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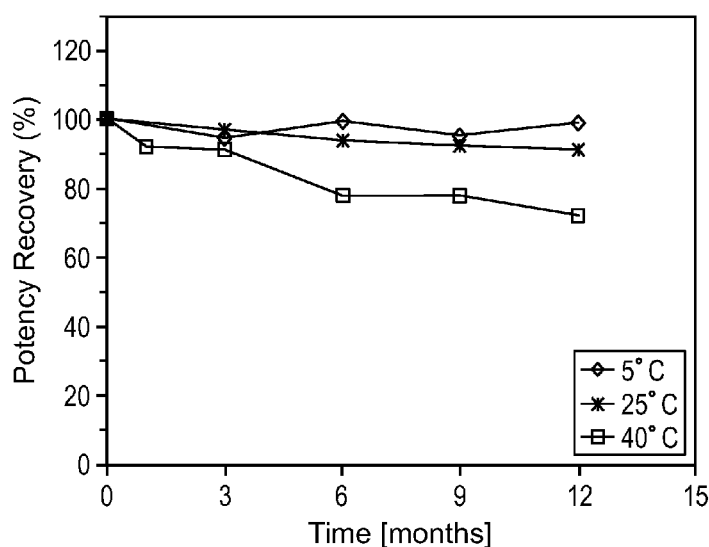
(54) **Title:** RECOMBINANT FACTOR VIII FORMULATIONS

FIG. 18

(57) **Abstract:** Provided are liquid and lyophilized recombinant Factor VIII formulations, including formulations for polymer-conjugated FVIII such as PEGylated Factor VIII.

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## RECOMBINANT FACTOR VIII FORMULATIONS

### BACKGROUND

[0001] This application claims the benefit of U.S. Provisional Application No. 61/779,495, filed March 15, 2013 and U.S. Provisional Application No. 61/869,191, filed August 23, 2013, both of which are hereby incorporated herein by reference in their entireties.

[0002] Hemophilia A is caused by deficiencies in coagulation factor VIII ("FVIII") and is the most common hereditary coagulation disorder, with an estimated incidence of 1 per 5000 males. The current treatment for hemophilia A involves intravenous injection of recombinant or plasma-derived human FVIII. Injections of FVIII are either given on demand in response to a bleeding event or as a prophylactic therapy that is administered 2 to 4 times a week. Although numerous studies have shown that prophylactic therapy decreases the complications of hemophilia A, the need for frequent intravenous injections creates barriers to patient compliance and affects patient quality of life. The requirement for frequent injections is primarily due to the short circulating FVIII half-life of 12 to 14 hours in patients.

[0003] Covalent addition of long-chain polymers, such as polyethyleneglycol ("PEG"), has been shown to increase the half-life of protein therapeutics. PEGylation is the covalent attachment of PEG molecules to proteins.

[0004] U.S. Patent No. 5,763,401 (Nayar) discloses stable, albumin-free, lyophilized formulations of full-length recombinant FVIII ("FL-rFVIII"). U.S. Patent No. 7,632,921 (Pan et al.) and Mei *et al.*, *Rational design of a fully active, long-acting PEGylated factor VIII for hemophilia A treatment*, 116 BLOOD 270-279 (2010) disclose cysteine enhanced FVIII mutants that are covalently bound to one or more biocompatible polymers such as PEG. U.S. Patent No. 7,087,723 (Besman et al.) pertains to albumin-free FVIII formulations. Österberg et al., *Development of a freeze-dried albumin-free formulation of recombinant factor VIII SQ*, Pharmaceutical Research, vol. 14, No. 7 (1997), pp. 892-898 and Österberg et al., *B-domain deleted recombinant factor VIII formulation and stability*, Seminars in Hematology, vol. 38, No. 2, suppl. 4 (April 2001), pp. 40-43 discuss formulations of B-domain deleted FVIII for lyophilization, including formulations containing sodium chloride, sucrose, histidine, calcium chloride dehydrate and polysorbate 80. Fatouros et al., *Recombinant factor VIII SQ – influence of oxygen, metal ions, pH and ionic strength on its stability in aqueous solution*, Int. J. of Pharmaceutics 155 (1997) 121-131

discloses the properties of rFVIII SQ on storage in solutions without albumin. WO2011/027152 (Jezek et al.) discloses formulations of FVIII.

#### SUMMARY

**[0004a]** In a first aspect the present invention provides a rFVIII formulation comprising:

- (a) a range of from about 1 mM to about 5 mM divalent cation;
- (b) a range of from about 150 mM to about 250 mM sodium chloride or potassium chloride;
- (c) a range of from about 50 ppm to about 200 ppm of a non-ionic surfactant; and
- (d) a range of from about 100 IU/ml to about 5000 IU/ml of a rFVIII, wherein the rFVIII comprises an amino acid sequence that has one or more non-cysteine residues in the amino acid sequence of SEQ ID NO: 3 replaced with cysteine residues such that at least one pair of cysteine residues creates a disulphide bond not found in wild type FVIII;

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5.

**[0004b]** In a second aspect the present invention provides a rFVIII formulation comprising:

- (a) about 0 mM, or a range of from about 1 mM to about 20 mM histidine;
- (b) a range of from 0.5% to 20% of sucrose or trehalose;
- (c) a range of from about 1 mM to about 5 mM divalent cation;
- (d) a range of from about 10 mM to about 50 mM sodium chloride;
- (e) about 0 mM, or a range of from about 20 ppm to about 80 ppm of a non-ionic

surfactant;

- (f) about 0%, or a range of from about 1.0% to about 5.0% glycine; and
- (g) a range of from about 100 IU/ml to about 5000 IU/ml of conjugated rFVIII;

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5.

**[0004c]** In a third aspect the present invention provides a method of treating hemophilia A comprising administering a therapeutically effective amount of a rFVIII formulation of the first or second aspects to a patient in need thereof.

**[0004d]** In a fourth aspect the present invention provides use of a rFVIII formulation of the first or second aspects in the manufacture of a medicament for treating hemophilia A.

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**[0005]** In one embodiment, the invention concerns a rFVIII formulation comprising: (a) a range of from about 1 mM to about 5 mM divalent cation; (b) a range of from about 150 mM to about 250 mM sodium chloride or potassium chloride; (c) a range of from about 50 ppm to about 200 ppm of a non-ionic surfactant; and (d) a range of from about 100 IU/ml to about 5000 IU/ml of a rFVIII, wherein the rFVIII comprises an amino acid sequence that has one or more non-cysteine residues in the amino acid sequence of SEQ ID NO: 3 replaced with cysteine residues; wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5. In another embodiment, the invention concerns a rFVIII formulation comprising: (a) a range of from 10 mM to 100 mM MOPS; (b) a range of from 0.5% to 10% by weight of a sugar or a sugar alcohol; (c) a range of from 0.5 mM to 20 mM of a divalent cation; (d) a range of from 10 mM to 100 mM sodium chloride or potassium chloride; (e) a range of from 50 to 150 ppm of a non-ionic surfactant; and (f) a range of from about 100 IU/ml to about 1500 IU/ml of rFVIII; wherein the rFVIII formulation contains less than 5.0 % by weight of components other than rFVIII having primary or secondary amine groups.

**[0006]** In yet another embodiment, the invention concerns a rFVIII formulation comprising: (a) a range of from 10 mM to 100 mM MOPS; (b) a range of from 150 mM to 300 mM NaCl; (c) a range of from 1 mM to 20 mM divalent cation; and (d) a range of from about 100 IU/ml to about 5000 IU/ml of nonconjugated rFVIII. In a further embodiment, the invention concerns a rFVIII formulation comprising: (a) a range of from 10 mM to 100 mM MOPS or histidine; (b) a range of from 25 mM to 200 mM NaCl; (c) a range of from 1 mM to 20 mM divalent cation; and (d) a range of from about 100 IU/ml to about 5000 IU/ml of conjugated rFVIII. In yet a further embodiment, the invention concerns a rFVIII formulation comprising: (a) about 0 mM, or a range of from about 1 mM to about 20 mM histidine; (b) a range of from 0.5% to 20% of sucrose or trehalose; (c) a range of from about 1 mM to about 5 mM divalent cation; (d) about 0 mM, or a range of from about 10 mM to about 50 mM sodium chloride; (e) about 0 mM, or a range of from about 20 ppm to about 80 ppm of a nonionic surfactant; (f) about 0%, or a range of from about 1.0% to about 5.0%, glycine and (g) a range of from about 100 IU/ml to about 5000 IU/ml of conjugated rFVIII; wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5.

## BRIEF DESCRIPTION OF THE DRAWINGS

[0007] The skilled artisan will understand that the drawings, described below, are for illustration purposes only. The drawings are not intended to limit the scope of the disclosure provided herein or the scope of the claims in any way.

[0008] FIG. 1 shows schematically the domains of full-length human factor VIII and BDD-rFVIII.

[0009] FIG. 2 is a graph showing the relative turbidity of BDD-rFVIII mutants having domains linked by disulfide bonds. The turbidity was measured in buffer comprising increasing concentration of sodium chloride. Turbidity was measured by  $A_{340nm}$ . In addition to sodium chloride, the buffer comprised 20 mM histidine, 2.5 mM calcium chloride, 29 mM sucrose, 293 mM glycine and 80 ppm polysorbate 80.

[0010] FIG. 3 is a graph showing the relative turbidity of BDD-rFVIII mutants having domains linked by disulfide bonds. The turbidity was measured in buffer comprising increasing concentration of polysorbate 80. Turbidity was measured by  $A_{340nm}$ . In addition to polysorbate 80, the buffer comprised 20 mM histidine, 30 mM sodium chloride, 2.5 mM calcium chloride, 29 mM sucrose and 293 mM glycine.

[0011] FIG. 4 is a graph showing the relative turbidity of BDD-rFVIII mutants having domains linked by disulfide bonds. The turbidity was measured in buffer comprising increasing concentration of human serum albumin ("HSA"). Turbidity was measured by  $A_{340nm}$ . The buffer comprised 20 mM histidine, 30 mM sodium chloride, 2.5 mM calcium chloride, 29 mM sucrose, 293 mM glycine and 80 ppm polysorbate 80.

[0012] FIG. 5 shows the relative turbidity of BDD-rFVIII mutants having domains linked by disulfide bonds. The turbidity was measured in a buffer comprising increasing concentration of sodium chloride in combination with polysorbate 80 and HSA. Turbidity was measured by  $A_{340nm}$ . In addition to sodium chloride, HSA and polysorbate 80, the buffer comprised 20 mM histidine, 2.5 mM calcium chloride, 29 mM sucrose and 293 mM glycine.

[0013] FIG. 6 shows clarity changes for BDD-rFVIII mutants with disulfide bonds linking domains in solution before and after addition of excipients. From left to right: (1) combination of excipients (HSA, sodium chloride and polysorbate 80), (2) HSA, (3) sodium chloride, (4) polysorbate 80, and (5) before addition of HSA, polysorbate 80 and sodium chloride.

[0014] FIG. 7 is a graph showing liquid stability of full-length FVIII in histidine, MOPS and TEA buffers during 7 days storage at 40°C.

[0015] FIG. 8 is a graph showing rFVIII stability in MOPS and histidine buffer at 25°C.

[0016] FIG. 9 is a diagram showing the structure of PEGylated BDD-rFVIII. The chains protruding from the A3 region represent the PEG molecule.

[0017] FIG. 10 is a graph showing the effect of sodium chloride on the potency recovery of PEGylated BDD-rFVIII during 6 days storage at 23°C.

[0018] FIG. 11 is a graph showing the effect of sodium chloride on the potency recovery of unPEGylated BDD-rFVIII during 6 days storage at 23°C.

[0019] FIG. 12 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation A after 26 weeks. Formulation A contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 293 mM glycine, 29 mM sucrose and 80 ppm polysorbate 80.

[0020] FIG. 13 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation B after 26 weeks. Formulation B contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 346 mM glycine, 38 mM sucrose and 80 ppm polysorbate 80.

[0021] FIG. 14 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation C after 26 weeks. Formulation C contains 2.5 mM calcium chloride, 20 mM histidine, 234 mM sucrose and 80 ppm polysorbate 80.

[0022] FIG. 15 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation D after 26 weeks. Formulation D contains 2.5 mM calcium chloride, 20 mM histidine, 211 mM trehalose and 80 ppm polysorbate 80.

[0023] FIG. 16 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation A up to 30 months. Formulation A contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 293 mM glycine, 29 mM sucrose and 80 ppm polysorbate 80.

[0024] FIG. 17 is a graph showing normalized potency trends for PEGylated BDD-rFVIII in lyophilized Formulation B up to 13 weeks. Formulation B contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 346 mM glycine, 38 mM sucrose and 80 ppm polysorbate 80.

[0025] FIG. 18 is a graph showing the normalized potency trends for PEGylated BDD-rFVIII (200 IU/mL) in lyophilized Formulation A up to 12 months. Formulation A



contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 293 mM glycine, 29 mM sucrose and 80 ppm polysorbate 80.

[0026] FIG. 19 is a graph showing the normalized potency trends for PEGylated BDD-rFVIII (1200 IU/mL) in lyophilized Formulation A up to 9 months. Formulation A contains 2.5 mM calcium chloride, 30 mM sodium chloride, 20 mM histidine, 293 mM glycine, 29 mM sucrose and 80 ppm polysorbate 80.

[0027] FIG. 20 is the amino acid sequence of BDD-rFVIII SQ (SEQ ID NO: 3).

[0028] FIG. 21 is the amino acid sequence of FL-rFVIII (SEQ ID NO: 1).

#### DETAILED DESCRIPTION

[0029] For the purposes of interpreting this specification, the following definitions will apply, unless otherwise indicated. All references cited herein are incorporated by reference herein in their entireties.

[0030] Factor VIII: Factor VIII ("FVIII") is a coagulation factor that circulates as a heterodimer composed of a heavy chain of approximately 200 kDa and a light chain of 80 kDa. The heavy chain contains structurally related A1 and A2 domains, as well as a unique B domain, and light chain comprises the A3, C1, and C2 domains. *See, e.g.,* Mei et al., 116 BLOOD 270-279 (2010). *See also* FIG.1, showing the domains of FVIII. The term "Factor VIII" or "FVIII" as used herein refers to all Factor VIII molecules, whether derived from blood plasma or produced through the use of recombinant DNA techniques, that have some procoagulant activity characteristic of wild type human FVIII. As used herein, FVIII includes modified or truncated forms of wild type or recombinant Factor VIII that retain some or all of the procoagulant activity of wild type Factor VIII or activated wild type Factor VIII, including variants or truncated forms that have procoagulant activity exceeding the activity of wild type Factor VIII or activated wild type Factor VIII. FVIII also includes fusion products containing active Factor VIII, such as fusions with an immunoglobulin fragment or domain. Commercially available examples of therapeutic preparations containing FVIII include those sold under the trade name KOGENATE FS (available from Bayer Healthcare LLC, Berkeley, CA, U.S.A.).

[0031] Recombinant Factor VIII: Recombinant Factor VIII ("rFVIII") as used herein refers to FVIII that is produced using recombinant technology, or a biologically active derivative thereof, and does not include FVIII obtained from mammalian plasma.

[0032] Full-length, native human Factor VIII ("FL-FVIII") is a 2,351 amino acid, single chain glycoprotein. The expressed 2,351 amino acid sequence is provided as SEQ. ID. NO: 1. When the expressed polypeptide is translocated into the lumen of the endoplasmic

reticulum, however, a 19-amino acid signal sequence is cleaved, resulting in a second sequence. This second sequence, herein provided as SEQ. ID. NO: 2, lacks the leading 19 amino acids and is the sequence conventionally used by researchers to assign a numeric location (e.g., Arg<sup>372</sup>) to a given amino acid residue of FVIII. Thus, unless specifically noted, all assignments of a numeric location of an amino acid residue as provided herein are based on SEQ. ID. NO: 2. For example, as is conventional and as used herein, when referring to mutated amino acids in BDD rFVIII, the mutated amino acid is designated by its position in the sequence of full-length FVIII. For example, a BDD rFVIII mutant can include a K1808C amino acid substitution wherein the lysine (K) at the position analogous to 1808 in the full-length sequence (here, SEQ ID NO: 2) is substituted to cysteine (C).

