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(54) **METHOD FOR THE PRODUCTION OF HUMAN THROMBIN AND USES THEREOF**

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USPC 530/350; 435/6.1

See application file for complete search history.

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(57) **ABSTRACT**

The present invention disclosed the cloning and the purification of a mutant of prethrombin-2, which contains a thrombin cleavage site instead of a factor Xa cleavage site. The stable mutant prethrombin-2 is able to convert itself autocatalytically into active α -thrombin in the absence of ecarin or factor Xa. The new concept of signal amplification using self-replicating enzymes can be applied to improve sensitivity of α -thrombin assays and also for the preparation of different enzymes.

14 Claims, 4 Drawing Sheets

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Figure 1

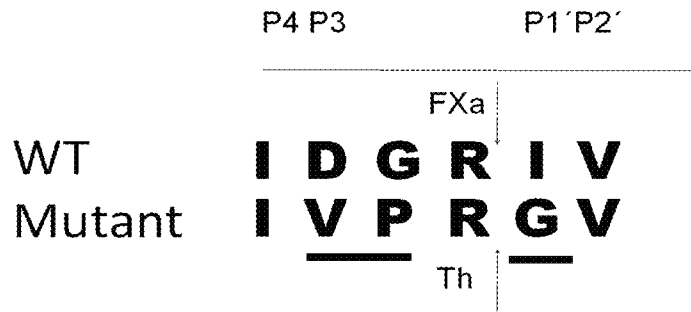


Figure 2

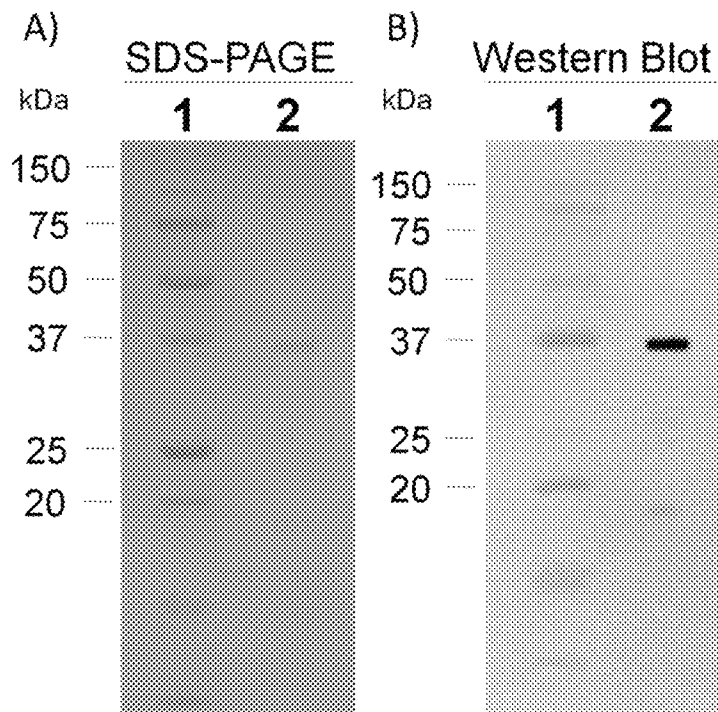


Figure 3

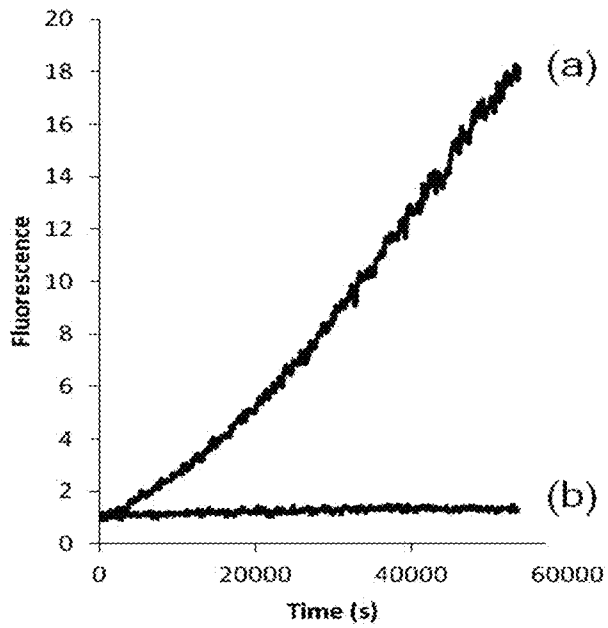


Figure 4

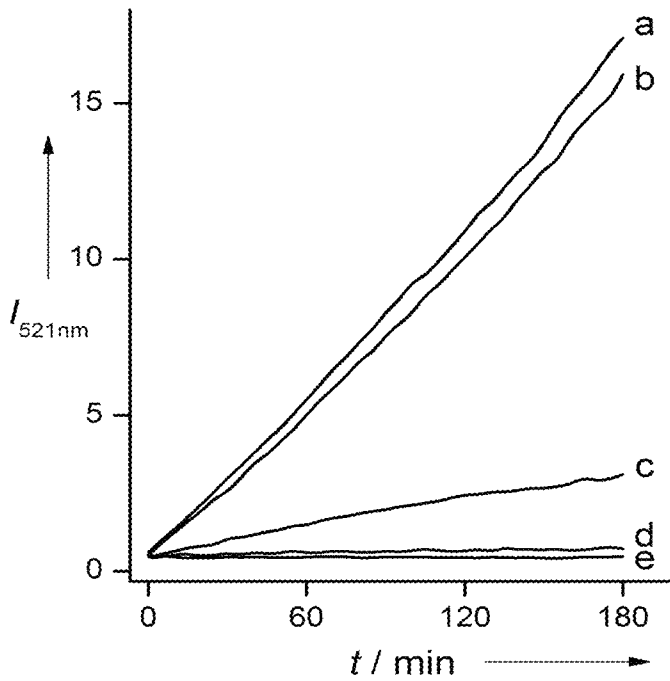


Figure 5

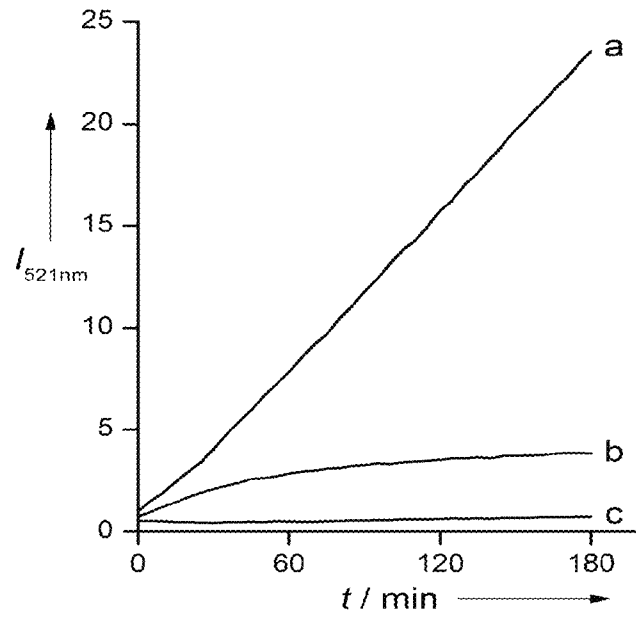


Figure 6

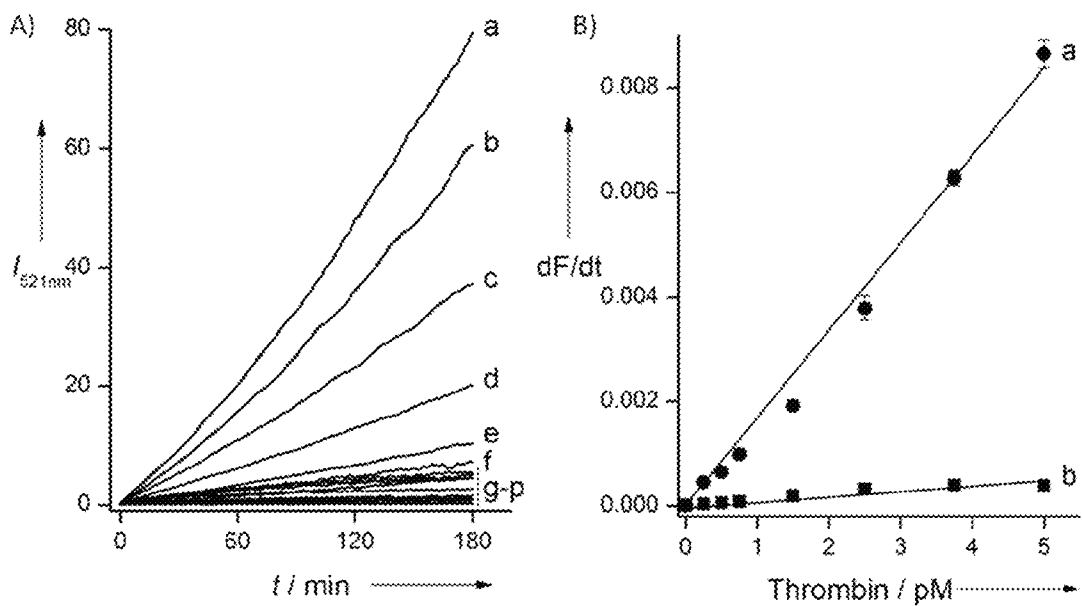
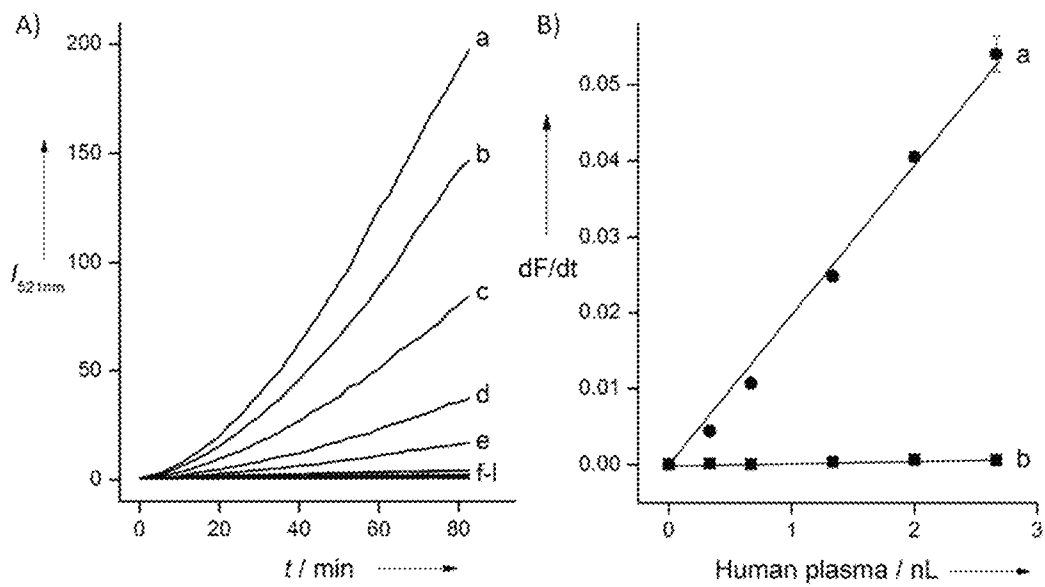


Figure 7



METHOD FOR THE PRODUCTION OF HUMAN THROMBIN AND USES THEREOF

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is filed under the provisions of 35 U.S.C. §371 and claims the priority of International Patent Application No. PCT/EP2012/071572 filed on 31 Oct. 2012 entitled "METHOD FOR THE PRODUCTION OF HUMAN THROMBIN AND USES THEREOF" in the name of Valery PAVLOV, et al, which claims priority to U.S. Provisional Patent Application No. 61/555,763 filed on Nov. 4, 2011 both of which are hereby incorporated by reference herein in their entirety.

