAAAAGGAGGAAACAAAGACCTCAG 720
GAGCAGTTTGTGTGTGTCTGGCTTCAAGAAGAAAATTCTAGACATTATGCCCGGC 780
AAGACCAAGCTCAGCTAAGACTACTTCTCCCAAGAAGATAATTGTATCAGAGGATGGGT 840
TGGATCAGTACAGGTGGTTTGAGGAGACGCTGACAGAGGACCATGGAAAGTGGGAGAGG 900
ACGCGGGCTCCTGGGCTTCCTCTGAGCTCAGCTCCAGGCACCAAGGCCACATAAGGA 960

Fig. 13

GGGTGAGTCCCTGGAGTGGA	CTACATTTTCATAACCGTTGAGGAGTTT	ATG GAA TTG GAG	1021
	Met Glu Leu Glu		4
AAA AGT GGT GCT CTC CTA GAA AGC GGG ACC TAT GAA GAC AAC TAC TAC GGT			1072
Lys Ser Gly Ala Leu Leu Glu Ser Gly Thr Tyr Glu Asp Asn Tyr Tyr Gly			21
ACC CCG AAG CCT CCA GCT GAA CCA CCA TTA AAT GTA ACA GAC CAG			1123
Thr Pro lys Pro Pro Ala Glu Pro Ala Pro Leu Leu Asn Val Thr Asp Gln			38
ATA CTT CCG GGA GCT ACT CCA AGT GCT GAG GGG AAG CGG AAA AGA AAT AAG			1174
Ile Leu Pro Gly Ala Thr Pro Ser Ala Glu Gly Lys Arg Lys Arg Asn Lys			55
TCA GTG ACC AAC ATG GAG AAA GCA AGT ATA GAG CCT CCA GAG GAG GAA GAA			1225
Ser Val Thr Asn Met Glu Lys Ala Ser Ile Glu Pro Pro Glu Glu Glu			72
GAA GAA AGG CCT GTA GTC AAT GGA AAC GGC GTG GTC ATA ACC CCA GAA TCC			1276
Glu Glu Arg Pro Val Val Asn Gly Asn Gly Val Val Ile Thr Pro Glu Ser			89
AGT GAA CAT GAA GAC AAA AGT GCA GGT GCC TCA GGG GAG ACA CCC TCC CAG			1327
Ser Glu His Glu Asp Lys Ser Ala Gly Ala Ser Gly Glu Thr Pro Ser Gln			106
CCT TAC CCT GCA CCC GTG TAC AGC CAG CCC GAA GAG CTC AAG GAC CAG ATG			1378
Pro Tyr Pro Ala Pro Val Tyr Ser Gln Pro Glu Glu Leu Lys Asp Gln Met			123

Fig. 14

GAC GAT ACA AAG CCA ACA AAG CCT GAG GAG AAC GAG GAC TCT GAT CCA TTG 1429
 Asp Asp Thr Lys Pro Thr Lys Pro Glu Glu Asn Glu Asp Ser Asp Pro Leu 140
 CCT GAT AAC TGG GAA ATG GCC TAC ACA GAG AAG GGG GAA GTC TAC TTC ATT 1480
 Pro Asp Asn Trp Glu Met Ala Tyr Thr Glu Lys Gly Glu Val Tyr Phe Ile 157
 GAC CAT AAC ACA AAG ACA ACA TCA TGG CTG GAT CCG CGA CTT GCG AAA AAG 1531
 Asp His Asn Thr Lys Thr Thr Ser Trp Leu Asp Pro Arg Leu Ala Lys Lys 174
 GCT AAA CCT CCA GAA GAG TGC AAA GAA AAT GAG CTT CCA TAT GGC TGG GAA 1582
 Ala Lys Pro Pro Glu Glu Cys Lys Glu Asn Glu Leu Pro Tyr Gly Trp Glu 191
 AAA ATC GAT GAT CCT ATA TAT GGC ACT TAC TAT GTT GAC CAC ATA AAT AGA 1633
 Lys Ile Asp Asp Pro Ile Tyr Gly Thr Tyr Tyr Val Asp His Ile Asn Arg 208
 AGA ACA CAG TTT GAA AAC CCT GTC CTG GAA GCA AAA AGG AAG CTA CAG CAA 1684
 Arg Thr Gln Phe Glu Asn Pro Val Leu Glu Ala Lys arg Lys Leu Gln Gln 225
 CAT AAC ATG CCC CAC ACA GAA CTT GGA GCA AAG CCC CTG CAG GCC CCA GGT 1735
 His Asn Met Pro His Thr Glu Leu Gly Ala Lys Pro Leu Gln Ala Pro Gly 242
 TTC CGA GAA AAG CCA CTC TTC ACC CGG GAT GCA TCC CAG TTG AAG GGA ACG 1786
 Phe Arg Glu Lys Pro Leu Phe Thr Arg Asp Ala Ser Gln Leu Lys Gly Thr 259