**[0033]** B-domain deleted (“BDD”) Factor VIII: As used herein, BDD or BDD-rFVIII is characterized by having an amino acid sequence with a deletion of all or part of the B-domain. In one embodiment, BDD is the molecule known as BDD-SQ, which contains a deletion of all but 14 amino acids of the B-domain of Factor VIII. In BDD-SQ, the first 4 amino acids of the B-domain (SEQ ID NO: 4) are linked to the 10 last residues of the B-domain (SEQ ID NO: 5). *See, e.g., Lind et al., Novel forms of B-domain-deleted recombinant factor VIII molecules*, 232 EUROPEAN JOURNAL OF BIOCHEMISTRY 19-27 (1995). *See also* FIG. 1 showing BDD by domain organization. BDD-SQ as used herein comprises the amino acid sequence of SEQ ID NO: 3. The B-domain of Factor VIII seems to be dispensable as a BDD molecule having a 90 kD A1-A2 heavy chain plus 80 kD light chain has been shown to be effective as a replacement therapy for hemophilia A. FVIII molecules having other portions of the B-domain deleted or all of the B-domain deleted are also included in the formulations and methods of the present invention.

**[0034]** BDD mutant or BDD-rFVIII mutant: BDD mutant or BDD-rFVIII mutant is a variant of BDD-SQ that maintains at least some of the FVIII procoagulant activity and differs from BDD-SQ by at least one amino acid residue. BDD mutant or BDD-rFVIII mutant includes variants that differ in amino acid sequence from BDD-SQ, for example, without limitation, by site-directed mutation of one or more amino acid residues. Without limitation, BDD mutant or BDD-rFVIII mutant includes the FVIII polypeptides with introduced cysteine residues disclosed in U.S. Patent No. 7,928,199 (Griffin et al.).

**[0035]** PEGylation: PEGylation is the covalent attachment of long-chain polyethylene glycol (PEG) molecules to proteins, such as by attaching a PEG that has an active functionality that binds to a site present on FVIII. One method used for PEGylation is the attachment of a functionalized PEG moiety to lysine residues or N-terminal amines that

are present in the native protein. Because FVIII contains many amine residues, amine-functionalized polymers are randomly conjugated to different sites on FVIII.

**[0036]** Site-directed PEGylation allows targeting of the PEG molecules to specific sites. These specific sites can include introduced surface-exposed cysteines to which the PEG polymer can be conjugated. *See* U.S. Patent No. 7,632,921 (Pan et al.). PEG may also be attached to FVIII by covalent linkage to a saccharide on FVIII. *See, e.g.,* U.S. Pat. App. Pub. 20110112028 (Turecek et al.). PEG may be attached to FVIII by enzymatic coupling of PEG to a glycan on FVIII, such as an O-glycan. Stennicke et al. disclose selective coupling of PEG to a unique O-glycan in the FVIII B-domain by incubating full-length FVIII with sialidase and excess CMP-SA-glycerol-PEG reagent in a buffer. Stennicke et al., “*A novel B-domain O-glycoPEGylated FVIII (N8-GP) demonstrates full efficacy and prolonged effect in hemophilic mice models,*” 121 (11) BLOOD 2108-16 (2013). U.S. Pat. App. Pub. 20130137638 (Bolt) discloses PEG attachment to a FVIII variant with a truncated B-domain. The FVIII molecule is covalently conjugated with a hydrophilic polymer via an O-linked oligosaccharide in the truncated B domain. U.S. Pat. App. Pub. 20120322738 (Behrens) discloses methods of conjugating polymers to FVIII, including covalently conjugating PEG to FVIII via an O-linked saccharide in the B-domain. As used herein, a PEGylated FVIII includes PEGylation by any method, including the various methods known in the art discussed above.

**[0037]** International Unit, IU: International Unit, or IU, is a unit of measurement of the blood coagulation activity (potency) of FVIII as measured by a standard assay. Standard assays include the one stage assay, as described in the art. *See, e.g.,* Lee et al., *An effect of predilution on potency assays of Factor VIII concentrates*, 30 THROMBOSIS RESEARCH 511-519 (1983). The one-stage assay is based upon the activated partial thromboplastin time (aPTT). FVIII acts as a cofactor in the presence of Factor IXa, calcium, and phospholipid in the enzymatic conversion of Factor X to Xa. In this assay, the diluted test samples are incubated at 37°C with a mixture of FVIII deficient plasma substrate and aPTT reagent. Calcium chloride is added to the incubated mixture and clotting is initiated. An inverse relationship exists between the time (seconds) it takes for a clot to form and logarithm of the concentration of FVIII:C. Activity levels for unknown samples are interpolated by comparing the clotting times of various dilutions of test material with a curve constructed from a series of dilutions of standard material of known activity and are reported in International Units per mL (IU/mL). Also useful are chromogenic assays, which may be purchased commercially, including the assay under the trade name COATEST SP FVIII (available from Chromogenix

AB, Molndal, Sweden). The chromogenic assay method consists of two consecutive steps where the intensity of color is proportional to the FVIII activity. In the first step, Factor X is activated to FXa by FIXa with its cofactor, FVIIIa, in the presence of optimal amounts of calcium ions and phospholipids. Excess amounts of Factor X are present such that the rate of activation of Factor X is solely dependent on the amount of FVIII. In the second step, Factor Xa hydrolyzes the chromogenic substrate to yield a chromophore and the color intensity is read photometrically at 405 nm. Potency of an unknown is calculated and the validity of the assay is checked with the slope-ratio statistical method. Activity is reported in International Units per mL (IU/mL).

**[0038]** Freeze-drying, freezing, lyophilizing: “Freeze-drying,” unless otherwise indicated by the context in which it appears, shall be used to denote the portion of a lyophilization process in which the temperature of a pharmaceutical preparation is raised in the primary and secondary drying phases in order to drive water out of the preparation. The “freezing” steps of a lyophilization process are those steps which occur prior to the primary and secondary drying stages. “Lyophilizing,” unless otherwise indicated, shall refer to the entire process of lyophilization, including both the freezing steps and the freeze-drying steps.

**[0039]** Within certain aspects of the present disclosure, formulations comprising rFVIII and BDD-rFVIII, including formulations comprising PEGylated FVIII and BDD-rFVIII, can be lyophilized according to methodology known in the art. For example, U.S. Patent Nos. 5,399,670 and 5,763,401 describe methodology for producing lyophilized FVIII formulations of enhanced solubility, which methodology may be employed to lyophilize the formulations described herein. The lyophilization process has a freezing phase, a primary drying phase, and a secondary drying phase. In the freezing phase, there is an annealing step. The freezing phase is performed at temperature not higher than -40°C, the annealing step occurs at temperature not higher than -15°C, the primary drying is performed at temperature not higher than 0°C, and the secondary drying is done at temperature not higher than 30°C. Once the set temperature is reached for the freezing temperature, annealing temperature, final freezing temperature, primary drying temperature, and secondary drying temperature, such temperature can be held for a reasonable time period as would be readily understood by one of skill in the art considering the particular protein sample involved, such as for one hour, two hours, three hours, or greater than three hours.

**[0040]** Anneal: The term “anneal” shall be used to indicate a step in the lyophilization process of a pharmaceutical preparation undergoing lyophilization, prior to the

freeze-drying of the preparation, in which the temperature of the preparation is raised from a lower temperature to a higher temperature and then cooled again after a period of time.

**[0041]** Bulking agent: For the purposes of this application, bulking agents are those chemical entities which provide structure to the “cake” or residual solid mass of a pharmaceutical preparation after it has been lyophilized and which protect it against collapse. A crystallizable bulking agent shall mean a bulking agent as described herein which can be crystallized during lyophilization.

**[0042]** Surfactant: As used herein, the term “surfactant” includes “non-ionic surfactants” such as polysorbates including polysorbate 20 and polysorbate 80, polyoxamers including poloxamer 184 or 188, pluronic polyols (sold under the trade name PLURONIC, manufactured by the BASF Wyandotte Corporation), and other ethylene/polypropylene block polymers. Non-ionic surfactants stabilize the rFVIII during processing and storage by reducing interfacial interaction and prevent protein from adsorption. The use of non-ionic surfactants permits the formulations to be exposed to shear and surface stresses without causing denaturation of the rFVIII. The formulations disclosed herein include formulations having one or more non-ionic surfactant(s), exemplified herein are formulations having a polysorbate, such as polysorbate 20 (sold under the trade name TWEEN 20) or polysorbate 80 (sold under the trade name TWEEN 80).

**[0043]** Osmolality: As used herein, the term “osmolality” refers to a measure of solute concentration, defined as the number of osmoles of solute per kg of solvent. A desired level of osmolality can be achieved by the addition of one or more stabilizer such as a sugar or a sugar alcohol including mannitol, dextrose, glucose, trehalose, and/or sucrose. Additional stabilizers that are suitable for providing osmolality are described in references such as the handbook of Pharmaceutical Excipients (Fourth Edition, Royal Pharmaceutical Society of Great Britain, Science & Practice Publishers) or Remingtons: The Science and Practice of Pharmacy (Nineteenth Edition, Mack Publishing Company). Formulations described herein have an osmolality ranging from about 240 mOsm/kg to about 450 mOsm/kg, or about 750 mOsm/kg, or about 1000 mOsm/kg, or from about 270 mOsm/kg to about 425 mOsm/kg, or from about 300 mOsm/kg to about 410 mOsm/kg.

**[0044]** Whenever appropriate, terms used in the singular also will include the plural and vice versa. The use of “a” herein means “one or more” unless stated otherwise or where the use of “one or more” is clearly inappropriate. The use of “or” means “and/or” unless stated otherwise. The use of “comprise,” “comprises,” “comprising,” “include,” “includes,” and “including” are interchangeable and not intended to be limiting. The term “such as” also

is not intended to be limiting. For example, the term “including” shall mean “including, but not limited to.” Furthermore, where the description of one or more embodiments uses the term “comprising,” those skilled in the art would understand that, in some specific instances, the embodiment or embodiments can be alternatively described using the language “consisting essentially of” and/or “consisting of.”

**[0045]** As used herein, the term “about” refers to +/- 10% of the unit value provided. As used herein, the term “substantially” refers to the qualitative condition of exhibiting a total or approximate degree of a characteristic or property of interest. One of ordinary skill in the biological arts will understand that biological and chemical phenomena rarely, if ever, achieve or avoid an absolute result because of the many variables that affect testing, production, and storage of biological and chemical compositions and materials, and because of the inherent error in the instruments and equipment used in the testing, production, and storage of biological and chemical compositions and materials. The term substantially is therefore used herein to capture the potential lack of completeness inherent in many biological and chemical phenomena.

**[0046]** The formulations of the invention described herein may be described in terms of the component concentrations by weight, such as by weight percent, or by molarity. It is to be understood that the invention also encompasses lyophilized preparations of these formulations that when reconstituted in suitable diluent, such as saline or water, for administration or storage have the concentrations reported. Ranges herein include the endpoints of the range.

**[0047]** Unless otherwise noted, percentage terms express weight/volume percentages and temperatures are in the Celsius scale.

#### COMPOSITION COMPONENTS

**[0048]** The FVIII compositions of the present invention may include stabilizing agents, buffering agents, sodium chloride, calcium salts, and, advantageously, other excipients. These excipients have been chosen in order to maximize the stability of FVIII in lyophilized preparations and/or in liquid preparations.

**[0049]** The bulking agents used in the present compositions are preferably selected from the group consisting of mannitol, glycine, and alanine. Mannitol, glycine, or alanine may be present in an amount of 1-5%, 2-3%, and 2.2-2.6%. Glycine may be the chosen bulking agent. Compositions are envisioned that do not contain a bulking agent.

[0050] The stabilizing agents used in the present compositions are selected from the group consisting of sugars or sugar alcohols, including without limitation sucrose, mannitol, dextrose, glucose and trehalose. These agents are present in the compositions in an amount of between 0.5-10%, 1-8%, 2-7%, 3-6%, 4-5%, 1-5%, 1-4%, 1-3%, or 1-2%. In compositions containing a bulking agent, sucrose is the preferred stabilizing agent in an amount of between 1-3%. In compositions lacking a bulking agent, sucrose or trehalose may be chosen as the stabilizing agent in an amount of about 8%. These sugars or sugar alcohols also function as cryo-protective agents.

[0051] In addition, buffers are present in certain of the inventive compositions. Buffers may be useful, for example, in FVIII formulations that are undergoing lyophilization, because it is believed that FVIII can be adversely affected by pH shifts during lyophilization. The buffering agents can be any physiologically acceptable chemical entity or combination of chemical entities which have the capacity to act as buffers, including histidine and MOPS (3-(N-morpholino) propanesulfonic acid). Histidine may be the chosen buffering agent in an amount of about 20 mM.

[0052] In order to preserve the activity of FVIII, the compositions of the present invention may also include calcium or another divalent cation able to interact with FVIII and maintain its activity, presumably by maintaining the association of the heavy and light chains of FVIII. Between 1 mM and 5 mM of a calcium salt can be used. The calcium salt can be calcium chloride, but can also be other calcium salts such as calcium gluconate, calcium glubionate, or calcium gluceptate. The FVIII compositions of the present invention also may include a surfactant, particularly a nonionic surfactant chosen from the group consisting of polysorbate 20 and polysorbate 80, polyoxamers including poloxamer 184 or 188, pluronic polyols (sold under the trade name PLURONIC, manufactured by the BASF Wyandotte Corporation), and other ethylene/polypropylene block polymers. The surfactant can be polysorbate 80 in an amount of about 80 ppm.

[0053] The FVIII used in the present compositions may be covalently attached to a biocompatible polymer, such as PEG. As used herein, the terms "polyethylene glycol" or "PEG" are interchangeable and include any water-soluble poly(ethylene oxide). PEG includes the following structure  $--(OCH_2CH_2)_n--$  where (n) is 2 to 4000. As used herein, PEG also includes  $--CH_2CH_2--O(CH_2CH_2O)_n--CH_2CH_2--$  and  $--(OCH_2CH_2)_nO--$ , depending upon whether or not the terminal oxygens have been displaced. The term "PEG" includes structures having various terminal or "end capping" groups, such as without limitation a hydroxyl or a C<sub>1-20</sub> alkoxy group such as methoxy. The term "PEG" also means a

polymer that comprises a majority, that is to say, greater than 50%, of --OCH<sub>2</sub>CH<sub>2</sub>--repeating subunits. With respect to specific forms, the PEG can take any number of a variety of molecular weights, as well as structures or geometries such as branched, linear, forked, and multifunctional. As used herein, the term "PEGylation" refers to a process whereby a polyethylene glycol (PEG) is covalently attached to a molecule such as a protein. When a functional group such as a biocompatible polymer is described as activated, the functional group reacts readily with an electrophile or a nucleophile on another molecule.

**[0054]** The biocompatible polymer used in the conjugates disclosed herein may be any of the polymers discussed herein or known in the art. The biocompatible polymer is selected to provide the desired improvement in pharmacokinetics. For example, the identity, size and structure of the polymer is selected so as to improve the circulation half-life of FVIII or decrease the antigenicity of FVIII without an unacceptable decrease in activity. The polymer can include PEG. For example, the polymer can be a polyethylene glycol terminally capped with an end-capping moiety such as hydroxyl, alkoxy, substituted alkoxy, alkenoxy, substituted alkenoxy, alkynoxy, substituted alkynoxy, aryloxy and substituted aryloxy. In some embodiments, the polymer can include methoxypolyethylene glycol such as methoxypolyethylene glycol having a size range from 3 kD to 200 kD.

**[0055]** The polymer can have a reactive moiety. For example, the polymer can have a sulfhydryl reactive moiety that can react with a free cysteine on a functional FVIII polypeptide to form a covalent linkage. Such sulfhydryl reactive moieties include thiol, triflate, tresylate, aziridine, oxirane, S-pyridyl, or maleimide moieties. The polymer can be linear and include a "cap" at one terminus that is not strongly reactive towards sulfhydryls (such as methoxy) and a sulfhydryl reactive moiety at the other terminus. The conjugate can include PEG-maleimide having a size range from 5 kD to 64 kD. Alternatively, the polymer can have an amine reactive moiety such as succinimidyl propionate, succinimidyl butanoate, benzotriazole carbonate, hydroxysuccinimide, aldehyde such as propionaldehyde, butyraldehyde, acetal, piperidone, methylketone, etc. (*see, e.g.* U.S. Patent 7,199,223 (Bossard)).

**[0056]** The FVIII molecule may be conjugated to a biocompatible polymer via conjugation of the polymer to the carbohydrate moieties of FVIII. *See* US Pat. App. Pub. 20110112028 (Turecek et al.). A FVIII molecule may be conjugated to a water-soluble polymer by conjugating a water soluble polymer to an oxidized carbohydrate moiety of FVIII. The water soluble polymer in some embodiments is selected from the group consisting of PEG, polysialic acid ("PSA") and dextran. In still another aspect, the activated



water soluble polymer is selected from the group consisting of PEG-hydrazide, PSA-hydrazine and aldehyde-activated dextran. In another aspect of the invention, the carbohydrate moiety is oxidized by incubation in a buffer comprising  $\text{NaIO}_4$ .