FIELD OF THE INVENTION

The invention refers to the field of biotechnology. Specifically, the invention relates to a stable recombinant mutated prethrombin-2 which is able to convert itself auto-catalytically into active α -thrombin that can be used as a haemostatic. Another aspect of the invention includes the application of mutated recombinant prethrombin-2 and the α -thrombin obtained by itself-autocatalytic activity, in diagnostics and/or prognostics of coagulation related-diseases.

BACKGROUND OF THE INVENTION

α -Thrombin plays an important role in hemostasis and hence is a highly useful protein for use as a hemostatic. Currently, α -thrombin is utilized in more than 1 000 000 patients in the USA each year [1]. The main source of α -thrombin is pooled human plasma collected from donors. The plasma is treated by complicated filtration and separation, however, no procedure is completely effective against viral particles derived from human blood [2].

Alternatively, recombinant α -thrombin can be produced that is devoid of the risks of viral particles. Thrombin is synthesized in the form of prothrombin which is one of proteolytic enzymes which are normally synthesized in their inactive form, known as proenzymes or zymogens, which can be cleaved by factor Xa or by ecarin, a protease found in snake poison, [3] at two places, Arg-Thr and Arg-Ile bonds. Cleavage at the first factor Xa site results in prethrombin-2, an inactive single-chain precursor that has the same size as α -thrombin. Activation of prethrombin-2 to α -thrombin occurs through internal rearrangement of the initial peptide chain upon cleavage of Arg-Ile bond with factor Xa or by ecarin [4]. The cleaved shorter peptide section does not leave the α -thrombin macromolecule but stays linked with the longer peptide sequence via an S—S bond.