Fig. 15

TTC CTC AGC ACC ACC CTC AAA AAG AGC AAC ATG GGC TTT GGG TTT ACC ATA 1837
 Phe Leu Ser Thr Thr Leu Lys Lys Ser Asn Met Gly Phe Gly Phe Thr Ile 276
 ATT GGT GGA GAC GAG CCG GAT GAG TTT CTA CAG GTG AAA AGT GTG ATC CCG 1888
 Ile Gly Gly Asp Glu Pro Asp Glu Phe Leu Gln Val Lys Ser Val Ile Pro 293
 GAT GGG CCT GCC GCA CAG CAG GAT GGG AAA ATG GAG ACA GGT GAT GTC ATT GTC 1939
 Asp Gly Pro Ala Ala Gln Asp Gly Lys Met Glu Thr Gly Asp Val Ile Val 310
 TAT ATT AAT GAA GTT TGT GTC CTT GGA CAC ACT CAT GCA GAT GTT GTC AAA 1990
 Tyr Ile Asn Glu Val Cys Val Leu Gly His Thr His Ala Asp Val Val Lys 327
 CTT TTC CAG TCT GTT CCT ATT GGT CAG AGT GTC AAC TTG GTG TTG TGT CGT 2041
 Leu Phe Gln Ser Val Pro Ile Gly Gln Ser Val Asn Leu Val Leu Cys Arg 344
 GGC TAC CCT TTG CCC TTT GAC CCT GAA GAT CCT GCT AAC AGC ATG GTG CCA 2092
 Gly Tyr Pro Leu Pro Phe Asp Pro Glu Asp Pro Ala Asn Ser Met Val Pro 361
 CCC CTT GCA ATA ATG GAG AGG CCA CCT CCG GTG ATG GTC AAT GGA AGA CAT 2143
 Pro Leu Ala Ile Met Glu Arg Pro Pro Val Met Val Asn Gly Arg His 378
 AAC TAT GAA ACA TAC TTG GAA TAC ATT TCT CGG ACC TCA CAG TCG GTC CCA 2194
 Asn Tyr Glu Thr Tyr Leu Glu Tyr Ile Ser Arg Thr Ser Gln Ser Val Pro 395