**[0057]** Suitable FVIII proteins to be used in the present invention have homology to specific known amino acid sequences. For example, suitable FVIII variants for use in the present invention are variants that have at least about 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, or 99 percent homology to known FVIII amino acid sequences, for example to the amino acid sequence of full-length FVIII (SEQ ID NO: 1) or that of BDD-FVIII (SEQ ID NO: 3). Also useful in the invention are genetic variants having defined sequence differences from a known FVIII sequence, such as FVIII molecules that comprise an amino acid sequence having 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 1-10, 1-5, 2-6 or 3-8 differences in amino acid sequence when compared to a native, known, or control sequence, such as the amino acid sequences of full-length FVIII (SEQ ID NO: 1) or BDD-SQ (SEQ ID NO: 3). Allelic variants are also useful in the present invention. Examples of allelic variants of FVIII are those disclosed in U.S. Patent Application Pub. No. 2010/0256062 (Howard et al.); Howard et al., "*African-Americans Express Multiple Haplotypic Forms of the Wildtype Factor VIII (FVIII) Protein: A Possible Role for Pharmacogenetics in FVIII Inhibitor Development?*" Blood, Vol. 104, 2004, Abstract 384; and Viel, K.R. et al., "*Inhibitors of Factor VIII in Black Patients with Hemophilia*," The New England Journal of Medicine, Vol. 360, 2009, pp. 1618-27. Allelic variants include those with amino acid substitutions such as histidine for arginine at position 484 (R484H), glycine for arginine at position 776 (R776G), glutamic acid for aspartic acid at position 1241 (D1241E), and valine for methionine at position 2238 (M2238V). The numbering systems used to designate the amino acid substitutions are based on SEQ ID NO: 2 herein.

**[0058]** Methods of alignment of nucleotide and amino acid sequences for comparison are well known in the art. Alignments for the present invention may be measured using a suitable method, including by using the local homology algorithm (BESTFIT) of Smith and Waterman, Adv. Appl. Math 2:482 (1981), which may conduct optimal alignment of sequences for comparison; by using the homology alignment algorithm (GAP) of Needleman and Wunsch, J. Mol. Biol. 48:443-53 (1970); or by using the search for similarity method (Tfasta and Fasta) of Pearson and Lipman, Proc. Natl. Acad. Sci. USA 85:2444 (1988). The alignments may be performed by using computerized implementations of these algorithms, including, but not limited to: CLUSTAL in the PC/Gene program by

Intelligenetics, Mountain View, California, GAP, BESTFIT, BLAST, FASTA, and TFASTA in the Wisconsin Genetics Software Package, Version 8 (available from Genetics Computer Group (GCG® programs (Accelrys, Inc., San Diego, CA).). The CLUSTAL program is well described by Higgins and Sharp, *Gene* 73:237-44 (1988); Higgins and Sharp, *CABIOS* 5:151-3 (1989); Corpet et al., *Nucleic Acids Res.* 16:10881-90 (1988); Huang et al., *Computer Applications in the Biosciences* 8:155-65 (1992), and Pearson et al., *Meth. Mol. Biol.* 24:307-31 (1994). One program to use for optimal global alignment of multiple sequences is PileUp (Feng and Doolittle, *J. Mol. Evol.*, 25:351-60 (1987)) which is similar to the method described by Higgins and Sharp, *CABIOS* 5:151-53 (1989). The BLAST family of programs can be used for database similarity searches, such as for identifying other suitable FVIII molecules. See *CURRENT PROTOCOLS IN MOLECULAR BIOLOGY*, Chapter 19, Ausubel et al., eds., Greene Publishing and Wiley-Interscience, New York (1995).

**[0059]** It is believed that the B-domain of FVIII is dispensable for activity, as discussed above. In certain embodiments, the FVIII used in the invention may have all or some of the B-domain deleted. Accordingly, the present invention applies to FVIII variants or nucleotide sequences encoding such variants that comprise an amino acid sequence or encode an amino acid sequence having at least about 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, or 99 percent homology to amino acids 1-740 of the full-length FVIII (SEQ ID NO: 1) and at least about 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, or 99 percent homology to amino acids 1689-2351 of the full-length FVIII (SEQ ID NO: 1). Alternatively, the present invention applies to FVIII variants or nucleotide sequences encoding such variants that comprise an amino acid sequence or encode an amino acid sequence having at least about 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, or 99 percent homology to the full-length of amino acid sequence SEQ ID NO: 1.

**[0060]** In certain embodiments of the invention, the FVIII may be the result of site-directed mutation, such as to create a binding site on FVIII to covalently attach a biocompatible polymer such as PEG. Site-directed mutation of a nucleotide sequence encoding polypeptide having FVIII activity may occur by any method known in the art. Methods include mutagenesis to introduce a cysteine codon at the site chosen for covalent attachment of the polymer. This may be accomplished using a commercially available site-directed mutagenesis kit such as the STRATAGENE CQUICKCHANGE II site-directed

mutagenesis kit, the CLONETECH TRANSFORMER site-directed mutagenesis kit no. K1600-1, the INVITROGEN GENTAYLOR site-directed mutagenesis system no. 12397014, the PROMEGA ALTERED SITES II in vitro mutagenesis system kit no. Q6210, or the TAKARA MIRUS BIO LA PCR mutagenesis kit no. TAK RR016. Conjugates described herein may be prepared by first replacing the codon for one or more amino acids on the surface of the functional FVIII polypeptide with a codon for cysteine, producing the cysteine mutant in a recombinant expression system, reacting the mutant with a cysteine-specific polymer reagent, and purifying the mutein. In this system, the addition of a polymer at the cysteine site can be accomplished through a maleimide active functionality on the polymer. *See, e.g.,* U.S. Patent 7,632,921 (Pan et al.).

**[0061]** The amount of sulfhydryl reactive polymer used should be at least equimolar to the molar amount of cysteines to be derivatized and can be present in excess. A 5-fold or a 10-fold molar excess of sulfhydryl reactive polymer can be used. Other conditions useful for covalent attachment are within the skill of those in the art.

**[0062]** The predefined site for covalent binding of the polymer, *e.g.*, PEG, can be selected from sites exposed on the surface of the rFVIII or BDD rFVIII polypeptide that are not involved in FVIII activity or involved in other mechanisms that stabilize FVIII *in vivo*, such as binding to vWF. Such sites are also best selected from those sites known to be involved in mechanisms by which FVIII is deactivated or cleared from circulation. Sites for substituting an amino acid with a cysteine include an amino acid residue in or near a binding site for (a) low density lipoprotein receptor related protein, (b) a heparin sulphate proteoglycan, (c) low density lipoprotein receptor and/or (d) FVIII inhibitory antibodies. By "in or near a binding site" means a residue that is sufficiently close to a binding site such that covalent attachment of a biocompatible polymer to the site would result in steric hindrance of the binding site. Such a site is expected to be within 20 Å of a binding site, for example.

**[0063]** The biocompatible polymer can be covalently attached to the rFVIII or BDD rFVIII polypeptide, or mutant variant thereof, at one or more of the FVIII amino acid positions 81, 129, 377, 378, 468, 487, 491, 504, 556, 570, 711, 1648, 1795, 1796, 1803, 1804, 1808, 1810, 1864, 1903, 1911, 2091, 2118 and 2284. One or more sites, such as one or two, on the functional FVIII polypeptide may be the predefined sites for polymer attachment. In particular embodiments, the polypeptide is mono-PEGylated or diPEGylated, meaning one PEG or two PEG molecules are attached to each FVIII, respectively.

**[0064]** Site directed PEGylation of a FVIII mutant can also be achieved by: (a) expressing a site-directed FVIII mutant wherein the mutant has a cysteine replacement for an

amino acid residue on the exposed surface of the FVIII mutant and that cysteine is capped; (b) contacting the cysteine mutant with a reductant under conditions to mildly reduce the cysteine mutant and to release the cap; (c) removing the cap and the reductant from the cysteine mutant; and (d) after the removal of the reductant, treating the cysteine mutant with PEG comprising a sulfhydryl coupling moiety under conditions such that PEGylated FVIII mutein is produced. The sulfhydryl coupling moiety of the PEG is selected from the group consisting of thiol, triflate, tresylate, aziridine, oxirane, S-pyridyl and maleimide moieties, and can be maleimide.

[0065] In another embodiment a biocompatible polymer such as, e.g., PEG, is covalently attached through use of a polymer functionalized with an amine-specific functional group. The polymer may be functionalized with, for example, mPEG tresylate or mPEG succinimidyl succinate such that it is reactive at lysines on FVIII. The coupling can occur at random lysines on FVIII by adding activated mPEG in a solid state to a solution of FVIII and rotating at room temperature. The degree of modification may be loosely controlled by the level of excess activated mPEG used. Rostin et al., "*B-Domain Deleted Recombinant Coagulation Factor VIII Modified with Monomethoxy Polyethylene Glycol*," 11 Bioconjug. Chem., 2000, pp. 387-396. Further examples of PEGylation conditions and reagents are provided in U.S. Patent 7,199,223 (Bossard) and U.S. Patent 4,970,300 (Fulton). The present invention is also directed to methods for covalently attaching a biocompatible polymer to FVIII in which one of the liquid formulations of the invention is the solution in which the reaction occurs.

[0066] The present disclosure also provides methods for the treatment of hemophilia A in a patient, comprising the administration to the patient in need thereof a therapeutically effective amount of one or more formulations described herein. These formulations may be administered to a patient via intravenous injection, subcutaneous injection, or through continuous infusion.

[0067] As used herein, the term "therapeutically effective amount" of a rFVIII formulation or a PEGylated rFVIII formulation refers to an amount of the formulation that provides therapeutic effect in an administration regimen to a patient in need thereof. For example, for replacement therapy for hemophilia A, an amount of between 10-30 IU/kg body weight of recombinant full-length FVIII for intravenous injection is recommended. For prophylaxis in a child with hemophilia A, 25 IU/kg body weight of recombinant full-length FVIII for intravenous injection is recommended. Prior to surgery, 15-30 IU/kg (minor surgery) or 50 IU/kg (major surgery) of recombinant full-length FVIII for intravenous

injection is recommended for a child with hemophilia A. Corresponding dosages for the various FVIII molecules used in the formulations of the invention can be determined by those of skill in the art. Preferably the therapeutic FVIII formulations of the invention are provided in single use dosages of 100, 250, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3400, 3600, 3800, 4000, or 5000 IU, or in a range between any two of these dosages, i.e., in a range of from 100 to 250 IU, from 100 to 500 IU, from 1000 to 2000 IU, etc., inclusive of the endpoints. Because of their low viscosity, the presently disclosed rFVIII and PEG-rFVIII formulations can be conveniently processed via, for example, ultrafiltration and sterile filtration and can be administered to a patient via injection, including intravenous injection, subcutaneous injection, and continuous infusion.

**[0068]** The FVIII compositions described in this application can be lyophilized and reconstituted in the indicated concentrations. These FVIII compositions can also be reconstituted in more dilute form. For example, a preparation according the present invention which is lyophilized and/or normally reconstituted in 2 ml of solution can also be reconstituted in a larger volume of diluent, such as 5 ml. This is particularly appropriate when the FVIII preparation is being injected into a patient immediately, since in this case the FVIII is less likely to lose activity, which may occur more rapidly in more dilute solutions of FVIII.

## **EMBODIMENT 1**

**[0069]** Recombinant FVIII is produced in the absence of plasma proteins that stabilize plasma-derived FVIII, such as von Willebrand factor (vWF). The absence of such stabilizing proteins makes rFVIII extremely labile. In addition, rFVIII is present at very low concentrations in therapeutic solutions (0.02 mg protein per ml for a therapeutic dose of 1000 IU BDD-SQ), which makes surface adsorption a cause for loss of activity.

**[0070]** One embodiment of the invention is a formulation of rFVIII, particularly BDD-rFVIII, and even more particularly BDD-rFVIII mutants with cross-linking between the domains, such as between the A1 and A2 or A3 domains. In one embodiment the formulation is of a FVIII having double cysteine mutations which cross-link the A2 and the A1 or the A3 domains, preferably the A2 and the A3 domains, such as through disulfide bridges as described in U.S. Patent 7,928,199 to Griffin et al. (issued Apr. 19, 2011), including without limitation mutants of FVIII, including mutants of BDD SQ (SEQ ID NO: 3), in which one or more cysteines have been introduced at one or more sites; such that at least one pair of cysteines creates a disulfide bond not found in wild type FVIII. In one

embodiment, the mutant FVIII comprises at least one pair of recombinantly introduced cysteines, wherein the pair of cysteines replaces a pair of residues selected from the group consisting of Met 662 and Asp 1828, Ser 268 and Phe 673, Ile 312 and Pro 672, Ser 313 and Ala 644, Met 662 and Lys 1827, Tyr 664 and Thr 1826, Pro 264 and Gln 645, Arg 282 and Thr 522, Ser 285 and Phe 673, His 311 and Phe 673, Ser 314 and Ala 644, Ser 314 and Gln 645, Val 663 and Glu 1829, Asn 694 and Pro 1980, and Ser 695 and Glu 1844. Suitable FVIII molecules for the formulations of the present embodiment suffer the disadvantage of aggregating in solution and/or show a high propensity for precipitation. These disadvantages create problems preparing a stable therapeutic dosage. Also, if the FVIII molecules are to be further processed, such as by covalent attachment of a biocompatible polymer such as PEG, the FVIII molecules are preferably in solution to provide good processing, such as good yields upon PEGylation, which requires that the FVIII be in suspension or solution and not be aggregated. In one embodiment, the FVIII formulations of the present application contain sodium chloride or potassium chloride in an amount sufficient to reduce or abolish precipitation and/or aggregation and to provide stability.

**[0071]** Formulations of Embodiment 1 may be as follows. A rFVIII formulation comprising:

- (a) a range of from about 0 mM to about 20 mM, from about 1mM to about 20mM, from about 1 mM to about 50 mM, from about 10 mM to about 50 mM, from about 10 mM to about 20mM, from about 10 mM to about 30mM, or from about 20 mM to about 50 mM histidine;
- (b) a range of from about 0 mM to about 29 mM, from about 1 mM to about 29 mM, from about 1 mM to about 300 mM, from about 10 mM to about 30 mM, from about 10 mM to about 100 mM, from about 10 mM to about 200 mM, from about 10 mM to about 50 mM, from about 29 mM to about 58 mM, from about 34 mM to about 58 mM, from about 58 mM to about 100 mM, or from about 100 mM to about 300 mM, or an amount of about 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 60, 70, 80, 90, 100, 200, or about 300 mM of a sugar or sugar alcohol;
- (c) a range of from about 1 mM to about 2 mM, from about 1 mM to about 2.5 mM, from about 1.5 mM to about 3.5 mM, or from about 1 mM to about 5 mM divalent cation such as a divalent calcium salt, including calcium chloride;
- (d) a range of from about 150 mM to about 250 mM, from about 150 mM to about 220 mM, from about 150 mM to about 200 mM, from about 150 mM to about 190 mM,

from about 170 mM to about 250 mM; from about 200 mM to about 220 mM, from about 170 mM to about 200 mM, from about 200 mM to about 250 mM, from about 170 mM to about 220 mM, from about 190 mM to about 220 mM, from about 210 mM to about 220 mM, from about 150 mM to about 180 mM, from about 150 mM to about 160 mM, or from about 220 mM to about 250 mM sodium chloride or potassium chloride;

(e) a range of from about 20 ppm to about 200 ppm, from about 20 ppm to about 50 ppm, from about 20 ppm to about 80 ppm, from about 50 ppm to about 80 ppm, from about 80 ppm to about 100 ppm, from about 80 ppm to about 200 ppm, from about 50 ppm to about 100 ppm, or from about 50 ppm to about 200 ppm of a non-ionic surfactant, or about 60, 65, 70, 75, 80, 85, 90, 95, 100, 105, 110, 115, 120, 125, 130, 135, 140, 145, 150, 160, 170, 180, 190, 200, or 210 ppm of a non-ionic surfactant;

(f) a range of from about 0 mM to about 50 mM, from about 1 mM to about 50 mM, from about 50 mM to about 100 mM, from about 100 mM to about 150 mM, from about 150 mM to about 293 mM, from about 150 mM to about 400 mM, from about 200 mM to about 300 mM; from about 250 mM to about 300 mM, or from about 200 mM to about 400 mM glycine, or about 100, 200, 210, 230, 240, 250, 260, 270, 280, 290, 293, 295, 300, 310, 320, 330, 340, 350, 360, 370, 380, 390, or about 400 mM glycine; and

(g) a range of from about 100 IU/ml to about 5000 IU/ml, from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 100 IU/ml to about 4000 IU/ml, from about 100 IU/ml to about 1200 IU/ml, from about 250 IU/ml to about 5000 IU/ml, from about 250 IU/ml to about 1000 IU/ml, from about 250 IU/ml to about 2000 IU/ml, from about 250 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, from about 1000 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 4000 IU/ml, or from about 1000 IU/ml to about 5000 IU/ml, or about 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3500, 3800, 4000, 4200, 4500, 4800, 5000, 5500, or 6000 IU/ml of a rFVIII selected from rFVIII, BDD-rFVIII, BDD-rFVIII mutants, and BDD-rFVIII mutants with cross-linking between FVIII domains,

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 6.5, from about pH 6.0 to about pH 7.0, from about pH 6.0 to about pH 7.5, from about pH 6.5 to about pH 7.5, or from about pH 7.0 to about pH 7.5, or a pH of about pH 6.0, 6.5, 7.0, 7.1, 7.2, 7.3, 7.4 or about pH 7.5.