Currently, it is impossible to directly express active recombinant α -thrombin from the gene fragment corresponding to prothrombin because the resulting protein will be always inactive prethrombin-2. The preparation of recombinant prethrombin-2 and its activation to α -thrombin by ecarin, has been described in the literature [5]. The main drawback of this procedure is the necessity to employ extremely dangerous ecarin, the primary reagent in the venom of the snake *Echis carinatus* [6] to activate α -thrombin. The removal of ecarin, significantly diminishes the yield and increases costs of α -thrombin preparation. Still there is no 100% assurance that recombinant α -thrombin is not contaminated with ecarin. Factor Xa can be used instead of lethal ecarin but it requires

for its optimal operation a complex with factor V, platelet phospholipids and calcium [7], that also must be separated from α -thrombin.

On the other hand, enzymes, such as, α -thrombin, have been commonly used in different bioanalytical assays for detection and amplification of signal. They are employed in quantification of glucose [8], H₂O₂ [9], pesticides [10], cholesterol [11], and ethanol [12], and are the basis for ELISAs (enzyme linked immunosorbent assays) [13]. Enzyme amplification techniques [14] have been used to improve the sensitivity of several bioanalytical assays. In this sense, the state of the art discloses an assay including a double amplification cascade in which ecarin, converted prothrombin to α -thrombin to digest an artificial fluorogenic substrate [15]. As mentioned above, prothrombin is one of proteolytic enzymes which are normally synthesized in their inactive form, known as proenzymes or zymogens [16]. When the product of the proenzyme cleavage reaction catalyzes the same reaction, the process is called an autocatalytic activation [17]. Some examples of natural autocatalytic enzymes are trypsinogen, pepsinogen, or the blood coagulation factor XII [18]. The autocatalytic behavior of these enzymes could be applied for analytical purposes. Moreover, tests for α -thrombin activity are used to evaluate the rate of blood coagulation, therefore it is important to develop sensitive methods to monitor its activity.

The signal obtained by a low concentration of an enzyme can be considerably amplified by means of an autocatalytic reaction. The use of zymogens in bioanalysis is limited since all known natural autocatalytic proenzymes are unstable in vitro and their preparations always contain traces of corresponding active enzymes [19]. To the best of our knowledge, there are still no commercially available kits based on natural autocatalytic proenzymes. Previously, the inventors used the self-replicating DNA machines based on endonucleases to create signal amplification networks [20, 21]. Unfortunately, DNA machines are not stable in body liquids and can not be applied for the analysis of blood derived samples.

In spite of considerable research into the signal amplification in bioassays, specifically, using self-replicating enzymes and also the preparation and production of these enzymes, more studies are still needed. Moreover, it has not yet been found a proenzyme stable in vitro without traces of corresponding active enzymes and also with autocatalytic activity, particularly these is not disclosures in the prior art about a recombinant prethrombin-2, with autocatalytic features, which show stability in vitro, without traces of α -thrombin.

DESCRIPTION OF THE INVENTION

Brief Description of the Invention

The present invention, overcomes these problems in the art by disclosing a new methodology for production of a mouse (SEQ ID NO: 1) and human (SEQ ID NO: 2) mutant recombinant stable prethrombin-2 which are able to convert itself auto-catalytically into active α -thrombin in the absence of ecarin or factor Xa or any other substance.

The method for obtaining a stable mutant recombinant prethrombin-2 consists of changing the cleavage site of a recombinant proenzyme by site directed mutagenesis to obtain a stable protein that can be cleaved by the corresponding active enzyme. Specifically, the inventors have modified the cleavage site of native or wild type (WT) mouse (SEQ ID NO: 3) and human (SEQ ID NO: 4) prethrombin-2 to a α -thrombin cleavage site to obtain a stable artificial autocatalytic enzyme. The mutant recombinant proenzyme is acti-

vated by α -thrombin triggering the autocatalytic reaction. The concept of signal amplification using self-replicating enzymes can be applied to improve sensitivity of α -thrombin assays and also for the preparation of different enzymes. Therefore, the present invention displays an autocatalytic mutant recombinant enzyme, prethrombin-2, and its application for the signal amplification in bioassays for the detection of α -thrombin/prothrombin and for the synthesis of recombinant α -thrombin.

Surprisingly, the mutated prethrombin-2 disclosed in the present invention is completely stable, without any trace of thrombin, or any other proteases or any undesired factor, component, contaminant, etc, which may initiate an undesired and/or uncontrolled cleavage of the mutant prethrombin-2. The invention described herein, allows total control of the autocatalysis of prethrombin-2 due to the unexpected stability shown by the mutated prethrombin-2 of the invention. This stability feature of mutated prethrombin-2 of present invention has as advantage its use in in vitro assays for detection of α -thrombin/prothrombin, with more accuracy as far as the stable mutated prethrombin-2 of the invention has not autocatalytic basal activity due to contaminant, as the ones previously mentioned, and its autocatalysis is only triggered by the amount of α -thrombin present in the assay sample.

Thus, in a first aspect the present invention refers to a stable and autocatalytic mutant prethrombin-2, from murine or human origins (SEQ ID NO: 1 or SEQ ID NO: 2, respectively) and their corresponding DNA encoding sequences (SEQ ID NO: 5 or SEQ ID NO: 6, respectively).

In a second aspect, the invention refers to the use of at least one of the mutated prethrombin-2 proenzymes or their nucleotide encoding sequences as mentioned above for the synthesis of α -thrombin.

In a third aspect, the present invention refers to the use of at least one of the mutated prethrombin-2 proenzymes for the manufacture of a kit for the in vitro diagnosis and/or prognosis of coagulation-related diseases.

In another aspect, the present invention refers to the use of at least one of the mutated prethrombin-2 proenzymes for the manufacture of a kit for detection of α -thrombin/prothrombin.

In another aspect, the invention refers to a kit which comprises at least one of the mutated prethrombin-2 proenzymes disclosed in the present invention. The kit disclosed herein also could include other reagents, i.e.: assay reagents, buffers and sterile saline or another pharmaceutically acceptable emulsion and suspension base. In addition, the kits may include instructional materials containing directions (e.g., protocols) for the practice of the assay methods of this invention.

In another aspect, the present invention refers to the use of the kit disclosed herein for the in vitro diagnosis and/or prognosis of coagulation-related diseases. The coagulation-related disease is selected from the group comprising: haemophilia, thrombosis, inherited prothrombin deficiency Type I or Type II, acquired prothrombin deficiency and von Willbrand disease.

In another aspect, the present invention refers to a method for preparing active α -thrombin using the mutant recombinant prethrombin-2 disclosed in the present invention.

In still another aspect, the invention refers to an in vitro method of diagnosis and/or prognosis of coagulation-related diseases which comprises determining in a sample from a subject, the expression or activity level of α -thrombin/prothrombin using the kit disclosed herein and the comparison of said expression level with respect to the expression values obtained from healthy controls. The α -thrombin/prothrombin

expression or activity levels are analyzed in a sample of blood. For this purpose is extremely important that the prethrombin-2 to be used in the amplification reaction wherein the in vitro assay is based was stable, because traces of thrombin or any other proteases contaminating the prethrombin-2 to be used in thrombin detection in biological samples would render artefacts.

Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this invention belongs. However, for ease of reference, some of these terms will now be defined.

The expression or activity level of a disease-associated α -thrombin/prothrombin is information in a number of ways. For example, a differential expression of a disease-associated α -thrombin/prothrombin compared to a control may be used as a diagnostic that a patient suffers from the disease. Expression or activity levels of a disease-associated α -thrombin/prothrombin may also be used to monitor the treatment and disease state of a patient. Furthermore, expression or activity levels of the disease-associated α -thrombin/prothrombin may allow the screening of drug candidates for altering a particular expression profile or suppressing an expression profile associated with the disease. In the present invention, the terms "expression or activity" in relation with α -thrombin/prothrombin levels are used interchangeably.

The term "sample" refers preferably to a sample of a fluid such as a solution and more preferably refers to a sample of a body fluid. Samples of body fluids can be obtained by well known techniques and include, preferably, samples of blood, plasma, serum, more preferably, samples of plasma.

The term "subject" as used herein refers to an animal, preferably a mammal, most preferably a human being, including both young and old human beings of both sexes which may suffer from or are predisposed to a pathology. The subject according to this aspect of the present invention may suffer from a pathology associated with abnormal expression or activity of coagulation pathway. The terms "subject" and "patient" could be used interchangeably throughout the present invention.

The term "stable prethrombin-2" means, within present invention, a prethrombin-2 in a sample free of thrombin or any other proteases or any undesired factor, component, contaminant, etc, which may trigger autocatalytic activity of prethrombin-2 itself.

FIGURE LEGENDS

FIG. 1. Sequence alignment of the cleavage site of wild type (WT) and mutant prethrombin-2. FXa cleavage site present in WT prethrombin-2, corresponding to residues IDGRIV, was changed to IVPRGV which corresponds to a α -thrombin (Th) cleavage site. Changed amino acids are underlined.

FIG. 2. Analysis of the purified mouse mutant prethrombin-2. A) 12% SDS-PAGE stained with Coomassie Brilliant Blue. B) Nitrocellulose membrane probed with anti-His-tag mouse monoclonal antibodies. Lane 1, molecular mass references; lane 2, 0.3 μ g mutant prethrombin-2.

FIG. 3. Stability assay of human mutant prethrombin-2. Evolution of fluorescence intensity in samples containing human wild type recombinant prethrombin-2 (1.6×10^{-6} M) (a) or human mutant recombinant prethrombin-2 (1.6×10^{-6} M) (b), with the fluorogenic substrate rhodamine 110, bis-(p-tosyl-L-glycyl-L-prolyl-L-arginine amide) (4×10^{-6} M) (In-vitrogen) in 50 mM Tris-HCl, pH 9.0, 150 mM NaCl.

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FIG. 4. Evolution of fluorescence intensity in samples containing a) 2.2×10^{-12} M human wild type α -thrombin and 2.1×10^{-7} M human mutant prethrombin-2; b) 2.2×10^{-12} M human wild type α -thrombin and 2.1×10^{-7} M mouse mutant prethrombin-2; c) 2.2×10^{-12} M human wild type α -thrombin; d) 2.1×10^{-7} M human mutant prethrombin-2; e) 2.1×10^{-7} M mouse mutant prethrombin-2.

FIG. 5. Time courses of mutant prethrombin-2 activation and self-replication. a) 5.38×10^{-9} M cleaved mutant prethrombin-2 and 2.3×10^{-7} M untreated mutant prethrombin-2; b) 5.38×10^{-9} M cleaved mutant prethrombin-2; c) 2.3×10^{-7} M untreated mutant prethrombin-2.

FIG. 6. A) Evolution of fluorescence intensity in the presence (curves a-h) of 5.3×10^{-7} M mutant prethrombin-2 or its absence (curves i-p). The samples contained different concentrations of human α -thrombin: a) and i) 5×10^{-12} M; b) and j) 3.75×10^{-12} M; c) and k) 2.5×10^{-12} M; d) and l) 1.5×10^{-12} M; e) and m) 7.5×10^{-13} M; f) and n) 5×10^{-13} M; g) and o) 2.5×10^{-13} M; h) and p) 0 M. B) Calibration curve of α -thrombin in the presence (curve a) and absence (curve b) of mutant prethrombin-2.

FIG. 7. A) Evolution of fluorescence intensity in human plasma samples in the presence of 1.5×10^{-6} M mutant prethrombin-2 (curves a-f) or its absence (curves g-l). The samples contained 6×10^{-8} M ecarin and different volumes of human plasma: a) and g) 2.6 nL; b) and h) 2 nL; c) and i) 1.3 nL; d) and j) 0.67 nL; e) and k) 0.33 nL; f) and l) 0 nL. B) Calibration curve of human plasma in the presence (curve a) or absence (curve b) of mutant prethrombin-2.

DETAILED DESCRIPTION OF THE INVENTION

Present invention relates to a mutant prethrombin-2 characterized by having SEQ ID NO: 1 or SEQ ID NO: 2 or their nucleotide encoding sequences characterized by SEQ ID NO: 5 or SEQ ID NO: 6, respectively.

In a particular embodiment of the invention, the mutant prethrombin-2 is SEQ ID NO: 2 or their nucleotide encoding sequence characterized by SEQ ID NO: 6.

In another particular embodiment of the invention, the mutant prethrombin-2 characterized by having SEQ ID NO: 2 or by their nucleotide encoding sequence: SEQ ID NO: 6 which is from human origin.

In another particular embodiment of the invention, the mutant prethrombin-2 disclosed in the present invention is characterized by having autocatalytic activity and by being stable in vitro.

Present invention also relates to the mutated prethrombin-2 mentioned above for use in the synthesis of α -thrombin or in the in vitro detection of α -thrombin/prothrombin in biological samples.

Present invention also discloses the use of the mutated prethrombin-2 mentioned above for the manufacture of a kit for the in vitro diagnosis and/or prognosis of coagulation-related diseases.

In a particular embodiment of the kit of the present invention, the coagulation-related disease is selected from the group comprising: haemophilia, thrombosis, inherited prothrombin deficiency Type I or Type II, acquired prothrombin deficiency and von Willebrand disease.

Present invention also discloses a mutant prethrombin-2 mentioned above for use in the in vitro diagnosis and/or prognosis of coagulation-related diseases. In a particular embodiment, the coagulation-related disease is selected from the group comprising: haemophilia, thrombosis, inherited prothrombin deficiency Type I or Type II, acquired prothrombin deficiency and von Willebrand disease.