Fig. 16

GAT ATT ACA GAC CGG CCA CCT CAT TCT TTG CAC TCC ATG CCA GCT GAC GGC 2245
 Asp Ile Thr Asp Arg Pro Pro His Ser Leu his Ser Met Pro Ala Asp Gly 412
 CAG CTA GAT GGC ACG TAT CCA CCA CCC GTC CAT GAC GAC AAT GTG TCT ATG 2296
 Gln Leu Asp Gly Thr Tyr Pro Pro Pro Val His Asp Asp Asn Val Ser Met 429
 GCT TCG TCT GGA GCC ACT CAA GCT GAA CTT ATG ACC TTA ACC ATT GTG AAA 2347
 Ala Ser Ser Gly Ala Thr Gln Ala Glu Leu Met Thr Leu Thr Ile Val Lys 446
 GGT GCC CAG GGA TTT GGC TTT ACT ATT GCC GAC AGT CCC ACG GGA CAG CGG 2398
 Gly Ala Gln Gly Phe Gly Phe Thr Ile Ala Asp Ser Pro Thr Gly Gln Arg 463
 GTG AAA CAA ATC CTT GAC ATT CAG GGA TGC CCT GGG CTG TGT GAA GGA GAC 2449
 Val Lys Gln Ile Leu Asp Ile Gln Gly Cys Pro Gly Leu Cys Glu Gly Asp 480
 CTC ATT GTT GAG ATC AAC CAA CAG AAT GTA CAG AAC CTG AGC CAT ACA GAA 2500
 Leu Ile Val Glu Ile Asn Gln Gln Asn Val Gln Asn Leu Ser His Thr Glu 497
 GTA GTG GAT ATA CTT AAG GAC TGC CCC GTT GGA AGT GAG ACT TCT TTA ATC 2551
 Val Val Asp Ile Leu Lys Asp Cys Pro Val Gly Ser Glu Thr Ser Leu Ile 514
 ATC CAT CGA GGA GGT TTC TTT TCT CCA TGG AAA ACT CCA AAG CCT ATG ATG 2602
 Ile His Arg Gly Gly Phe Phe Ser Pro Trp Lys Thr Pro Lys Pro Met Met 531

Fig. 17

GAC CGA TGG GAG AAC CAA GGC AGT CCA CAA ACA AGT TTA TCT GCT CCG GCC 2653
 Asp Arg Trp Glu Asn Gln Gly Ser Pro Gln Thr Ser Leu Ser Ala Pro Ala 548
 GTC CCA CAG AAC CTG CCC TTC CCA CCT GCC CTT CAC AGG AGC TCC TTT CCT 2704
 Val Pro Gln Asn Leu Pro Phe Pro Pro Ala Leu His Arg Ser Ser Phe Pro 565
 GAT TCA ACA GAG GCC TTT GAC CCA CGG AAG CCT GAC CCA TAT GAG CTC TAC 2755
 Asp Ser Thr Glu Ala Phe Asp Pro Arg Lys Pro Asp Pro Tyr Glu Leu Tyr 582
 GAG AAA TCG AGA GCC ATT TAT GAA AGT AGG CAA CAA GTG CCA CCC AGG ACC 2806
 Glu Lys Ser Arg Ala Ile Tyr Glu Ser Arg Gln Gln Val Pro Pro Arg Thr 599
 AGT TTT CGA ATG GAT TCC TCT GGT CCA GAT TAT AAG GAA CTG GAT GTT CAC 2857
 Ser Phe Arg Met Asp Ser Ser Gly Pro Asp Tyr Lys Glu Leu Asp Val His 616
 CTT CGG AGG ATG GAG TCT GGA TTT GGC TTT AGA ATC CTT GGG GGA GAT GAA 2908
 Leu Arg Arg Met Glu Ser Gly Phe Gly Phe Arg Ile Leu Gly Gly Asp Glu 633
 CCT GGA CAG CCT ATT TTG ATC GGA GCC GTC ATT GCC ATG GGC TCA GCT GAC 2959
 Pro Gly Gln Pro Ile Leu Ile Gly Ala Val Ile Ala Met Gly Ser Ala Asp 650
 AGA GAC GGC CGT CTA CAC CCA GAT GAG CTT GTC TAT GTC GAT GGG ATC 3010
 Arg Asp Gly Arg Leu His Pro Gly Asp Glu Leu Val Tyr Val Asp Gly Ile 667