[0072] In another version of Embodiment 1, the invention pertains to a rFVIII formulation comprising:

- (a) a range of from about 0 mM to about 20 mM, from about 10 mM to about 50 mM, or from about 10 mM to about 30mM histidine;
- (b) a range of from about 1 mM to about 29 mM, from about 10 mM to about 30 mM, from about 10 mM to about 100 mM, from about 10 mM to about 200 mM, from about 10 mM to about 50 mM, from about 29 mM to about 58 mM, or from about 34 mM to about 58 mM, or an amount of about 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, or 40 mM of a sugar or sugar alcohol;
- (c) a range of from about 1 mM to about 2 mM, from about 1 mM to about 2.5 mM, from about 1.5 mM to about 3.5 mM, or from about 1 mM to about 5 mM divalent cation such as a divalent calcium salt, including calcium chloride;
- (d) a range of from about 150 mM to about 200 mM, from about 150 mM to about 220 mM, from about 170 mM to about 250 mM sodium chloride or potassium chloride;
- (e) a range of from about 20 ppm to about 80 ppm, from about 80 ppm to about 100 ppm, from about 50 ppm to about 100 ppm, or from about 50 ppm to about 200 ppm of a non-ionic surfactant, or about 70, 75, 80, 85, 90, 95, 100, 105, 110, 115, or 120 ppm of a non-ionic surfactant;
- (f) a range of from about 1 mM to about 50 mM, from about 150 mM to about 300 mM, from about 150 mM to about 400 mM, from about 200 mM to about 300 mM; or from about 250 mM to about 300 mM, or about 250, 260, 270, 280, 290, 293, 295, 300, 310, or about 320 mM glycine; and
- (g) a range of from about 100 IU/ml to about 5000 IU/ml, from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 100 IU/ml to about 4000 IU/ml, from about 100 IU/ml to about 1200 IU/ml, from about 250 IU/ml to about 5000 IU/ml, from about 250 IU/ml to about 1000 IU/ml, from about 250 IU/ml to about 2000 IU/ml, from about 250 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, from about 1000 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 4000 IU/ml, or from about 1000 IU/ml to about 5000 IU/ml, or about 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3500, 3800, 4000, 4200, 4500, 4600, 4800, 5000, 5500, or 6000 IU/ml of a rFVIII selected from rFVIII, BDD-rFVIII, BDD-rFVIII mutants, and BDD-rFVIII mutants with cross-linking between FVIII domains,



wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5, from about pH 6.5 to about pH 7.5, or from about pH 7.0 to about pH 7.5, or a pH of about pH 6.0, 6.5, 7.0, 7.1, 7.2, 7.3, 7.4 or about pH 7.5.

[0073] In certain embodiments the sugar or sugar alcohol is sucrose and sodium chloride is present.

[0074] In another version of Embodiment 1, the invention pertains to a rFVIII formulation comprising:

- (a) a range of from about 10 mM to about 30 mM histidine;
- (b) a range of from about 10 mM to about 30 mM, from about 10 mM to about 100 mM, from about 10 mM to about 200 mM, from about 10 mM to about 50 mM, from about 29 mM to about 58 mM, or from about 34 mM to about 58 mM of a sugar or sugar alcohol;
- (c) a range of from about 1 mM to about 2 mM, from about 1 mM to about 2.5 mM, from about 1.5 mM to about 3.5 mM, or from about 1 mM to about 5 mM of a divalent calcium salt, including calcium chloride;
- (d) a range of from about 150 mM to about 220 mM, from about 170 mM to about 250 mM sodium chloride or potassium chloride;
- (e) a range of from about 50 ppm to about 200 ppm of a non-ionic surfactant;
- (f) a range of from about 1 mM to about 50 mM, from about 150 mM to about 300 mM, from about 150 mM to about 400 mM, from about 200 mM to about 300 mM; or from about 250 mM to about 300 mM glycine; and
- (g) a range of from about 100 IU/ml to about 5000 IU/ml, from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 100 IU/ml to about 4000 IU/ml, from about 100 IU/ml to about 1200 IU/ml, from about 250 IU/ml to about 5000 IU/ml, from about 250 IU/ml to about 1000 IU/ml, from about 250 IU/ml to about 2000 IU/ml, from about 250 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, from about 1000 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 4000 IU/ml, or from about 1000 IU/ml to about 5000 IU/ml, or about 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3500, 3800, 4000, 4200, 4500, 4600, 4800, 5000, 5500, or 6000 IU/ml of a rFVIII selected from rFVIII, BDD-rFVIII, BDD-rFVIII mutants, and BDD-rFVIII mutants with cross-linking between FVIII domains, wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5, from about pH 6.5

to about pH 7.5, or from about pH 7.0 to about pH 7.5, or a pH of about pH 6.0, 6.5, 7.0, 7.1, 7.2, 7.3, 7.4 or about pH 7.5.

**[0075]** In yet another version of Embodiment 1, the invention pertains to a rFVIII formulation comprising:

- (a) a range of from about 10 mM to about 30 mM histidine;
- (b) a range of from about 10 mM to about 50 mM of sucrose;
- (c) a range of from about 1.5 mM to about 3.5 mM calcium chloride;
- (d) a range of from about 150 mM to about 220 mM or from about 170 mM to about 220 mM sodium chloride;
- (e) a range of from about 70 ppm to about 90 ppm of a non-ionic surfactant;
- (f) a range of from about 200 mM to about 300 mM or from about 250 mM to about 300 mM glycine; and
- (g) a range of from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 250 IU/ml to about 1000 IU/ml, from about 250 IU/ml to about 2000 IU/ml, from about 250 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, or from about 1000 IU/ml to about 3000 IU/ml of a rFVIII selected from BDD-rFVIII, BDD-rFVIII mutants, and BDD-rFVIII mutants with cross-linking between FVIII domains, wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5, from about pH 6.5 to about pH 7.5, or from about pH 7.0 to about pH 7.5, or a pH of about pH 6.0, 6.5, 7.0, 7.1, 7.2, 7.3, 7.4 or about pH 7.5.

**[0076]** The rFVIII formulations of embodiment 1 may optionally contain albumin, such as HSA. In certain embodiments, HSA is present at a range of from about 10 to about 50 mg/mL, from about 15 to about 30 mg/mL, from about 20 to about 30 mg/mL or from about 25 to about 30 mg/mL.

## **EMBODIMENT 2**

**[0077]** During covalent addition of a biocompatible polymer to FVIII it was observed that buffer components may interfere with the covalent addition. For example, when FVIII was covalently coupled to PEG using PEG functionalized to have an amine-reactive group that would add at lysine residues, amine-containing components in the reaction buffer were observed to interfere with the reaction. Accordingly, the present invention includes improved liquid FVIII formulations or buffers in which the polymer addition reaction to FVIII may occur. In one version of Embodiment 2, the liquid FVIII formulations do not comprise, or comprises less than 10% by weight, or less than 5% by

weight, or less than 1% by weight, or less than 0.5% by weight or only a trace amount of components with primary or secondary amine groups, other than FVIII. The inventive FVIII formulations of this embodiment include formulations that avoid the use of histidine and glycine. Histidine and glycine contain amines that may interfere with the PEGylation process.

**[0078]** One version of Embodiment 2 of the invention is a formulation of rFVIII having buffer capacity at pH 6-7 that does not form an insoluble complex or chelate with calcium chloride (an important rFVIII stabilizer) and does not contain components with primary or secondary amine groups, or contains such components at a weight percent of 10% or less, 5% or less, 1% or less, or in trace amounts. This formulation may include MOPS in a range of from 10 mM to 100 mM, in a range of from 10 mM to 70 mM, in a range of from 10 mM to 50 mM, in a range of from 10 mM to 40 mM, in a range of from 10 mM to 30 mM, in a range of from 12 mM to 30 mM, in a range of from 14 mM to 30 mM, in a range of from 16 mM to 30 mM, in a range of from 18 mM to 30 mM, in a range of from 20 mM to 28 mM, in a range of from 12 mM to 28 mM, in a range of from 12 mM to 26 mM, in a range of from 12 mM to 24 mM, in a range of from 12 mM to 22 mM, in a range of from 14 mM to 22 mM, or in a range of from 18 mM to 22 mM, or may contain about 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30 mM MOPS. This formulation includes rFVIII in a range of from about 100 IU/ml to about 1000 IU/ml, from about 100 IU/ml to about 500 IU/ml, from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 2000 IU/ml, from about 500 IU/ml to about 2500 IU/ml, from about 500 IU/ml to about 1200 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 1500 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, from about 1000 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2500 IU/ml, from about 1000 IU/ml to about 1500 IU/ml, from about 1000 IU/ml to about 6000 IU/ml, or from about 1000 IU/ml to about 5000 IU/ml or about 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3500, 3800, 4000, 4200, 4500, 4800, 5000, 5500, or 6000 IU/ml of rFVIII. It is also possible that the invention may be used with rFVIII formulations having higher activity than 6000 IU/ml.

**[0079]** In one version of Embodiment 2, the rFVIII formulation comprises FVIII or BDD that is recombinantly produced. In another version of Embodiment 2, the formulation comprises recombinantly produced full-length FVIII, such as FVIII comprising the amino

acid sequence of SEQ ID NO: 1 or an allelic variant thereof. In another version of Embodiment 2, the formulation comprises a mutant of BDD or a mutant of FL-FVIII.

**[0080]** This formulation may also include a sugar or a sugar alcohol such as sucrose in a range of from 0.5% to 10%, in a range of from 0.6% to 10%, in a range of from 0.7% to 10%, in a range of from 0.8% to 10%, in a range of from 0.9% to 10%, in a range of from 1.0% to 10%, in a range of from 0.6% to 5%, in a range of from 0.6% to 2.5%, in a range of from 0.6% to 2.0%, in a range of from 0.6% to 1.5%, in a range of from 0.6% to 1.2%, in a range of from 0.8% to 1.2%, in a range of from 0.9% to 1.2%, or in a range of from 0.9% to 1.1% by weight, or at about 0.8, 0.9, 1.0, 1.1, 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, or 2.0% by weight. This formulation may also include a divalent cation such as a calcium salt, such as calcium chloride, in a range of from 0.5 mM to 20 mM, in a range of from 1 mM to 10 mM, in a range of from 1 mM to 5 mM, in a range of from 1.5 mM to 5 mM, in a range of from 2 mM to 5 mM, in a range of from 2.5 mM to 5 mM, in a range of from 3 mM to 5 mM, in a range of from 3.5 mM to 5 mM, in a range of from 4 mM to 5 mM, in a range of from 1.5 mM to 4.5 mM, in a range of from 1.5 mM to 4 mM, in a range of from 1.5 mM to 3.5 mM, in a range of from 1.5 mM to 3 mM, in a range of from 1.5 mM to 2.5 mM, in a range of from 2 mM to 3 mM, in a range of from 2.2 mM to 2.8 mM, or in a range of from 2.4 mM to 2.6 mM. This formulation may also include sodium chloride or potassium chloride in a range of from 10 mM to 100 mM, in a range of from 10 mM to 70 mM, in a range of from 10 mM to 50 mM, in a range of from 15 mM to 50 mM, in a range of from 20 mM to 50 mM, in a range of from 25 mM to 50 mM, in a range of from 30 mM to 50 mM, in a range of from 15 mM to 45 mM, in a range of from 15 mM to 40 mM, in a range of from 15 mM to 35 mM, in a range of from 20 mM to 45 mM, in a range of from 20 mM to 40 mM, in a range of from 25 mM to 40 mM, in a range of from 25 mM to 35 mM, in a range of from 25 mM to 30 mM, or in a range of from 30 mM to 35 mM. This formulation may also include a non-ionic surfactant such as polysorbate 20 or polysorbate 80 in a range of from 50 to 150 ppm, in a range of from 60 ppm to 150 ppm, in a range of from 70 ppm to 150 ppm, in a range of from 80 ppm to 150 ppm, in a range of from 60 ppm to 140 ppm, in a range of from 60 ppm to 130 ppm, in a range of from 60 ppm to 120 ppm, in a range of from 60 ppm to 110 ppm, in a range of from 60 ppm to 100 ppm, in a range of from 60 ppm to 90 ppm, in a range of from 70 ppm to 90 ppm, in a range of from 70 ppm to 80 ppm, and in a range of from 80 ppm to 90 ppm. This composition provides acceptable stability to rFVIII in solution, and can be used as a reaction buffer during the conjugation of a polymer to FVIII using a polymer functionalized to be active at amine residues.

[0081] In one version of Embodiment 2, the invention is related to a rFVIII formulation comprising

- (a) MOPS in a range of from 12 mM to 28 mM, in a range of from 12 mM to 22 mM, or in a range of from 18 mM to 22 mM;
- (b) FVIII in a range of from 100 IU/ml to 3000 IU/ml, or in a range of from 1000-1500 IU/ml;
- (c) sucrose in a range of from 0.5% to 5%, in a range of from 0.6% to 2.5%, or in a range of from 0.9% to 1.1%;
- (d) sodium chloride or potassium chloride in a range of from 10 mM to 50 mM, in a range of from 15 mM to 35 mM, or in a range of from 25 mM to 35 mM;
- (e) a divalent calcium salt, such as calcium chloride, in a range of from 1 mM to 5 mM, in a range of from 1.5 mM to 3.5 mM, or in a range of from 2.4 mM to 2.6 mM; and
- (f) non-ionic surfactant such as polysorbate 20 or polysorbate 80 in a range of from 60 ppm to 100 ppm, or in a range of from 70 ppm to 90 ppm;

wherein the rFVIII formulation contains less than 10%, less than 5%, less than 1%, less than 0.5%, or less than a trace level, or is essentially free, of a component having a primary or secondary amine group.

[0082] The invention also is directed to a method of conjugating an amine-reactive biocompatible polymer, such as an amine-reactive PEG, to FVIII comprising suspending or dissolving the FVIII in a rFVIII formulation of Embodiment 2, adding the amine-reactive polymer, and incubating the resulting mixture under conditions of time and temperature such that conjugation occurs. Such conditions preferably are at about ambient temperature. The polymer may be added at excess molar amounts (1-100-fold excess) over the FVIII. The polymer and FVIII may be conjugated by incubation together for several hours with rotation or stirring.

[0083] Although the above formulations of Embodiment 2 have been shown to be useful as reaction buffers during polymer addition involving amine-reactive functional groups, it is envisioned that the formulations are also useful in other contexts outside of such reactions and therefore that the formulations may be used when stable FVIII formulations are required.

### EMBODIMENT 3

[0084] In Embodiment 3, the rFVIII formulations comprise NaCl, MOPS, a divalent calcium ion or another divalent cation, and optionally a nonionic surfactant and/or optionally

a sugar or a sugar alcohol. The formulations of Embodiment 3 in particular are shown to provide storage without aggregation of FVIII molecules that are not conjugated to a biocompatible polymer, such as FVIII not covalently attached to PEG and not covalently attached to any polymer other than glycans present in wild-type FVIII. The formulations of Embodiment 3 are particularly suitable for non-PEGylated BDD. As used herein, “nonconjugated FVIII” refers to FVIII that is not conjugated to a polymer other than to a glycan associated with a native mammalian glycosylation pattern resulting from the host cell in which the FVIII is produced. For example, “nonconjugated FVIII” includes wild type human FVIII that is recombinantly produced in a mammalian host cell such as a BHK cell or a CHO cell such as the marketed products KOGENATE® and RECOMBINATE® FVIII.