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Present invention also relates to a method for preparing α -thrombin which comprises the following steps:

1. Modify, preferably by site-direct mutagenesis, the DNA sequence encoding for the factor Xa cleavage site in wild type prethrombin-2 gene to the DNA sequence encoding for the α -thrombin cleavage site.
2. Insert the mutated DNA prethrombin-2 into an expression vector.
3. Transform a host cell with the vector disclosed in previous step.
4. Culturing the transformed host cell of previous step in an appropriate culture medium.
5. Harvest transformed host cells and solubilize proteins in inclusion bodies.
6. Isolating the supernatant to obtain purified mutant prethrombin-2.
7. Contacting the purified mutant prethrombin-2 obtained in previous step with α -thrombin.
8. Purifying the α -thrombin obtained in previous step 6.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the amino acid sequence of factor Xa cleave site in wild type prethrombin-2 is SEQ ID NO: 17.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the amino acid sequence of the thrombin cleave site in mutant prethrombin-2 is SEQ ID NO: 18.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the ADN sequence of the wild type prethrombin-2 is selected from the group comprising SEQ ID NO: 5 or SEQ ID NO: 6.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the vector is preferably a plasmid. Any vector or plasmid known in the prior art for the same purpose could be used. In this sense, any vector suitable for expression in bacteria containing T7 promoter and coding a N-terminal His tag may be employed. On the other hand, any vector suitable for expression in yeast, containing galactose or methanol inducible promoters and coding a His tag may also be used in present invention. More particularly the plasmid is pET-TEV-prethr-2 plasmid or pQE9-hprethr-2 plasmid.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the host cell is selected from the group comprising: bacterial cells, animal cells and/or yeast.

In a more preferred embodiment of the invention the bacterial host cell is *E. coli*, preferably, *E. coli* XL1-blue, or *E. coli* BL21 or any strain derived thereof.

In a more preferred embodiment of the invention the host cell is a yeast selected from the group comprising: *Saccharomyces* sp. or *Pichia pastoris* or any strain derived thereof.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the ratio prethrombin-2: α -thrombin of step 7 ranged $1:10^{-6}$.

In a preferred embodiment of the invention, the method disclosed herein it is characterized in that the purification of α -thrombin comprises at least one step of techniques of affinity chromatography.

Present invention also discloses a kit for detecting α -thrombin/prothrombin in a sample comprising the mutant prethrombin-2, murine or human as disclosed herein.

Present invention also discloses the use of the kit mentioned above for the in vitro diagnosis and/or prognosis of coagulation-related diseases.

In a preferred embodiment, the coagulation-related disease is selected from the group comprising: haemophilia, throm-

basis, inherited prothrombin deficiency Type I or Type II, acquired prothrombin deficiency and von Willebrand disease.

Present invention also disclosed an in vitro method of diagnosis and/or prognosis of coagulation-related diseases which comprises the determination in a sample from a subject of the expression or activity level of α -thrombin/prothrombin using the kit mentioned above and the comparison of said expression or activity level with respect to the values for the same expression or activity obtained from healthy controls.

In a preferred embodiment of the in vitro method, the coagulation-related disease is selected from the group comprising: haemophilia, thrombosis, inherited prothrombin deficiency: Type I or Type II, acquired prothrombin deficiency and von Willebrand disease.

In a preferred embodiment of the in vitro method, the sample of the subject is selected from blood, plasma and/or serum, more preferably, samples of plasma.

Although any methods and materials similar or equivalent to those described herein can also be used in the practice or testing of the present invention, the preferred methods and materials are now described. It will be appreciated by those of skill in the art that, in light of the present disclosure, numerous modifications and changes can be made in the particular embodiments exemplified without departing from the intended scope of the invention.

The following examples illustrate the invention and should not be considered in a limiting sense, but rather in an illustrative sense of the invention.

Example 1

Cloning and Site Specific Mutations of Prethrombin-2 Gene

The present invention discloses the change the cleavage site of a recombinant proenzyme, prethrombin-2, by site directed mutagenesis to obtain a stable artificial autocatalytic enzyme. To achieve this purpose, the inventors have cloned the WT gene of prethrombin-2 and modified the factor Xa cleavage site in it to a thrombin cleavage site to obtain a stable artificial autocatalytic enzyme. The mutation strategy relies on converting the factor Xa cleavage site (Arg-Ile bond) into the α -Thrombin cleavage site by directed mutagenesis to create a self-replicative protease.

Briefly, for this approach, the sequences corresponding to human and mouse prethrombin-2 genes were cloned and five single mutations were introduced by, as mentioned above, site-directed mutagenesis in order to create a mutant prethrombin-2.

First, the vector pCMV-SPORT6 containing the full length mouse prothrombin cDNA was obtained from Geneservice (UK). The prethrombin-2 fragment was amplified from the same vector by PCR using PfuTurbo (Stratagene) and PCR nucleotide mix (Promega). Amplification primers were SEQ ID NO: 9 and SEQ ID NO: 10 containing BamHI and XhoI restriction sites respectively. The PCR product was run on an agarose gel, purified with QIAquick gel extraction kit (Qiagen) and digested with BamHI and XhoI (Takara). Afterwards, the fragment was purified using the QIAquick PCR purification kit (Qiagen) and ligated into the pET-TEV expression vector, a pET-19b vector modified by introducing a TEV protease cleavage site (generously provided by Dr. Lars Backman) giving pET-TEV-prethr-2 plasmid. The correctness of the inserted sequence was verified (MWG Germany). pET-TEV-prethr-2 plasmid was used as a template to mutate the prethrombin-2 gene. Site-direct specific mutations were introduced using the following primers: SEQ ID NO: 11

and SEQ ID NO: 12 with QuikChange II Site-Directed Mutagenesis Kit (Stratagene). The correctness of the mutant sequence was verified (MWG Germany).

The vector pOTB7 containing the full length human prothrombin cDNA was obtained from Geneservice (UK). The prethrombin-2 fragment was amplified from the same vector by PCR using PfuTurbo (Stratagene) and PCR nucleotide mix (Promega). Amplification primers were SEQ ID NO: 13 and SEQ ID NO: 14 containing Sall and HindIII restriction sites respectively. The PCR product was run on an agarose gel, purified with QIAquick gel extraction kit (Qiagen) and digested with Sall and HindIII (Takara). Afterwards, the fragment was purified using the QIAquick PCR purification kit (Qiagen) and ligated into the pQE9 expression vector (Qiagen), giving pQE9-hprethr-2 plasmid. The correctness of the inserted sequence was verified (MWG Germany). pQE9-hprethr-2 plasmid was used as a template to mutate the human prethrombin-2 gene. Site-direct specific mutations were introduced using the following primers SEQ ID NO: 15 and SEQ ID NO: 16 with QuikChange II Site-Directed Mutagenesis Kit (Stratagene). The correctness of the mutant sequence was verified (MWG Germany).

As a result, the, the FXa cleavage site present in wild type (WT) prethrombin-2, corresponding to residues with SEQ ID NO: 17 was changed to residues with SEQ ID NO: 18 which corresponds to the thrombin cleavage site. FIG. 1 shows the alignment of the FXa and thrombin cleavage sites of mutant prethrombin-2 and WT respectively. Changed amino acids are underlined. Thrombin selectively cleaves Arg-Gly bonds in fibrinogen and other polypeptides. Studies on the thrombin cleavage site from 30 different polypeptides revealed that the optimum cleavage site has the structure of P4-P3-Pro-Arg-P1'-P2' where P4 and P3 are hydrophobic amino acids and P1' and P2' are non acidic amino acids. It was also observed that polypeptides which contained Gly at P1' were especially susceptible to thrombin cleavage [24].