Fig. 18

CCA GTG GCT GGC AAG ACC CAC CGC TAT GTC ATC GAC CTC ATG CAC CAC GCG 3061
 Pro Val Ala Gly Lys Thr His Arg Tyr Val Ile Asp Leu Met His His Ala 684
 GCC CGC AAT GGG CAG GTT AAC CTC ACT GTG AGA AGA AAG GTG CTA TGT GGA 3112
 Ala Arg Asn Gly Gln Val Asn Leu Thr Val Arg Arg Lys Val Leu Cys Gly 701
 GGG GAG CCC TGC CCA GAG AAT GGG AGG AGT CCA GGC TCT GTA TCA ACT CAC 3163
 Gly Glu Pro Cys Pro Glu Asn Gly Arg Ser Pro Gly Ser Val Ser Thr His 718
 CAC AGC TCT CCG CGC AGT GAC TAT GCC ACC TAC TCC AAC AGC AAC CAC GCC 3214
 His Ser Ser Pro Arg Ser Asp Tyr Ala Thr Tyr Ser Asn Ser Asn His Ala 735
 GCC CCC AGC AGC AAT GCC TCA CCT CCT GAA GGC TTT GCC TCA CAC AGC TTG 3265
 Ala Pro Ser Ser Asn Ala Ser Pro Pro Glu Gly Phe Ala Ser His Ser Leu 752
 CAG ACC AGT GAT GTG GTC ATT CAC CGC AAA GAA AAC GAA GGG TTT GGC TTC 3316
 Gln Thr Ser Asp Val Val Ile His Arg Lys Glu Asn Glu Gly Phe Gly Phe 769
 GTC ATC ATC AGC TCT CTG AAC AGG CCT GAG TCT GGA GCC ACC ATA ACT GTG 3367
 Val Ile Ile Ser Ser Leu Asn Arg Pro Glu Ser Gly Ala Thr Ile Thr Val 786
 CCC CAT AAA ATT GGA CGA ATC ATT GAT GGG AGC CCT GCA GAT CGC TGT GCC 3418
 Pro His Lys Ile Gly Arg Ile Ile Asp Gly Ser Pro Ala Asp Arg Cys Ala 803

Fig. 19

AAA CTC AAA GTG GGC GAC CGT ATC TTA GCA GTC AAC GGC CAG TCT ATC ATC 3469
 Lys Leu Lys Val Gly Asp Arg Ile Leu Ala Val Asn Gly Gln Ser Ile Ile 820
 AAC ATG CCT CAC GCT GAC ATT GTG AAG CTC ATC AAG GAC GCC GGT CTC AGT 3520
 Asn Met Pro His Ala Asp Ile Val Lys Leu Ile Lys Asp Ala Gly Leu Ser 837
 GTC ACC CTT CGC ATC ATT CCT CAG GAG GAG CTC AAC AGC CCA ACA TCA GCA 3571
 Val Thr Leu Arg Ile Ile Pro Gln Glu Glu Leu Asn Ser Pro Thr Ser Ala 854
 CCC AGT TCA GAG AAA CAG AGC CCC ATG GCC CAG CAG CAC AGC CCT CTG GCC 3622
 Pro Ser Ser Glu Lys Gln Ser Pro Met Ala Gln Gln His Ser Pro Leu Ala 871
 CAG CAG AGT CCT CTG GCC CAG CCA AGC CCC GCC ACC CCC AAC AGC CCA GTC 3673
 Gln Gln Ser Pro Leu Ala Gln Pro Ser Pro Ala Thr Pro Asn Ser Pro Val 888
 GCA CAG CCA GCT CCT CCC CAA CCT CTC CAG CTG CAA GGA CAC GAA AAT AGT 3724
 Ala Gln Pro Ala Pro Pro Gln Pro Leu Gln Leu Gln Gly His Glu Asn Ser 905
 TAC AGG TCA GAA GTT AAA GCG AGG CAA GAT GTG AAG CCA GAC ATC CGG CAG 3775
 Tyr Arg Ser Glu Val Lys Ala Arg Gln Asp Val Lys Pro Asp Ile Arg Gln 922
 CCT CCC TTC ACA GAC TAC AGG CAG CCC CCG CTG GAC TAC AGG CAG CCC CCG 3826
 Pro Pro Phe Thr Asp Tyr Arg Gln Pro Pro Leu Asp Tyr Arg Gln Pro Pro 939