**[0085]** One version of Embodiment 3 of the compositions described herein is a composition that provides stability for FVIII and contains sodium chloride in a range of from 150 mM to 300 mM, from 150 mM to 275 mM, from 150 mM to 250 mM, from 150 mM to 225 mM, from 150 mM to 200 mM, from 150 mM to 175 mM, from 175 mM to 300 mM, from 175 mM to 275 mM, from 175 mM to 250 mM, from 175 mM to 225 mM, from 175 mM to 200 mM, from 175 mM to 190 mM; from 200 mM to 300 mM, from 200 mM to 275 mM, from 200 mM to 250 mM, from 200 mM to 225 mM, from 200 mM to 210 mM, from 250 mM to 300 mM; or about 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, or 300 mM. The compositions also include MOPS buffer in a range of from 10 mM to 100 mM, in a range of from 10 mM to 60 mM, in a range of from 10 mM to 50 mM, in a range of from 10 mM to 40 mM, in a range of from 10 mM to 30 mM, in a range of from 12 mM to 30 mM, in a range of from 14 mM to 30 mM, in a range of from 16 mM to 30 mM, in a range of from 18 mM to 30 mM, in a range of from 12 mM to 28 mM, in a range of from 12 mM to 26 mM, in a range of from 12 mM to 24 mM, in a range of from 16 mM to 24 mM, in a range of from 18 mM to 24 mM, in a range of from 20 mM to 24 mM, or in a range of from 18 mM to 22 mM. The compositions also include a divalent cation such as calcium chloride in a range of from 1 mM to 20 mM, in a range of from 5 mM to 10 mM, in a range of from 1 mM to 30 mM, in a range of from 6 mM to 30 mM, in a range of from 7 mM to 30 mM, in a range of from 8 mM to 30 mM, in a range of from 5 mM to 20 mM, in a range of from 5 mM to 25 mM, or in a range of from 9 mM to 12 mM. The amount of rFVIII present in the formulations of Embodiment 3 may be the same as the amount provided in Embodiment 1.

**[0086]** The compositions may also include a sugar or sugar alcohol such as sucrose in a range of from 0.5% to 10%, in a range of from 0.6% to 10%, in a range of from 0.7% to

10%, in a range of from 0.8% to 10%, in a range of from 0.9% to 10%, in a range of from 1.0% to 10%, in a range of from 0.5% to 5%, in a range of from 0.6% to 5%, in a range of from 0.7% to 5%, in a range of from 0.8% to 5%, in a range of from 0.9% to 5%, in a range of from 1.0% to 5%, in a range of from 0.5% to 2.5%, in a range of from 0.6% to 2.5%, in a range of from 0.5% to 2.0%, in a range of from 0.5% to 1.5%, in a range of from 0.6% to 1.2%, in a range of from 0.8% to 1.2%, in a range of from 0.9% to 1.2%, or in a range of from 0.9% to 1.1%. The compositions may also include a non-ionic surfactant such as polysorbate 80 in a range of from 20 ppm to 250 ppm, in a range of from 50 ppm to 250 ppm, in a range of from 50 ppm to 150 ppm, in a range of from 60 ppm to 150 ppm, in a range of from 70 ppm to 150 ppm, in a range of from 80 ppm to 150 ppm, in a range of from 60 ppm to 140 ppm, in a range of from 60 ppm to 130 ppm, in a range of from 60 ppm to 120 ppm, in a range of from 60 ppm to 110 ppm, in a range of from 60 ppm to 100 ppm, in a range of from 70 ppm to 110 ppm, in a range of from 70 ppm to 105 ppm, in a range of from 70 ppm to 100 ppm, in a range of from 80 ppm to 100 ppm, or in a range of from 90 ppm to 110 ppm.

[0087] In certain versions of Embodiment 3 the FVIII formulation is free of histidine and/or (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid) ("HEPES") and/or albumin, or contains less than 0.1%, less than 0.5%, less than 0.8%, less than 1.0%, or less than 5.0% by weight of histidine, and/or HEPES, and/or albumin. One version of Embodiment 3 is a FVIII formulation essentially free of histidine, HEPES and albumin.

#### **EMBODIMENT 4**

[0088] Polymer-conjugated FVIII, such as PEGylated FVIII, may be more hydrophilic than the corresponding unconjugated FVIII. Accordingly, formulations for conjugated FVIII such as PEGylated FVIII may require different components than those identified for unconjugated FVIII. Applicants prepared a PEGylated FVIII in a buffer that contained elevated levels of NaCl (200 mM). Such elevated levels of sodium chloride were observed to impose difficulties during lyophilization. Applicants discovered compositions of the present invention for polymer-conjugated FVIII that avoid undesirably high levels of NaCl, avoid the formation of aggregates and substantially retain potency of FVIII when stored over six days at ambient temperature. The present application provides the unexpected result that sodium chloride concentration can be reduced from 200 mM to 50 mM and still achieve potency of the rFVIII after storage at ambient temperature. In Embodiment 4, the rFVIII formulations comprise a buffer such as histidine or MOPS, NaCl, a divalent calcium ion or another divalent cation, and optionally a nonionic surfactant and/or optionally a sugar or a sugar alcohol. The formulations of Embodiment 4 in particular are shown to provide

storage without aggregation of FVIII molecules that are conjugated to a biocompatible polymer, particularly a hydrophilic biocompatible polymer such as PEG. As used herein, “conjugated FVIII” refers to FVIII that is conjugated to a polymer other than to a glycan associated with a native mammalian glycosylation pattern resulting from the host cell in which the FVIII is produced.

**[0089]** One version of Embodiment 4 described herein is a rFVIII composition that contains sodium chloride in a range of from 25 mM to 200 mM, in a range of from 25 mM to 175 mM, in a range of from 25 mM to 150 mM, in a range of from 25 mM to 125 mM, in a range of from 25 mM to 100 mM, in a range of from 25 mM to 75 mM, in a range of from 25 mM to 50 mM, in a range of from 40 mM to 55 mM, in a range of from 25 mM to 35 mM, in a range of from 25 mM to 30 mM, in a range of from 30 mM to 60 mM, in a range of from 50 mM to 200 mM, in a range of from 50 mM to 175 mM, in a range of from 50 mM to 150 mM, in a range of from 50 mM to 125 mM, in a range of from 50 mM to 100 mM, or in a range of from 50 mM to 75 mM. If the formulation is to be subjected to lyophilization, then lower levels of NaCl from those provided above are preferred. The amount of rFVIII present in the formulations of Embodiment 4 may be the same as the amount provided in Embodiment 1.

**[0090]** The compositions also include a buffering agent such as histidine or MOPS buffer in a range of from 10 mM to 100 mM, in a range of from 10 mM to 60 mM, in a range of from 10 mM to 50 mM, in a range of from 10 mM to 40 mM, in a range of from 10 mM to 30 mM, in a range of from 12 mM to 30 mM, in a range of from 14 mM to 30 mM, in a range of from 16 mM to 30 mM, in a range of from 18 mM to 30 mM, in a range of from 12 mM to 28 mM, in a range of from 12 mM to 26 mM, in a range of from 12 mM to 24 mM, in a range of from 16 mM to 24 mM, in a range of from 18 mM to 24 mM, in a range of from 20 mM to 24 mM, or in a range of from 18 mM to 22 mM. The compositions also include a divalent cation such as calcium chloride in a range of from 1 mM to 20 mM, in a range of from 5 mM to 10 mM, in a range of from 1 mM to 30 mM, in a range of from 6 mM to 30 mM, in a range of from 7 mM to 30 mM, in a range of from 8 mM to 30 mM, in a range of from 5 mM to 20 mM, in a range of from 5 mM to 25 mM, or in a range of from 9 mM to 12 mM.

**[0091]** The compositions may also include a sugar or sugar alcohol such as sucrose or trehalose in a range of from 0.5% to 10%, in a range of from 0.6% to 10%, in a range of from 0.7% to 10%, in a range of from 0.8% to 10%, in a range of from 0.9% to 10%, in a range of from 1.0% to 10%, in a range of from 0.5% to 5%, in a range of from 0.6% to 5%, in a range of from 0.7% to 5%, in a range of from 0.8% to 5%, in a range of from 0.9% to 5%,



in a range of from 1.0% to 5%, in a range of from 0.5% to 2.5%, in a range of from 0.6% to 2.5%, in a range of from 0.5% to 2.0%, in a range of from 0.5% to 1.5%, in a range of from 0.6% to 1.2%, in a range of from 0.8% to 1.2%, in a range of from 0.9% to 1.2%, or in a range of from 0.9% to 1.1%. The compositions may also include a non-ionic surfactant such as polysorbate 80 in a range of from 20 ppm to 250 ppm, in a range of from 50 ppm to 250 ppm, in a range of from 50 ppm to 150 ppm, in a range of from 60 ppm to 150 ppm, in a range of from 70 ppm to 150 ppm, in a range of from 80 ppm to 150 ppm, in a range of from 60 ppm to 140 ppm, in a range of from 60 ppm to 130 ppm, in a range of from 60 ppm to 120 ppm, in a range of from 60 ppm to 110 ppm, in a range of from 60 ppm to 100 ppm, in a range of from 70 ppm to 110 ppm, in a range of from 70 ppm to 105 ppm, in a range of from 70 ppm to 100 ppm, in a range of from 80 ppm to 100 ppm, or in a range of from 90 ppm to 110 ppm.

**[0092]** In certain versions of Embodiment 4 the FVIII formulation is free of histidine and/or HEPES and/or albumin, or contains less than 0.1%, less than 0.5%, less than 0.8%, less than 1.0%, or less than 5.0% by weight of histidine, and/or HEPES, and/or albumin. One version of Embodiment 3 is a FVIII formulation essentially free of histidine, HEPES and albumin.

#### **EMBODIMENT 5**

**[0093]** The invention also includes rFVIII formulations suitable for lyophilization. In certain versions of this embodiment, the FVIII formulations are particularly suitable for lyophilization of conjugated FVIII, PEGylated FVIII, PEGylated BDD, or PEGylated BDD mutants. The rFVIII formulations of this embodiment comprise (1) sodium chloride, and/or sucrose, and/or trehalose, (2) glycine and/or sucrose and/or trehalose; and (3) a divalent cation such as calcium chloride, and optionally contain (1) a nonionic surfactant, and/or (2) histidine, and if NaCl is present, then optionally also a sugar or a sugar alcohol, including without limitation sucrose and/or trehalose.

**[0094]** The invention includes formulations of Embodiment 5 as follows. A rFVIII formulation comprising:

- (a) about 0mM, or a range of from about 1mM to about 20mM, from about 1 mM to about 50 mM, from about 10 mM to about 50 mM, from about 10 mM to about 20mM, from about 10 mM to about 30mM, or from about 20 mM to about 50 mM histidine;
- (b) a range of from 0.5% to 20%, a range of from 1.0% to 20%, a range of from 0.6% to 10%, a range of from 0.7% to 10%, a range of from 0.8% to 10%, a

range of from 0.9% to 10%, a range of from 1.0% to 10%, a range of from 0.5% to 5%, a range of from 0.6% to 5%, a range of from 0.7% to 5%, a range of from 0.8% to 5%, a range of from 0.9% to 5%, a range of from 1.0% to 5%, a range of from 0.5% to 2.5%, a range of from 0.6% to 2.5%, a range of from 0.5% to 2.0%, a range of from 0.5% to 1.5%, a range of from 0.6% to 1.4%, a range of from 0.8% to 1.4%, a range of from 0.9% to 1.2%, a range of from 3.0% to 9.0%, a range of from 5.0% to 9.0%, a range of from 6.0% to 8.0%, a range of from 7.0% to 9.0%, or a range of from 0.9% to 1.1%, or about 1.0%, 1.1%, 1.2%, 1.3%, 1.4%, 1.5%, 1.6%, 1.7%, 1.8%, 1.9%, 2.0%, 3.0%, 4.0%, 5.0%, 6.0%, 7.0%, 7.5%, 8.0%, 8.5%, 9.0%, 9.5%, 10.0%, 12.0%, or 15.0% of sucrose or trehalose;

- (c) a range of from about 1 mM to about 5 mM, from about 1 mM to about 3 mM, from about 1.5 mM to about 3.5 mM, or from about 1 mM to about 2.5 mM divalent cation such as a divalent calcium salt, including calcium chloride;
- (d) about 0 mM, or a range of from about 10 mM to about 50 mM, from about 10 mM to about 40 mM, from about 10 mM to about 35 mM, from about 10 mM to about 30 mM; from about 10 mM to about 20 mM, from about 20 mM to about 50 mM, from about 20 mM to about 40 mM, or from about 20 mM to about 80 mM sodium chloride;
- (e) about 0 mM, or a range of from about 20 ppm to about 50 ppm, from about 20 ppm to about 80 ppm, from about 50 ppm to about 80 ppm, from about 80 ppm to about 100 ppm, from about 80 ppm to about 200 ppm, or from about 50 ppm to about 100 ppm of a non-ionic surfactant, or about 60, 65, 70, 75, 80, 85, 90, 95, 100, 105, 110, 115, 120, 125, 130, 135, 140, 145, 150, 160, 170, 180, 190, or 200 ppm of a non-ionic surfactant;
- (f) about 0%, or a range of from about 1.0% to about 5.0%, a range of from about 1.0% to about 4.0%, a range of from about 1.0% to about 3.0%, a range of from about 1.0% to about 2.0%, a range of from about 1.0% to about 1.5%, a range of from about 1.0% to about 1.4%, a range of from about 0.5% to about 5.0%, a range of from about 0.5% to about 4.0%, a range of from about 0.5% to about 3.0%, a range of from about 0.5% to about 2.0%, a range of from about 0.5% to about 1.5% glycine, or about 1.5%, 1.8%, 2.0%, 2.1%, 2.2%, 2.3%, 2.4%, 2.5%, 2.6%, 2.7%, 2.8%, 2.9%, 3.0%, 3.3%, 3.5%, or 4.0% glycine and

(g) a range of from about 100 IU/ml to about 5000 IU/ml, from about 100 IU/ml to about 2000 IU/ml, from about 100 IU/ml to about 3000 IU/ml, from about 100 IU/ml to about 4000 IU/ml, from about 100 IU/ml to about 1200 IU/ml, from about 250 IU/ml to about 5000 IU/ml, from about 250 IU/ml to about 1000 IU/ml, from about 250 IU/ml to about 2000 IU/ml, from about 250 IU/ml to about 3000 IU/ml, from about 500 IU/ml to about 1000 IU/ml, from about 500 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 2000 IU/ml, from about 1000 IU/ml to about 3000 IU/ml, from about 1000 IU/ml to about 4000 IU/ml, or from about 1000 IU/ml to about 5000 IU/ml, or about 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 1200, 1400, 1600, 1800, 2000, 2200, 2400, 2600, 2800, 3000, 3200, 3500, 3800, 4000, 4200, 4500, 4600, 4800, 5000, 5500, or 6000 IU/ml of rFVIII;

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 6.5, from about pH 6.0 to about pH 7.0, from about pH 6.0 to about pH 7.5, from about pH 6.5 to about pH 7.5, or from about pH 7.0 to about pH 7.5, or a pH of about pH 6.0, 6.5, 7.0, 7.1, 7.2, 7.3, 7.4 or about pH 7.5.

[0095] In one version of Embodiment 5, the rFVIII formulation comprises sodium chloride and contains less than 2.0% sucrose or sucrose in a range of from 0.5% to 2.0%, and contains less than 1.0%, less than 0.5%, less than 0.1% or no trehalose. In this version, NaCl may be present at a range of from about 10 mM to about 50 mM, from about 10 mM to about 40 mM, from about 10 mM to about 35 mM, from about 10 mM to about 30 mM; from about 10 mM to about 20 mM, from about 20 mM to about 50 mM, from about 20 mM to about 40 mM, or from about 20 mM to about 80 mM sodium chloride. In this version of Embodiment 5, glycine is present at a range of from about 1.0% to about 5.0%, a range of from about 1.0% to about 4.0%, a range of from about 1.0% to about 3.0%, a range of from about 1.0% to about 2.0%, a range of from about 1.0% to about 1.5%, a range of from about 1.0% to about 1.4%, a range of from about 0.5% to about 5.0%, a range of from about 0.5% to about 4.0%, a range of from about 0.5% to about 3.0%, a range of from about 0.5% to about 2.0%, a range of from about 0.5% to about 1.5%, or at about 1.5%, 1.8%, 2.0%, 2.1%, 2.2%, 2.3%, 2.4%, 2.5%, 2.6%, 2.7%, 2.8%, 2.9%, 3.0%, 3.3%, 3.5%, or 4.0% and sucrose is present at a range of from 0.5% to 5%, a range of from 0.6% to 5%, a range of from 0.7% to 5%, a range of from 0.8% to 5%, a range of from 0.9% to 5%, a range of from 1.0% to 5%, a range of from 0.5% to 2.5%, a range of from 0.6% to 2.5%, a range of from 0.5% to 2.0%, a range of from 0.5% to 1.5%, a range of from 0.6% to 1.4%, a range of from 0.8% to 1.4%, a range of from

0.9% to 1.2%, or a range of from 0.9% to 1.1%, or about 1.0%, 1.1%, 1.2%, 1.3%, 1.4%, 1.5%, 1.6%, 1.7%, 1.8%, 1.9%, 2.0%, 3.0%, 4.0% sucrose. In this version of Embodiment 5, histidine is present at a range of from about 1 mM to about 20mM, from about 1 mM to about 50 mM, from about 10 mM to about 50 mM, from about 10 mM to about 20mM, from about 10 mM to about 30mM, or from about 20 mM to about 50 mM and a non-ionic surfactant such as polysorbate 20 or polysorbate 80 is present at a range of from about 20 ppm to about 50 ppm, from about 20 ppm to about 80 ppm, from about 50 ppm to about 80 ppm, from about 80 ppm to about 100 ppm, from about 80 ppm to about 200 ppm, or from about 50 ppm to about 100 ppm of a non-ionic surfactant, or about 60, 65, 70, 75, 80, 85, 90, 95, 100, 105, 110, 115, 120, 125, 130, 135, 140, 145, 150, 160, 170, 180, 190, or 200 ppm. In this version of Embodiment 5, trehalose is present at less than 1.0%, less than 0.5%, less than 0.1% by weight or is not present.