Example 2

Expression and Purification of Mutant Prethrombin-2

The plasmid contained the mutated sequences of human and mouse prethrombin-2 were used to transform *Escherichia coli* cells and the human and mouse mutant prothrombin-2 proteins were expressed and purified.

Particularly, *Escherichia coli* BL21 (DE3) cells were transformed by heat shock with the plasmid containing the mutant mouse prethrombin-2 gene. Cells were grown at 37° C. in Luria-Bertani media supplemented with 100 μ g/ml ampicillin to reach an OD600 \approx 0.7. Protein expression was induced by adding 0.1 mM isopropyl thio- β -D-galactoside. Cells were grown for 4 h at 37° C. and harvested by centrifugation. Protein purification from inclusion bodies was performed as described in Soejima et al. [25] with some modifications. After protein refolding, the sample was dialyzed against 50 mM Tris-HCl, pH 7.6, 150 mM NaCl, during two days at 4° C. without stirring. The dialyzed sample was centrifuged and the resulting supernatant was filtered. The pass-through fraction was loaded on a nickel HisTrap™ HP affinity column (GE Healthcare) controlled by an ÄKTA purifier equipment (GE Healthcare). Protein was eluted with a linear gradient from 0 to 500 mM of imidazole in 50 mM Tris-HCl, pH 7.6, 150 mM NaCl. Protein purity was investigated by 12% SDS-PAGE.

Electrophoretic analysis of the purified protein showed a single band corresponding to the expected size (FIG. 2A).

Protein identity was confirmed by Western blot using antibodies against His-tagged polypeptides (FIG. 2B).

For the purification of the human mutated prethrombin-2, *E. coli* XL1-blue cells were transformed by heat shock with the plasmid containing the mutant human prethrombin-2 gene. Cells were grown at 37° C. in Luria-Bertani media supplemented with 100 µg/ml ampicillin to reach an OD₆₀₀≈0.6. Protein expression was induced by adding 1 mM isopropyl thio-β-D-galactoside. Cells were grown for 3 h at 37° C. and harvested by centrifugation. Protein purification from inclusion bodies was performed as described before for the mouse mutant prethrombin-2. After protein refolding, the sample was dialyzed against 50 mM Tris-HCl, pH 7.0, at 4° C. without stirring. The dialyzed sample was centrifuged and the resulting supernatant was filtered. The pass-through fraction was loaded on a HiTrap™ HeparinHP affinity column (GE Healthcare) controlled by an ÄKTA purifier equipment (GE Healthcare). Protein was eluted with a linear gradient from 0 to 2 M NaCl in 50 mM Tris-HCl, pH 7.0. Protein purity was also investigated by 12% SDS-PAGE (FIG. 2A).

Western Blot

Proteins, mouse mutant prethrombin-2 and human mutant prethrombin-2, were separated by 12% SDS-PAGE and electroblotted onto a nitrocellulose membrane. The membrane was then incubated for 1 h in phosphate-buffered saline (PBS), containing 0.05% Tween 20, and 10% defatted milk. Afterwards, the membrane was washed with PBS containing 0.05% Tween 20 and incubated with anti-His mouse monoclonal antibodies (GE Healthcare) for 1 h. Then, the membrane was washed with PBS containing 0.05% Tween 20 and further incubated with secondary rabbit antibodies antimouse IgG conjugated to alkaline phosphatase (Sigma-Aldrich) for 1 h. After washing with PBS containing 0.05% Tween 20, protein bands were developed with 5-bromo-4-chloro-3-indolyl phosphate/nitro blue tetrazolium solution (Fluka) (FIG. 2B).

Example 3

Stability Assay of Mutant Prethrombin-2

Stability assays were performed by mixing human WT or human mutant prethrombin-2 (1.6×10^{-6} M) both produced by recombination in *E. coli*, with the fluorogenic substrate rhodamine 110, bis-(p-tosyl-L-glycyl-L-prolyl-L-arginine amide) (4×10^{-6} M) (Invitrogen) in 50 mM Tris-HCl, pH 9.0, 150 mM NaCl. The fluorescence of the resulting solutions was monitored using $\lambda_{\text{excitation}}=498$ nm and $\lambda_{\text{emission}}=521$ nm.

Stability assay measures in real time the enzymatic activity of active α-thrombin in the presence of the fluorogenic substrate (p-tosyl-Gly-Pro-Arg)2-rhodamine 110 (Invitrogen). This substrate contains the amino acid sequence recognized by α-thrombin, which is a peptidase. Thrombin cleaves the amino acid releasing the fluorescent rhodamine. During the test, it was measured the increase in fluorescence over time. The more enzyme activity of α-thrombin higher the cleavage speed, and therefore the release of rhodamine, which causes the increase in fluorescence. FIG. 3 shows that the WT human recombinant prethrombin-2 (a) is less stable than the mutant human recombinant prethrombin-2 discloses in the present invention (b). The test with mutated recombinant prethrombin-2 of the invention shows no increase in fluorescence for 15 h (55 000 sec). This implies that in preparation of mutated recombinant prethrombin-2 of the invention there are not residues of active α-thrombin, neither are being produced during the time that test takes place. On the other hand, the

test with WT human recombinant prethrombin-2 displays an increase in the fluorescence from the beginning of the test. This increase in fluorescence indicates the presence of active α-thrombin in the preparation; even though WT human recombinant prethrombin-2 was also obtained by genetic engineering in *E. coli*. Therefore the stability in vitro shown by mutant human prethrombin of present invention was due to the mutagenesis operated herein.

Example 4

α-Thrombin Activity Assays

In order to verify whether human α-thrombin could cleave mutated recombinant human and mouse prethrombins-2, were incubated both mutated recombinant proteins with WT human α-thrombin (Sigma Aldrich) in the presence of the fluorogenic substrate (p-tosyl-Gly-Pro-Arg)2-rhodamine 110 (Invitrogen). Thrombin activity assays were performed in a Varioskan Flash microplate reader (Thermo Scientific) using black microwell plates at room temperature. Samples (100 µl final volume) were incubated in 50 mM Tris-HCl, pH 9.0, 150 mM NaCl with 4×10^{-6} M rhodamine 110, bis-(p-tosyl-L-glycyl-L-prolyl-L-arginine amide) (Invitrogen) and the fluorescence of the resulting solution was monitored using $\lambda_{\text{excitation}}=498$ nm and $\lambda_{\text{emission}}=521$ nm.