Fig. 20

GGA GGA GAC TAC TCA CAG CCC CCA CCC TTG GAC TAC AGG CAG CAC TCT CCA 3877
 Gly Gly Asp Tyr Ser Gln Pro Pro Pro Leu Asp Tyr Arg Gln His Ser Pro 956
 GAC ACC AGG CAG TAC CCT CTG TCA GAC TAC AGG CAG CCA CAG GAT TTT GAT 3928
 Asp Tyr Arg Gln Tyr Pro Leu Ser Asp Tyr Arg Gln Pro Gln Asp Phe Asp 973
 TAT TTC ACT GTG GAC ATG GAG AAA GGA GCC AAA GGA TTT GGA TTC AGC ATT 3979
 Tyr Phe Thr Val Asp Met Glu Lys Gly Ala Lys Gly Phe Gly Phe Ser Ile 990
 CGT GGA GGA AGG GAA TAC AAG ATG GAT CTG TAT GTG TTG AGA TTG GCA GAG 4030
 Arg Gly Gly Arg Glu Tyr Lys Met Asp Leu Tyr Val Leu Arg Leu Ala Glu 1007
 GAT GGG CCA GCC ATA AGG AAC GGC AGG ATG AGG GTA GGA GAT CAG ATC ATT 4081
 Asp Gly Pro Ala Ile Arg Asn Gly Arg Met Arg Val Gly Asp Gln Ile Ile 1024
 GAA ATA AAT GGG GAA AGC ACA CGA GAC ATG ACC CAC GCC AGA GCA ATA GAA 4132
 Glu Ile Asn Gly Glu Ser Thr Arg Asp Met Thr His Ala Arg Ala Ile Glu 1041
 CTC ATC AAG TCT GGA GGA AGA AGA GTG CGG CTG CTG AAG AGA GGC ACG 4183
 Leu Ile Lys Ser Gly Gly Arg Arg Val Arg Leu Leu Lys Arg Gly Thr 1058
 GGG CAG GTC CCG GAG TAT GGA ATG GTA CCT TCC AGC CTC TCC ATG TGC ATG 4234
 Gly Gln Val Pro Glu Tyr Gly Met Val Pro Ser Ser Leu Ser Met Cys Met 1075

Fig. 21

AAA AGT GAC AAG CAT GGG TCC CCA TAT TTC TAC TTA CTG GGC CAC CCT AAA 4285
 Lys Ser Asp Lys His Gly Ser Pro Tyr Phe Tyr Leu Leu Gly His Pro Lys 1092
 GAC ACG ACG AAC CCC ACG CCT GGA GTG CTG CCG CTG CCG CCC CAG GCC 4336
 Asp Thr Thr Asn Pro Thr Pro Gly Val Leu Pro Pro Pro Gln Ala 1109
 TGC CGG AAG TAGGCGTCTCCCTCGAAGACATCCTCTCTCCATTCTCTCCATCAGCCCC 4400
 Cys Arg Lys 1112
 ACCCTCCGACCCCTTCCCACCAGATAGGCCCCAGACCCAACTTGGGATATCCAAAGGGAACA 4460
 CGACGTTAGGAACCAAGGAGCTTTCGGCCCGCGCCAGAGAAGCAGCGCCTGGGGGA 4520
 GCAGAGGAGCGCTCGGCGAGCCCCGACGCGAGTGC CGGCCCCAGGCTGGAGGAGGTGCC 4580
 CGCGGCCAGGGCGCGGAGGCCCGCAGGCCCGCTCGGAGCGGCCGACGGGAAGGA 4640
 GCGGCTGCGGCGCGGTGAGGGCCTCGGGGCCCGCGCGCGGAGGCCGAGGCCAA 4700
 GGTGGGTGTCCGCTCGGGGCCCCGACCCGACGCGCGGCCACGGGGCGGCCAGCGCG 4760
 CAAGGCGACGATGGCGCCCGGGCCCTGGAAGGTGCCGGCTCCGACAAAGCTGCCGGCGC 4820
 CCTGCAGCCTGGCGCCTCGGGCCCGGGCAGATGAGCCCCAAGCGAGGGCCCCGCCCG 4880
 CCTCCACGAGCCGATCTTCTGGGTTCGGTCTCAGGCGTTTAATTATTCCACTG 4940
 TCACACGCATAGATCTATACGAGCGCCGGAAGCCCGGAGCGCGCGGTGCGACGCGCGT 5000