[0096] In another version of Embodiment 5, sodium chloride is present at less than 1.0%, less than 0.5%, less than 0.1% by weight or is not present. In this version, sucrose or trehalose is present a range of from 0.5% to 20%, a range of from 1.0% to 20%, a range of from 0.6% to 10%, a range of from 0.7% to 10%, a range of from 0.8% to 10%, a range of from 0.9% to 10%, a range of from 1.0% to 10%, a range of from 3.0% to 9.0%, %, a range of from 5.0% to 9.0%, a range of from 6.0% to 8.0%, %, or a range of from 7.0% to 9.0%, or about 5.0%, 6.0%, 7.0%, 7.5%, 8.0%, 8.5%, 9.0%, 9.5%, 10.0%, or 12.0%. In this version of Embodiment 5, glycine is present at less than 1.0%, less than 0.5%, less than 0.1% by weight or is not present.

[0097] Aspects of the present disclosure may be further understood in light of the following examples, which should not be construed as limiting the scope of the present teachings in any way.

### EXAMPLES

#### Example 1: Effect of Sodium Chloride, Polysorbate 80, and Human Serum Albumin on BDD-rFVIII Protein Solubility and Stability

##### Effect of Sodium Chloride

[0098] Studies were performed on BDD mutants having introduced cysteine residues that permit the stabilization of FVIII by formation of at least one disulfide bond between different domains of FVIII. In particular, BDD-SQ (SEQ ID NO: 3) was mutated at Tyr664Cys:Thr1826Cys to create the C664-BDD mutant used in this example. For methods of preparation, see U.S. Patent 7,928,199 (Griffin et al.). When the C664-BDD mutant was

formulated in a buffer containing histidine, unacceptable levels of precipitation were observed.

[0099] A study was performed to determine whether the precipitation observed when the C664-BDD mutant was placed in histidine buffer could be reversed. The buffer solution in which precipitation was observed contained 20 mM histidine, 30 mM sodium chloride, 2.5 mM calcium chloride, 29 mM sucrose, 293 mM glycine and 80 ppm polysorbate 80. The C664 BDD mutant was present at 145 IU/ml. The aim of the study was to develop a formulation that stabilizes BDD-rFVIII mutants. Solubilizers and stabilizers, such as sodium chloride, Polysorbate 80, and human serum albumin (HSA) were tested to either increase the solubility of the mutants or to improve the stability by reducing protein aggregation. Results are shown in FIGs. 2-6. The experiments shown in FIGs. 2 and 5 both involved modification of the NaCl concentration, and the results in each instance showed remarkable turbidity decline from a solution containing 30 mM NaCl when compared to a solution containing about 120 mM NaCl. The study established that as the sodium chloride concentration increased, the turbidity of the solution comprising the mutants decreased, suggesting that sodium chloride reversed the precipitation process. When the sodium chloride concentration was 176 mM or higher, the cloudy solution turned to a clear solution and the turbidity dropped from 0.169 to 0.029, which is more than 80% based on  $A_{340\text{ nm}}$  measurements (FIG. 2). These results demonstrated that sodium chloride was an effective solubilizer for the BDD-rFVIII mutants and can reverse their precipitation. In summary, higher sodium chloride concentrations improved the solubility of the BDD-rFVIII mutants. Table 2 shows preferred formulations. "BDD-rFVIII mutants" in Table 2 refers to a formulation of BDD-SQ mutated at Tyr664Cys:Thr1826Cys. "Full-length rFVIII" in Table 2 refers to a formulation of FVIII that has the amino acid sequence of SEQ ID NO: 2 (full-length FVIII).

Table 2

*Formulation Composition for full-length rFVIII and BDD-rFVIII mutants*

Composition	BDD-rFVIII mutants	Full-length rFVIII
Sodium chloride (mM)	220	30
Sucrose (mM)	29	29
Histidine (mM)	20	20

Glycine (mM)	293	293
Calcium chloride (mM)	2.5	2.5
Polysorbate 80 (ppm)	80	80

Example 2: Formulation Development for rFVIII PEGylation through Random Lysine Coupling

**[0100]** PEG polymer was conjugated to the full-length rFVIII of SEQ ID NO: 1 using random lysine coupling. In this type of coupling, the reactive groups are primarily the N-terminal amine or the  $\epsilon$ -amino group of lysine in a protein. Other primary or secondary amine groups in the formulation could interfere with the reaction. Because many full-length and BDD-rFVIII formulations comprise amino acids, such as glycine and histidine, new formulations were developed for PEGylation of these molecules. While glycine was used as a bulking agent in the full-length rFVIII formulation and could be eliminated during PEGylation, histidine served as a buffer component and needed to be replaced with another buffer.

**[0101]** A suitable buffer system meets the following criteria: (1) it provides buffer capacity at pH 6-7; (2) it does not form insoluble complex or chelate with calcium chloride, an important rFVIII stabilizer; and (3) it does not comprise primary or secondary amine groups.

**[0102]** Several commonly used buffers were considered for random PEGylation of rFVIII. As shown in Table 3, only two buffer systems, tri-ethanolamine ("TEA") and MOPS were selected for further investigation.

Table 3

Buffers Considered for Random PEGylation of rFVIII

Buffer at pH 7	Ca <sup>2+</sup> ppt.	Ca <sup>2+</sup> chelating	Amine group	pH change during freezing
Citrate		X		
Phosphate	X			X

Buffer at pH 7	Ca <sup>2+</sup> ppt.	Ca <sup>2+</sup> chelating	Amine group	pH change during freezing
Histidine			X	
TRIS			X	
Carbonate	X			
Triethanolamine (TEA)				
MOPS or MOPSO				
HEPES			X	

[0103] For this study, full-length rFVIII was dialyzed against the formulations listed in Table 4. The dialyzed rFVIII in the three formulations was placed at 40°C (Figure 7) or 25°C (Figure 8) to establish stability at accelerated conditions and the results are shown in Figures 7 and 8.

Table 4

*Buffers Evaluated for Random PEGylation of rFVIII*

	<i>NaCl</i> (mM)	<i>CaCl<sub>2</sub></i> (mM)	<i>Tween 80</i> (ppm)	<i>Glycine</i> (mM)	<i>Sucrose</i> (mM)	<i>Sodium Azide</i> (%)	<i>Buffer Agent</i> (20mM)
1	30	2.5	80	--	29	0.05	TEA
2	30	2.5	80	--	29	0.05	MOPS
3	30	2.5	80	293	29	0.05	Histidine

*Example 3: PEGylation for BDD-rFVIII*

[0104] BDD-rFVIII encounters formulation challenges due to its propensity for aggregation. Therefore, one of the objectives with designing a formulation for PEGylated rFVIII was to ensure its stability in solution. The working formulation for the PEGylated BDD-rFVIII comprised 200 mM sodium chloride, 20 mM MOPS, 10 mM CaCl<sub>2</sub>, 100 ppm polysorbate 80 and 29 mM sucrose. 200 mM sodium chloride will impose difficulties during freeze-drying. Accordingly, the solubility and potency of the PEGylated BDD-rFVIII were evaluated as a function of sodium chloride concentration in the range of 50 and 250 mM.

[0105] The buffer composition used for the study is shown in Table 5 and the data are summarized in Figures 10 and 11. The PEGylated BDD used in this example comprised the amino acid sequence of SEQ ID NO: 3 with one amino acid mutation to create a free cysteine at which PEG was added. This is shown graphically in FIG. 9. The PEGylated BDD-rFVIII retained more than 87% potency in the formulation comprising 50-150 mM sodium chloride during 6 days storage at 23°C. UnPEGylated BDD-rFVIII retained 70% potency in the same formulation during 6 days storage at 23°C. Both molecules remained soluble during the study with no visual detection of precipitates or opalescence. These and earlier data suggest that 100 mM sodium chloride can be used for further formulation development.

Table 5

*Composition of the Formulation Used for Evaluating the Effect of Sodium Chloride*

MOPS (mM)	NaCl (mM)	CaCl <sub>2</sub> (mM)	Polysorbate 80 (ppm)	Sucrose (mM)
20	250	10	100	29
20	200	10	100	29
20	150	10	100	29
20	100	10	100	29
20	50	10	100	29
20	25	10	100	29
20	0	10	100	29



[0106] The effect of sodium chloride on the solubility and aggregation of PEGylated and unPEGylated BDD-rFVIII was investigated.

[0107] UV absorbance of PEGylated BDD-rFVIII in MOPS buffer comprising 25 mM, 55 mM, 75 mM, 125 mM and 200 mM sodium chloride showed no scattering of the PEGylated BDD-rFVIII at all sodium chloride concentration tested, suggesting lack of aggregation. In contrast, the unPEGylated-rFVIII showed considerable scattering at 25 mM, 55 mM and 75 mM sodium chloride most likely due to formation of soluble aggregates. When sodium chloride concentration was increased to 125 mM and 200 mM, no scattering was observed. It was concluded, therefore, that higher salt concentrations prevented aggregate formation.

Example 4: Development of Freeze-Drying Formulation for PEGylated BDD-rFVIII

[0108] Four candidate formulations were screened for lyophilization of PEGylated BDD-rFVIII. The PEGylated BDD used in this example comprised the amino acid sequence of SEQ ID NO: 3 with one amino acid mutation to create a free cysteine at which PEG was added. The aim was to evaluate the stability of the lyophilized drug product in these formulations and to select a formulation for the leading stability study. The formulations that were screened were (1) Formulation A, which had been successful for unPEGylated full-length rFVIII, (2) Formulation B, comprising increased solids content compared to Formulation A, (3) Formulation C with sucrose instead of the NaCl used in Formulation A, and (4) Formulation D with trehalose instead of the NaCl used in Formulation A. The last two formulations provided an amorphous matrix for the lyophilized drug product.

[0109] Stability was evaluated at three storage temperatures (5°C, 25°C and 40°C). Table 6 shows the formulation composition for PEGylated BDD-rFVIII used for stability evaluation.

[0110] The concentrations of sucrose and glycine were increased from 29 mM and 293 mM in Formulation A to 38 mM and 346 mM in Formulation B. The additional solids were added to enhance the mechanical strength of the freeze-dried cake and improve the appearance of the final drug product.

Table 6:

*Formulation Composition for PEGylated BDD-rFVIII Used in Stability Evaluation*

Component	Formulation A	Formulation B	Formulation C	Formulation D
Calcium Chloride	2.5 mM	2.5 mM	2.5 mM	2.5 mM
Sodium Chloride	30 mM	30 mM	X	X
Histidine	20 mM	20 mM	20 mM	20 mM
Glycine	293 mM	346 mM	X	X
Polysorbate 80	80 ppm	80 ppm	80 ppm	80 ppm
Sucrose	29 mM	38 mM	234 mM	X
Trehalose	X	X	X	211 mM
PEGylated BDD-rFVIII concentration (IU/mL)	100 IU/mL	100 IU/mL	100 IU/mL	100 IU/mL

1: pH = 6.8 for all formulations

[0111] Formulations C and D were designed to provide an alternate matrix compared to the other two formulations. Formulations A and B formed a crystalline matrix upon freeze-drying due to the presence of sodium chloride and glycine as structural stability and bulking agents. The concentrations of sucrose and trehalose were increased to 234 mM and 211 mM, respectively, in lieu of including sodium chloride and glycine. This resulted in an amorphous matrix for the freeze-dried drug product.

[0112] The stability program for each of the four candidate formulations was set up for a 26 week time period. Stability was evaluated by potency, moisture content, percent high molecular weight (HMW) impurities and total product related impurities by SEC-HPLC. The potency recovery data for the four formulations are summarized in Figures 12-15.

[0113] The data of potency recovery, moisture content by Karl Fischer, and percent aggregates and product related impurities by SEC-HPLC (tested at 26 weeks) for the four formulations demonstrate that rFVIII is stable in the four formulations.

[0114] Stability for PEGylated BDD-rFVIII was further evaluated with Formulations A and B (see Table 6 for formulation composition). Two drug product lots were prepared at lab-scale and were placed on stability at 5°C and 25°C and 40°C. Potency by the chromogenic assay, percent high molecular weight impurities and total product related impurities by SEC-HPLC, and moisture by Karl Fischer were employed for drug product stability evaluation. Target concentrations and ranges of the components used in Formulation A are presented in Table 7.

Table 7: Target Concentrations and Ranges of the Components Used in the Formulation A

Component	Formulation A Target Concentrations	Low and High Concentration Range
Calcium Chloride	2.5 mM	1.5 mM to 3.5 mM
Sodium Chloride	30 mM	21 mM to 43 mM
Histidine	20 mM	15 mM to 27 mM
Glycine	293 mM	240 mM to 386 mM
Polysorbate 80	80 ppm	57 ppm to 103 ppm
Sucrose	29 mM	20 mM to 41 mM
PEGylated BDD-rFVIII concentration (IU/mL)	200 IU/mL 400 IU/mL 1200 IU/mL	188 IU/mL to 250 IU/mL 376 IU/mL to 500 IU/mL 1128 IU/mL to 1500 IU/mL

[0115] These data demonstrated comparable drug product stability in the two formulations. The study with Formulation A was continued up to 30 months, whereas the study with Formulation B was terminated at 3 months (Figures 16 and 17, respectively). rFVIII concentration in Figure 18 and 19 was 400 IU/mL.

[0116] Formulation A was selected for further development and was tested with PEGylated rFVIII at concentrations of 200 IU/mL and 1200 IU/mL. The potency profiles at 200 IU/mL and 1200 IU/mL are shown in Figures 18 and 19, respectively. The data demonstrate that Formulation A provides continuous stability for the PEGylated rFVIII.

**[0117]** Throughout this specification and the claims which follow, unless the context requires otherwise, the word "comprise", and variations such as "comprises" or "comprising", will be understood to imply the inclusion of a stated integer or step or group of integers or steps but not the exclusion of any other integer or step or group of integers or steps.

**[0118]** The reference in this specification to any prior publication (or information derived from it), or to any matter which is known, is not, and should not be taken as an acknowledgment or admission or any form of suggestion that that prior publication (or information derived from it) or known matter forms part of the common general knowledge in the field of endeavour to which this specification relates.

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**The claims defining the invention are as follows:**

1. A rFVIII formulation comprising:
  - (a) a range of from about 1 mM to about 5 mM divalent cation;
  - (b) a range of from about 150 mM to about 250 mM sodium chloride or potassium chloride;
  - (c) a range of from about 50 ppm to about 200 ppm of a non-ionic surfactant; and
  - (d) a range of from about 100 IU/ml to about 5000 IU/ml of a rFVIII, wherein the rFVIII comprises an amino acid sequence that has one or more non-cysteine residues in the amino acid sequence of SEQ ID NO: 3 replaced with cysteine residues such that at least one pair of cysteine residues creates a disulphide bond not found in wild type FVIII;

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5.
2. The rFVIII formulation of claim 1 further comprising:
  - (a) a range of from about 10 mM to about 50 mM histidine;
  - (b) a range of from about 10 mM to about 100 mM of a sugar or sugar alcohol; and
  - (c) a range of from about 150 mM to about 400 mM glycine.
3. A rFVIII formulation comprising:
  - (a) about 0 mM, or a range of from about 1 mM to about 20 mM histidine;
  - (b) a range of from 0.5% to 20% of sucrose or trehalose;
  - (c) a range of from about 1 mM to about 5 mM divalent cation;
  - (d) a range of from about 10 mM to about 50 mM sodium chloride;
  - (e) about 0 mM, or a range of from about 20 ppm to about 80 ppm of a non-ionic surfactant;
  - (f) about 0%, or a range of from about 1.0% to about 5.0% glycine; and
  - (g) a range of from about 100 IU/ml to about 5000 IU/ml of conjugated rFVIII;

wherein the rFVIII formulation has a pH in a range of from about pH 6.0 to about pH 7.5.
4. The rFVIII formulation of claim 3, wherein
  - (a) sodium chloride is present in a range of from about 10 mM to about 50 mM;
  - (b) sucrose is present in a range of from 0.5% to 2.0%;

- (c) glycine is present in a range of from about 1.0% to about 5.0%;
- (d) histidine is present in a range of from about 1mM to about 20mM; and
- (e) a non-ionic surfactant is present in a range of from about 20 ppm to about 80 ppm.