The results obtained from the experiment display that in the presence of WT human α-thrombin, the protease activity increased considerably with time (FIG. 4, curves a, b), compared to the samples where only WT α-thrombin was added (FIG. 4, curve c). On the other hand, in the absence of WT α-thrombin, both mutants did not present enzymatic activity (FIG. 4, curves d, e), demonstrating their stability under the experimental conditions. These results clearly indicate that WT α-thrombin is able to cleavage both mutant recombinant prethrombins-2 to generate active enzymes with α-thrombin activity. Exogenous α-thrombin activates the mutant recombinant prethrombin-2 converting it into endogenous α-thrombin which in turn cleaves other macromolecules of prethrombin-2 and the fluorogenic substrate in the course of this autocatalytic reaction.

Example 5

Mutant Prethrombin-2 Self-Replication Assays

Another experiment proved that generated endogenous α-thrombin is able to cleavage mutated recombinant prethrombin-2 to maintain the autocatalytic process. A solution of mutated prethrombin-2 was treated with α-thrombin immobilized on agarose. Mouse mutant prethrombin-2 was cleaved using Thrombin CleanCleave™ Kit (Sigma-Aldrich) as described in manufacturer's directions. The protein solution was then filtered through a 0.22 µm filter to eliminate any remaining. To investigate the autocatalytic reaction, 5.38×10^{-9} M cleaved mouse mutant prethrombin-2 was mixed with and without 2.3×10^{-7} M untreated mouse mutant prethrombin-2 and with the fluorogenic substrate (p-tosyl-Gly-Pro-Arg)2-rhodamine 110 (Invitrogen). Then, the evolution of the fluorescence intensity was monitored using $\lambda_{\text{excitation}}=498$ nm and $\lambda_{\text{emission}}=521$ nm.

FIG. 5 shows that in the presence of the cleaved mutant prethrombin-2 and excess of the untreated mutant prethrombin-2 (FIG. 5 curve a), the fluorescence signal increased

considerably in comparison with the sample where only the cleaved mutant prethrombin-2 was included (FIG. 5 curve b). The untreated mutant prethrombin-2 treated with the buffer solution used to wash α -thrombin on agarose demonstrated no activity (FIG. 5 curve c). Consequently, growth in the protease activity was not caused by exogenous α -thrombin that could have detached from agarose. This experiment confirms the ability of endogenous α -thrombin derived from mutated prethrombin-2 to participate in the autocatalytic reaction.

Example 6

Detection of α -thrombin

Different concentrations of human α -thrombin (Sigma-Aldrich) were mixed with and without untreated 5.3×10^{-7} M mouse mutant prethrombin-2 disclosed herein. The reactions were carried out in 50 mM Tris-HCl, pH 9.0, 150 mM NaCl with 4×10^{-6} M rhodamine 110, bis-(p-tosyl-L-glycyl-L-prolyl-L-arginine amide) (Invitrogen) and the fluorescence of the resulting solution was monitored using $\lambda_{\text{excitation}}=498$ nm and $\lambda_{\text{emission}}=521$ nm.

FIG. 6A shows evolution of the fluorescence intensities for varying amounts of human α -thrombin mixed with the fluorogenic substrate in the presence (FIG. 6 curves a-h) or absence (FIG. 6 curves i-p) of mutated prethrombin-2. The fluorescence intensities demonstrated exponential growth in the presence of the mutant, pointing to the autocatalytic character of the reaction. In such case calibration plots of the first derivative of fluorescence intensity with respect to time dF/dt , representing the rate of reaction after 2 hours, versus analyte concentration (FIG. 6B) are more informative than the conventional plots [26]. On the basis of the obtained calibration curves we calculated the detection limit of the systems operating without mutated prethrombin-2 (0.488 pM, S/N=3, n=3) and with the mutant (10 fM, S/N=3, n=3).

Example 7

Detection of Prothrombin in Human Plasma

The present invention displays the advantage of the amplified assay for prothrombin over a conventional non amplified quantification method (FIG. 7). Varying concentrations of pooled human plasma (Sigma-Aldrich) were mixed with 6×10^{-8} M ecarin (Sigma-Aldrich) with or without 1.5×10^{-6} M mouse mutant prethrombin-2. The reactions were carried out in 50 mM Tris-HCl, pH 9.0, 150 mM NaCl with 4×10^{-6} M rhodamine 110, bis-(p-tosyl-L-glycyl-L-prolyl-L-arginine amide) (Invitrogen) and the fluorescence of the resulting solution was monitored using $\lambda_{\text{excitation}}=498$ nm and $\lambda_{\text{emission}}=521$ nm.

The detection limits of amplified (7.7 pL, S/N=3, RSD=12%, n=3) and non amplified (710 pL, S/N=3, RSD=6%, n=3) assays were calculated according to the calibration plots (FIG. 7B) depicting dF/dt at fixed time of 60 min versus volume of human plasma per microplate well. Taking into consideration that human plasma usually contains 90 $\mu\text{g/ml}$ of prothrombin [27] the assay with autocatalytic amplification allowed to detect as low as 0.693 pg of prothrombin per microplate well. The conventional non amplified assay allowed to quantify as low as 63.9 pg of the analyte per well. Thus, employment of autocatalytic amplification cascade allows diminishing the volume of human plasma needed for the prothrombin assay by two orders of magnitude.

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 <220> FEATURE:
 <221> NAME/KEY: mutation
 <222> LOCATION: (182)..(182)
 <223> OTHER INFORMATION: Mutated residues obtained by directed
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 <220> FEATURE:
 <221> NAME/KEY: mutation
 <222> LOCATION: (184)..(185)
 <223> OTHER INFORMATION: Mutated residues obtained by directed
 mutagenesis
 <220> FEATURE:

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<221> NAME/KEY: mutation
<222> LOCATION: (190)..(191)
<223> OTHER INFORMATION: Mutated residues obtained by directed
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ctgttcgaga agaagtcgct ggaggacaaa accgaaagag agctcctgga atcctacatc    180
gtcccgcgcg gtgtggaggg ctcgatgca gagatcgga tgtcaccttg gcagggtgatg    240
ctttccgga agagtcccca ggagctgctg tgtggggcca gcctcatcag tgaccgctgg    300
gtcctcaccg ccgccactg cctcctgtac ccgccctggg acaagaactt caccgagaat    360
gacctctggt tgcgcattgg caagcactcc cgcaccaggt acgagcgaaa cattgaaaag    420
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cgggacattg cctgatgaa gctgaagaag cctgttgctt tcagtgacta cattcacctt    540
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gtgacaggct ggggcaacct gaaggagacg tggacagcca acgttggtaa ggggcagccc    660
agtgtcctgc aggtggtgaa cctgcccatt gtggagcggc cggctctgca ggactccacc    720
cggatccgca tcaactgaca catgttctgt gctggttaca agcctgatga agggaaacga    780
gggatgcct gtgaaggatg cagtggggga ccctttgtca tgaagagccc cttaacaac    840
cgctggtatc aaatgggcat cgtctcatgg ggtgaaggct gtgaccggga tgggaaatat    900
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<210> SEQ ID NO 7
<211> LENGTH: 924
<212> TYPE: DNA
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cttcttgact cttacataga cgggcgcata gtggagggct gggacgctga gaagggtatc    180
gccccctgga aggtgatgct ttttcggaag agtccccaag agctgctgtg tggggccagc    240
cttatcagtg accgatgggt cctcactgct gcccaactgca ttctgtaacc acctgggac    300
aagaacttca ctgagaatga cctcctggtg cgcattggca agcattcccg aaccagatat    360
gagcggaaat ttgaaaagat ctccatgctg gaaaagatct acgtccacc cagatataac    420
tggcgggaga acctagaccg cgatatcgct ctgctcaagc taaagaaacc tgtacccttc    480
agtgactata ttcaccccgt gtgtttgcca gacaagcaga cagtaaccag cttgctccgg    540
gctggttata aagggcgggt gacaggctgg ggcaaccttc gggagacatg gacaaccaac    600
atcaatgaga tacagcccag cgtcctgcag gtggtgaacc tgcccattgt agagcggcca    660
gtgtgcaagg cctccacccc gattcgaatt actgacaaca tgttctgtgc tggcttcaag    720
gtgaatgaca ccaagcgagg agatgcttgt gaaggtgaca gtggaggacc tttgtcatg    780
aagagcccct ttaacaaccg ctggtatcaa atgggtattg tctcatgggg tgaaggatgt    840
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 aaagtcattg atcaatttgg atag 924

<210> SEQ ID NO 8
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 <213> ORGANISM: Homo sapiens

<400> SEQUENCE: 8

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 ctgttcgaga agaagtcgct ggaggacaaa accgaaagag agctcctgga atcctacatc 180
 gacgggcgca ttgtggaggg ctcggatgca gagatcgga tgtcaccttg gcagggtgatg 240
 cttttccgga agagtcccca ggagctgctg tgtggggcca gcctcatcag tgaccgctgg 300
 gtctcaccg ccgccactg cctcctgtac ccgccctggg acaagaactt caccgagaat 360
 gacctctcgg tgcgcattgg caagcactcc cgcaccaggt acgagcgaaa cattgaaaag 420
 atatccatgt tggaaaagat ctacatccac cccaggtaca actggcggga gaacctggac 480
 cgggacattg cctgatgaa gctgaagaag cctgttgctt tcagtgacta cattcacctt 540
 gtgtgtctgc ccgacagga gacggcagcc agcttgcctc aggtggata caagggggcg 600
 gtgacaggct ggggcaacct gaaggagacg tggacagcca acgttggtaa ggggcagccc 660
 agtgtcctgc aggtggtgaa cctgcccatt gtggagcggc cggctctcaa ggactccacc 720
 cggatccgca tcaactgaca catgttctgt gctggttaca agcctgatga agggaaacga 780
 ggggatgcct gtgaaggatg cagtggggga ccctttgtca tgaagagccc cttaacaac 840
 cgctggtatc aatgggcat cgtctcatgg ggtgaaggct gtgaccgga tgggaaatat 900
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<210> SEQ ID NO 9
 <211> LENGTH: 28
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 <213> ORGANISM: Artificial Sequence
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 <223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 9

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<210> SEQ ID NO 10
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 <212> TYPE: DNA
 <213> ORGANISM: Artificial Sequence
 <220> FEATURE:
 <223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 10

ccgctcgagc tatccaaatt gatcaatgac 30

<210> SEQ ID NO 11
 <211> LENGTH: 45
 <212> TYPE: DNA
 <213> ORGANISM: Artificial Sequence
 <220> FEATURE:
 <223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 11

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<210> SEQ ID NO 12
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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 12

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<210> SEQ ID NO 13
<211> LENGTH: 25
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 13

acgcgtcgac accgccacca gtgag 25

<210> SEQ ID NO 14
<211> LENGTH: 27
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 14

cccaagcttc tactctcaa actgatc 27

<210> SEQ ID NO 15
<211> LENGTH: 44
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 15

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<210> SEQ ID NO 16
<211> LENGTH: 44
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 16

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<210> SEQ ID NO 17
<211> LENGTH: 6
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 17

Ile Asp Gly Arg Ile Val
1 5

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<210> SEQ ID NO 18
 <211> LENGTH: 6
 <212> TYPE: PRT
 <213> ORGANISM: Homo sapiens

<400> SEQUENCE: 18

Ile Val Pro Arg Gly Val
 1 5

The invention claimed is:

1. A mutant prethrombin-2 comprising (a) the polypeptide sequences SEQ ID NO: 1 or SEQ ID NO: 2 or (b) polypeptides encoded by SEQ ID NO: 5 or SEQ ID NO: 6.
2. The mutant prethrombin-2 according to claim 1 comprising (a) the polypeptide sequence SEQ ID NO: 2 or (b) polypeptide encoded by SEQ ID NO: 6.
3. The mutant prethrombin-2 according to claim 1 characterized by having autocatalytic activity.
4. A method for preparing α -thrombin which comprises the following steps:
 - (a) modifying a DNA sequence encoding for a factor Xa cleavage site represented by SEQ ID NO:17 in wild type prethrombin-2 gene represented by SEQ ID NO:7 or SEQ ID NO:8 to a DNA sequence encoding for an α -thrombin cleavage site represented by SEQ ID NO:18;
 - (b) inserting a mutated DNA prethrombin-2 of step (a) into an expression vector;
 - (c) transforming a host cell with the expression vector of step (b);
 - (d) culturing the transformed host cell in an appropriate culture medium;
 - (e) harvesting the transformed host cell and solubilizing proteins in inclusion bodies;
 - (f) purifying mutant prethrombin-2;
 - (g) contacting the purified mutant prethrombin-2 obtained with α -thrombin; and
 - (h) purifying the α -thrombin.
5. The method according to claim 4 wherein the expression vector is a plasmid.
6. The method according to claim 5 wherein the plasmid is pET-TEV-prethr-2 plasmid or pQE9-hprethr-2 plasmid.
7. The method according to claim 4 wherein the host cell is selected from the group consisting of bacterial cells, animal cells, and yeast.
8. The method according to claim 7 wherein the bacterial host cell is *E. coli*.
9. The method according to claim 7 wherein the host cell is yeast selected from the group consisting of *Saccharomyces* sp and *Pichia pastoris*.
10. The method according to claim 4 wherein the purification of α -thrombin comprises at least one step of affinity chromatography.
11. A kit for detecting α -thrombin/prothrombin in a sample comprising the mutant prethrombin-2 of claim 1, and instructions for use thereof.
12. The method according to claim 8 wherein the bacterial host cell is *E. coli* XL1-blue or *E. coli* BL21.
13. The mutant prethrombin-2 of claim 3, wherein the autocatalytic activity is located at the alpha-thrombin cleavage site represented by the amino acid sequence SEQ ID NO:18.
14. The mutant prethrombin-2 according to claim 3, characterized by being stable in vitro.

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