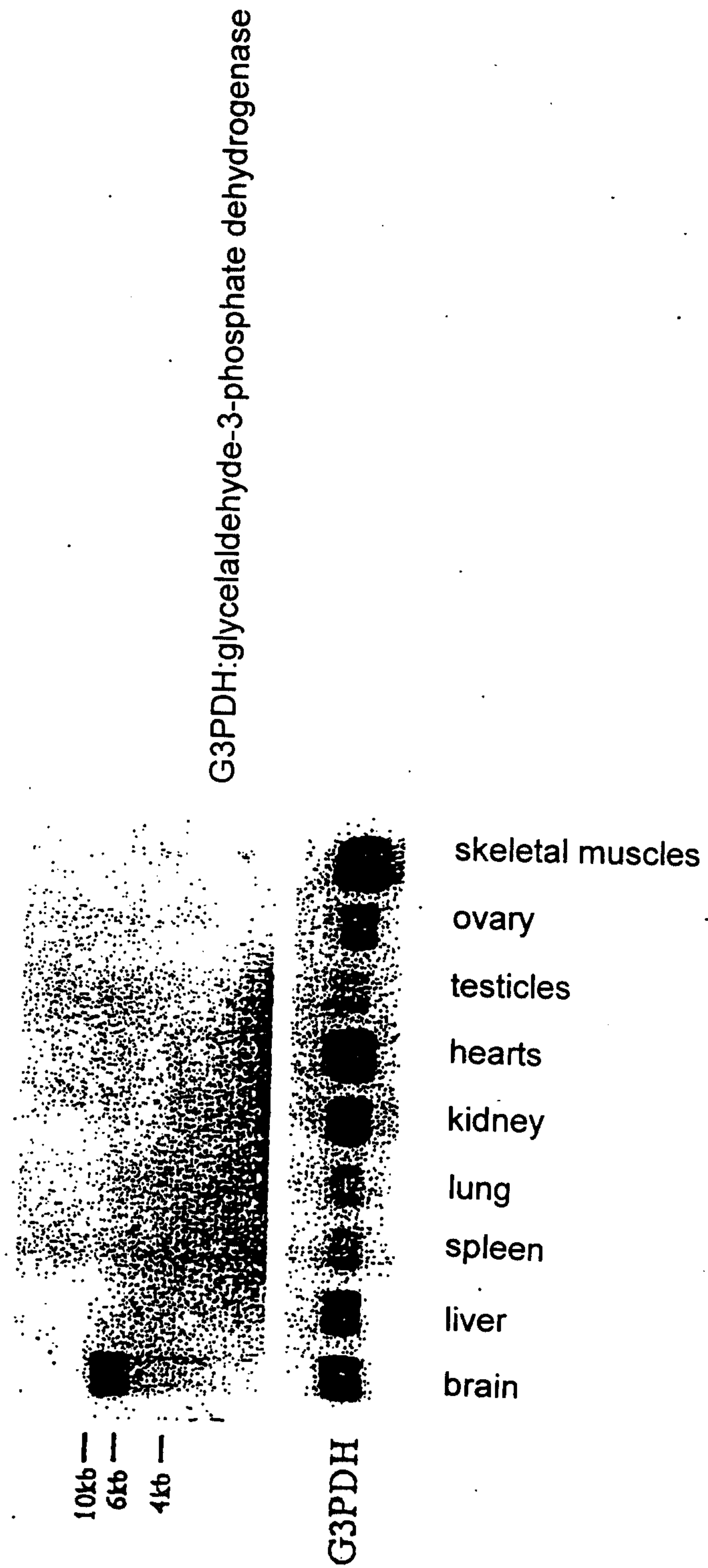
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Fig. 22

AGGCGGCACGGC	5060
CGGTGTGCGCGCAGGCAGACCTAAACTGATCCTAAAGCCCCGGTTC	
CATGGTGGAGCTTTGGCAGCTACGGAAGAACCAAAATCACGCAACATCACAGAGAGAC	5120
AGTGCAGTGTAGCTTTAGATTCAAAAAA	5156

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Fig. 23



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Fig. 24