5. The rFVIII formulation of claim 3, wherein

- (a) sodium chloride is present at less than 1.0% by weight;
- (b) sucrose or trehalose is present in a range of from 0.5% to 20%; and
- (c) glycine is present at less than 1.0% by weight or is not present.

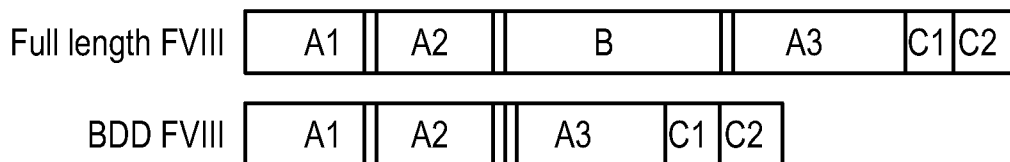
6. The rFVIII formulation of claim 5, wherein sucrose or trehalose is present in a range of from 1.0% to 10.0%.

7. A method of treating hemophilia A comprising administering a therapeutically effective amount of a rFVIII formulation of any one of claims 1 to 6 to a patient in need thereof.

8. Use of a rFVIII formulation of any one of claims 1 to 6 in the manufacture of a medicament for treating hemophilia A.

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## FIG. 1



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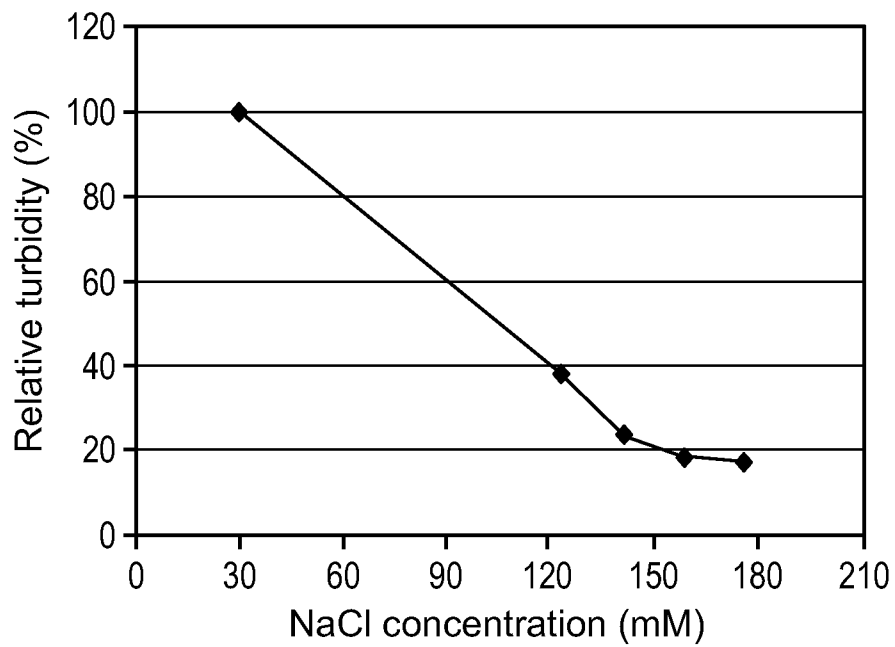


FIG. 2

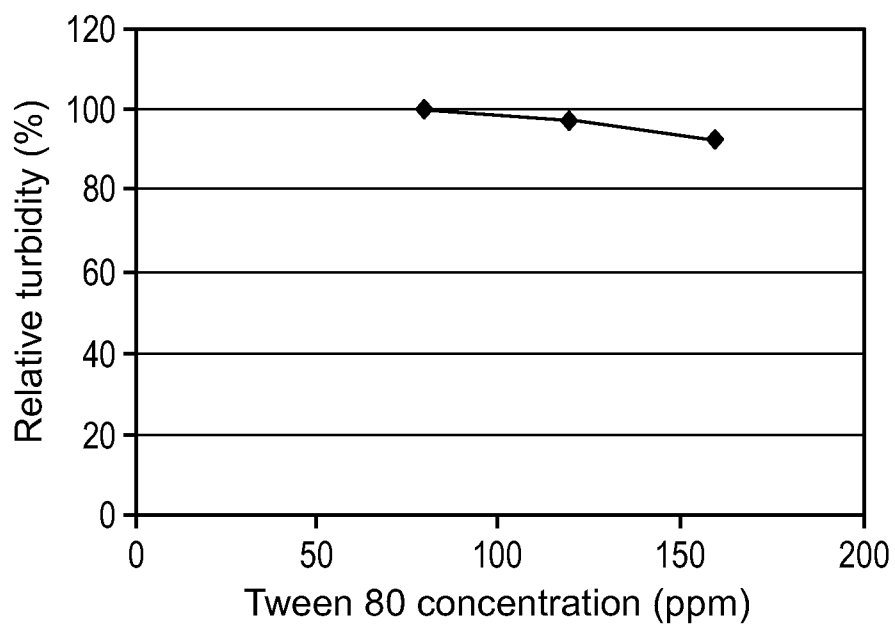


FIG. 3



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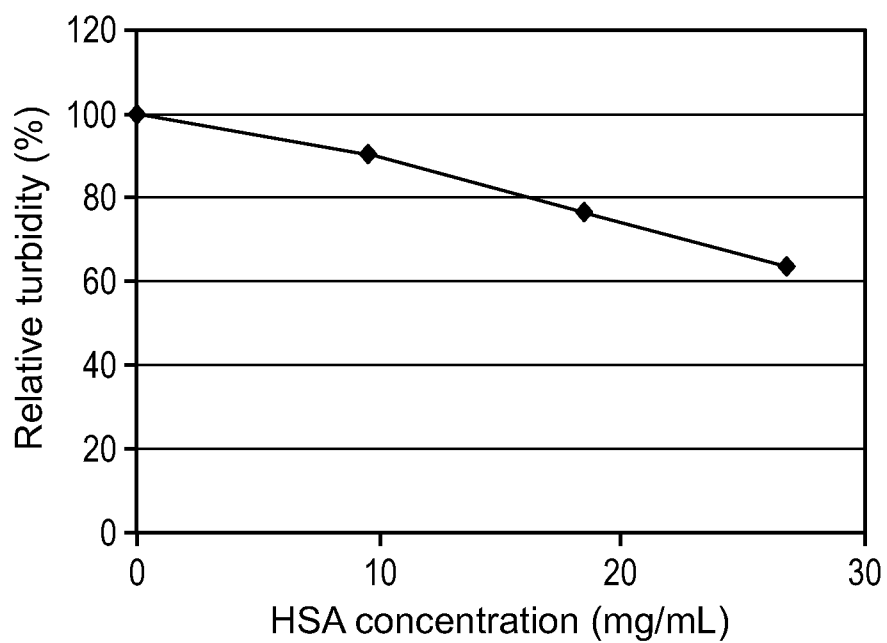


FIG. 4

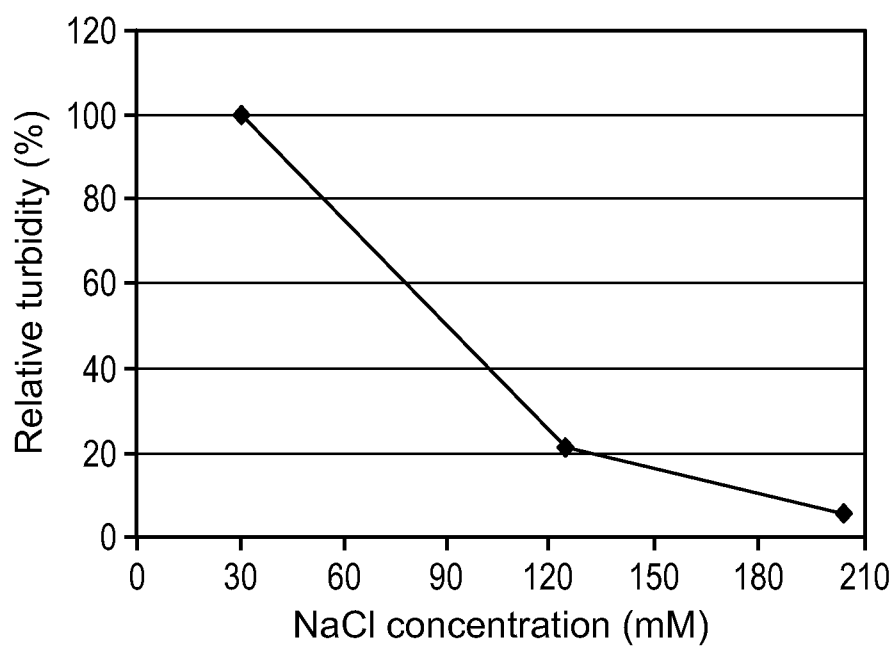


FIG. 5

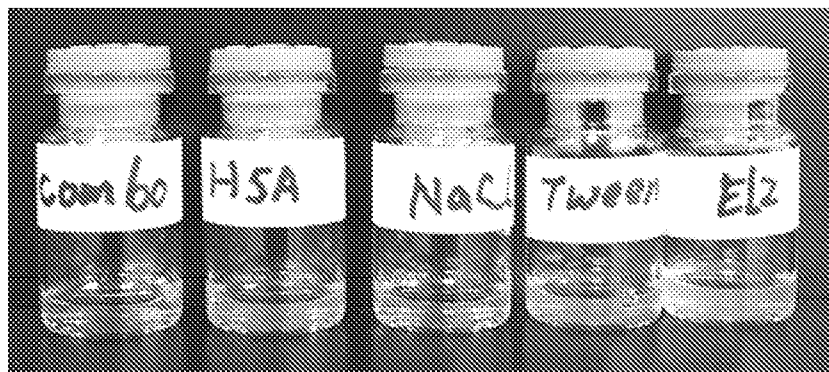


FIG. 6

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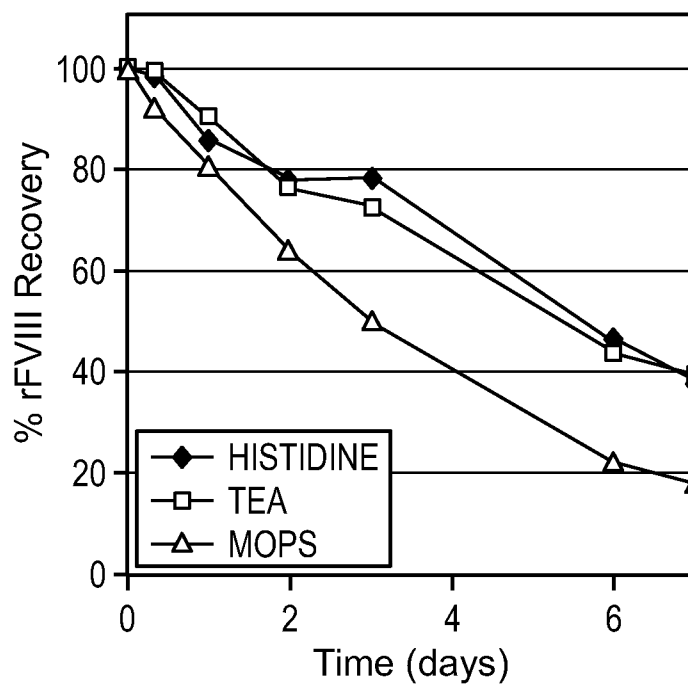


FIG. 7

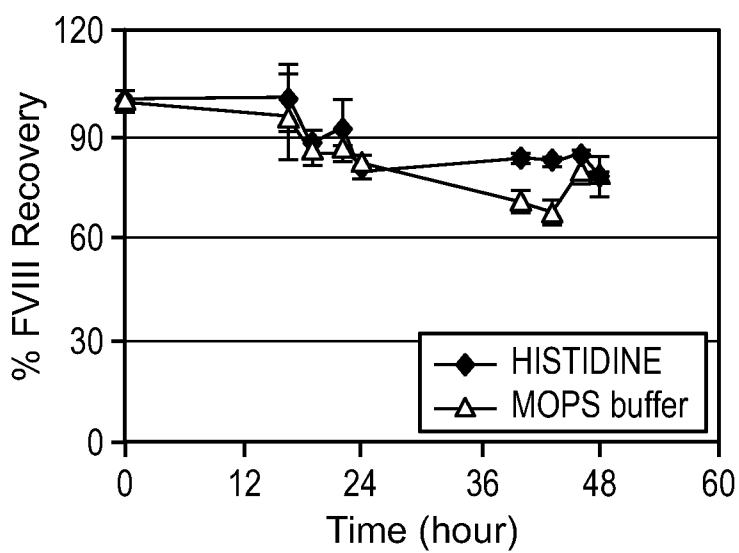
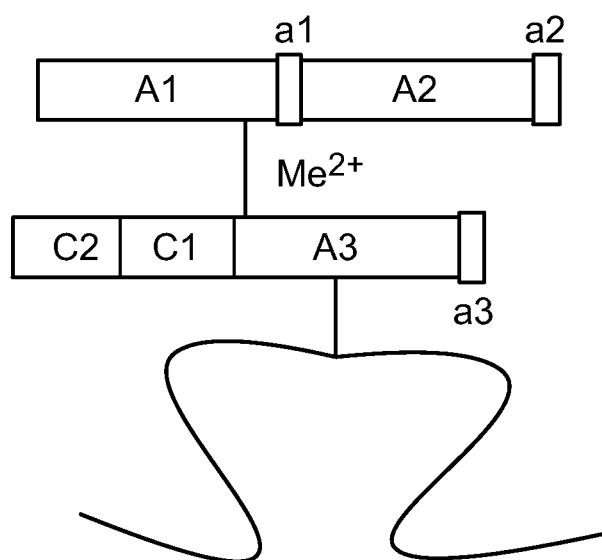


FIG. 8

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FIG. 9



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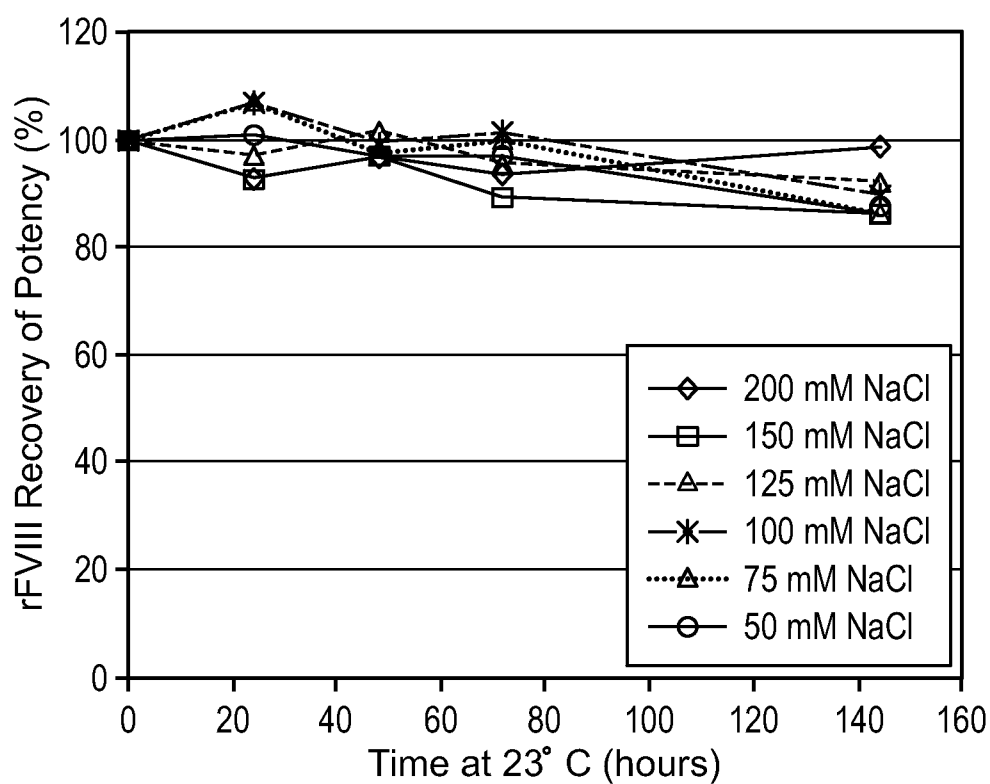


FIG. 10

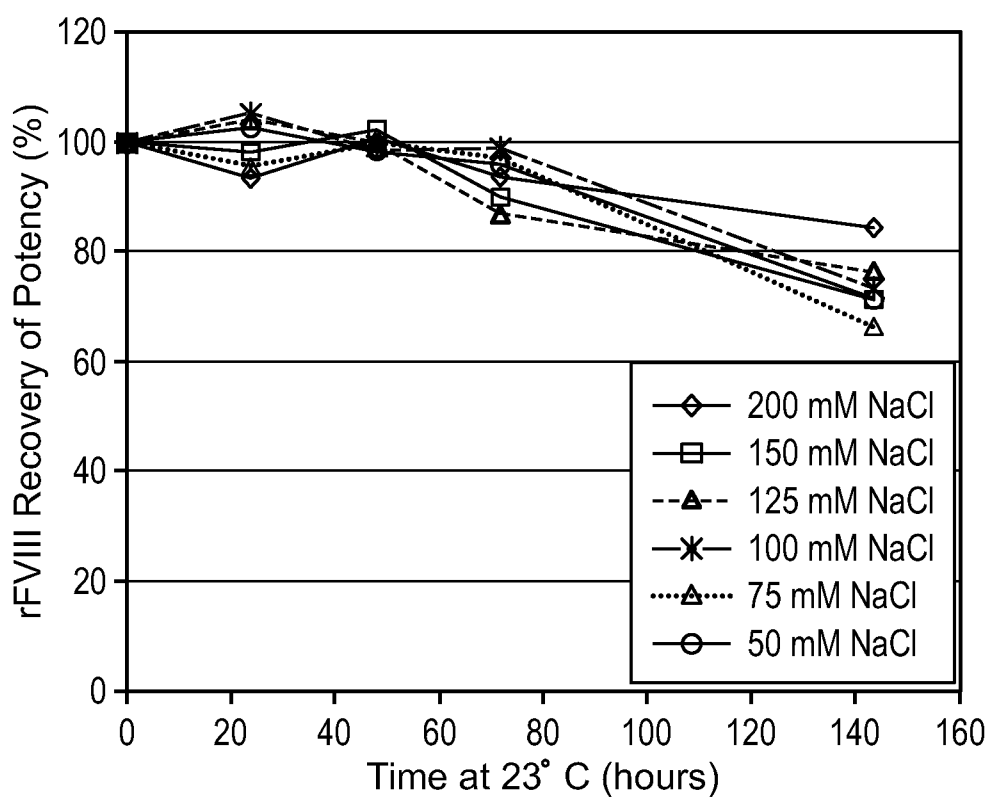


FIG. 11

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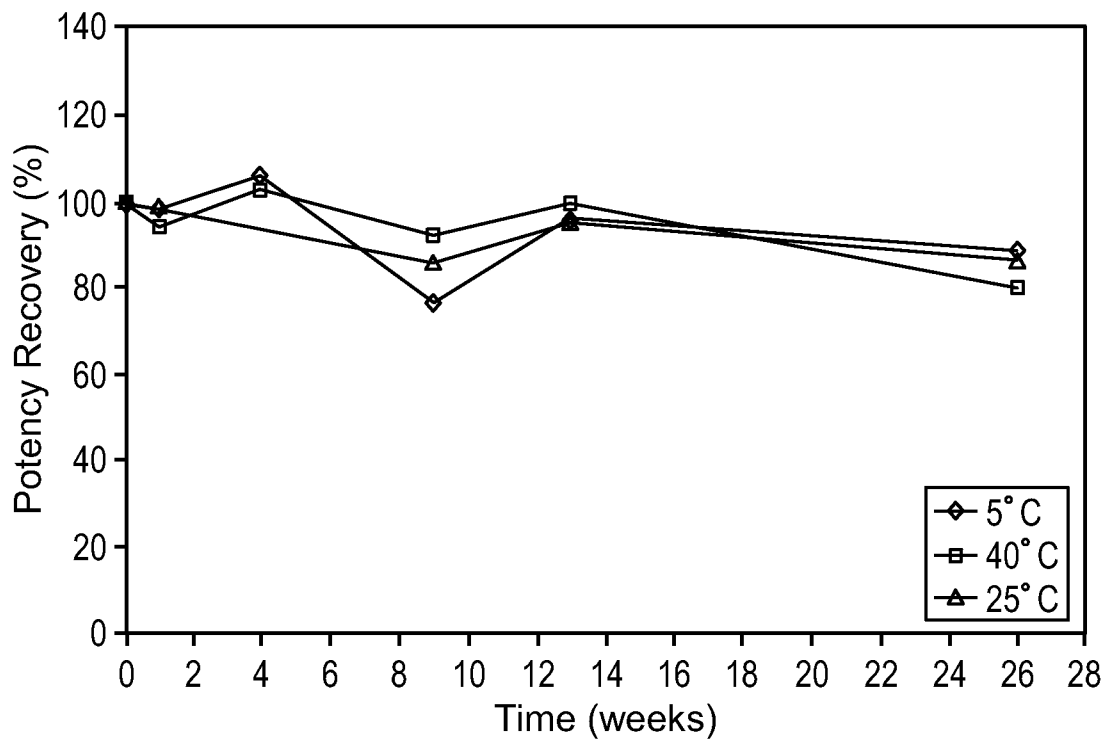


FIG. 12

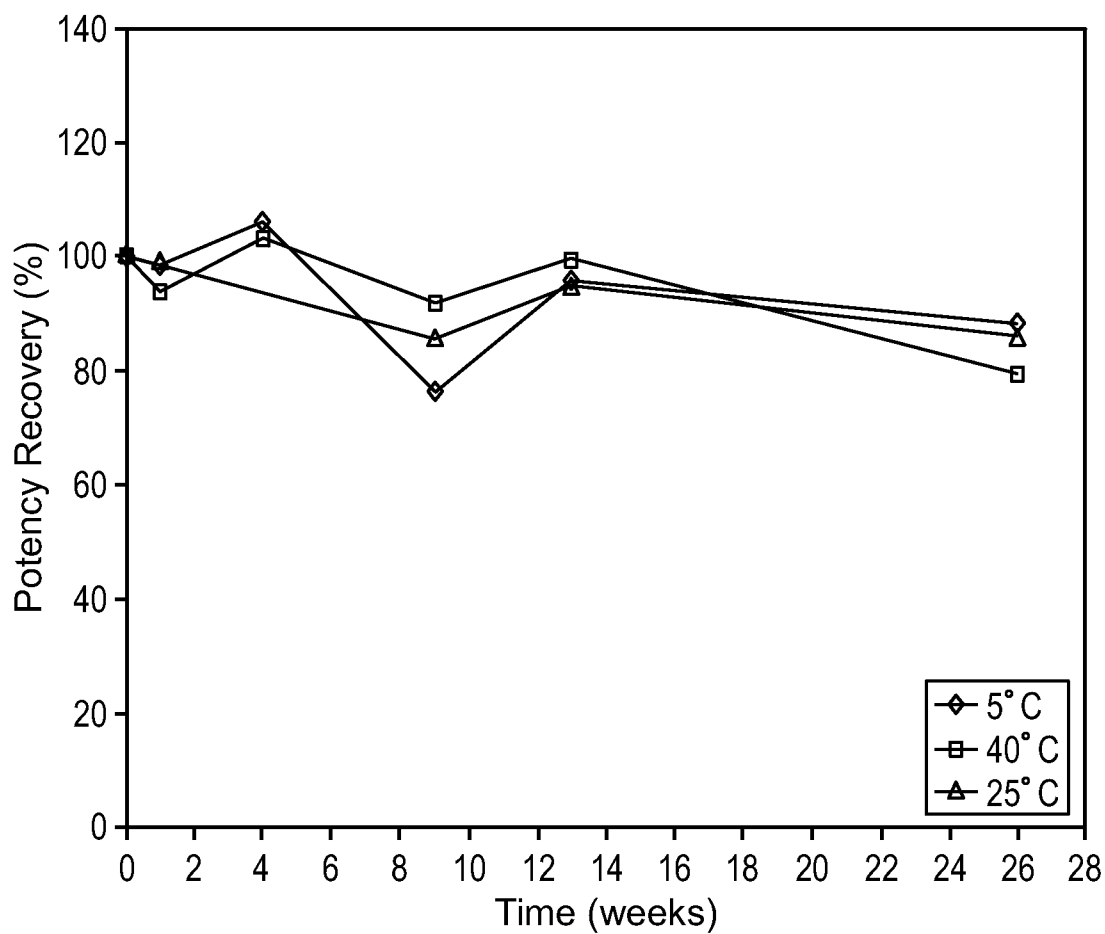


FIG. 13

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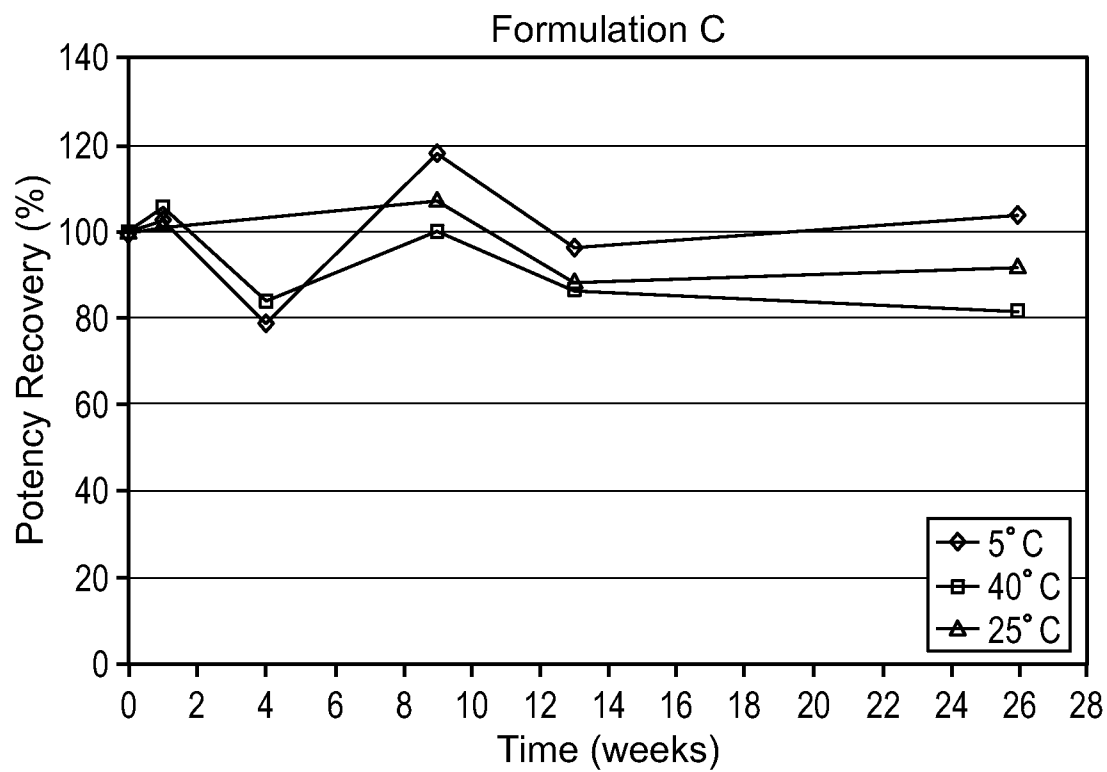


FIG. 14

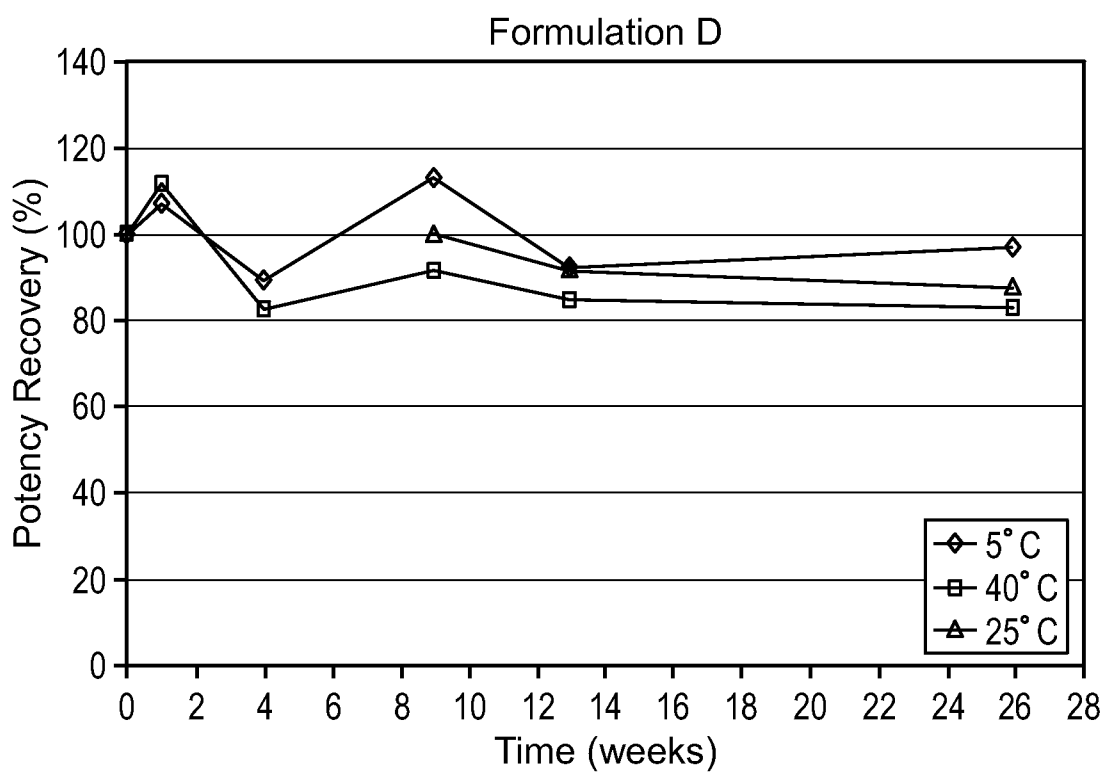


FIG. 15

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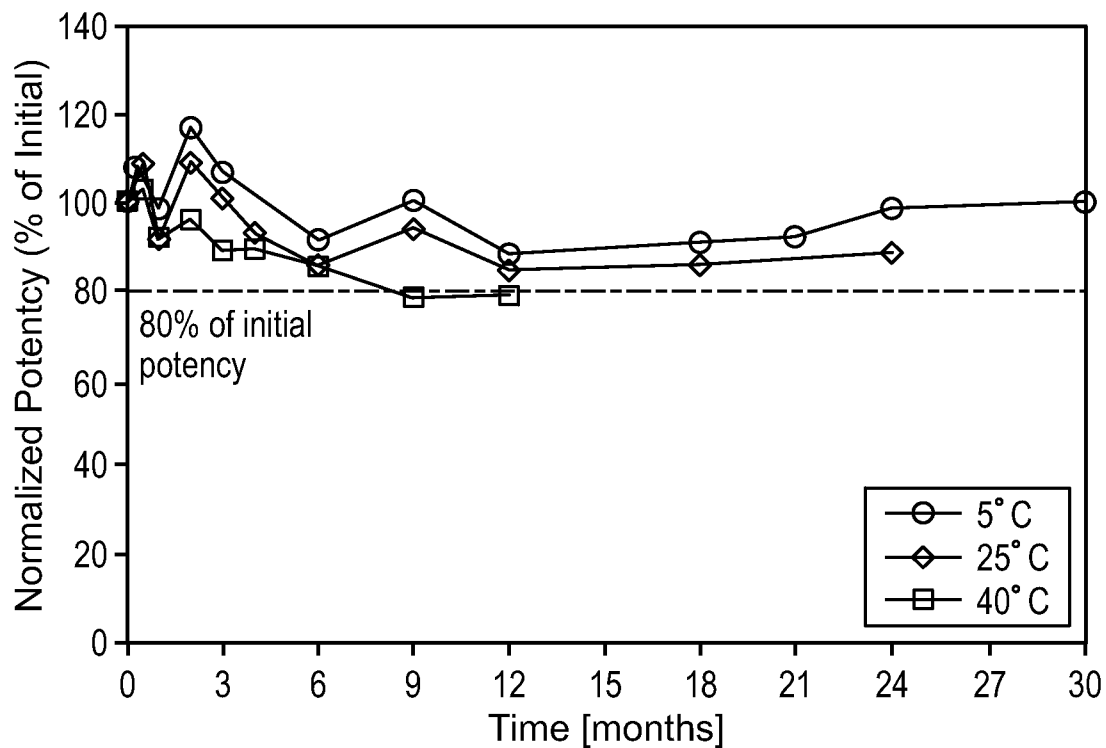


FIG. 16

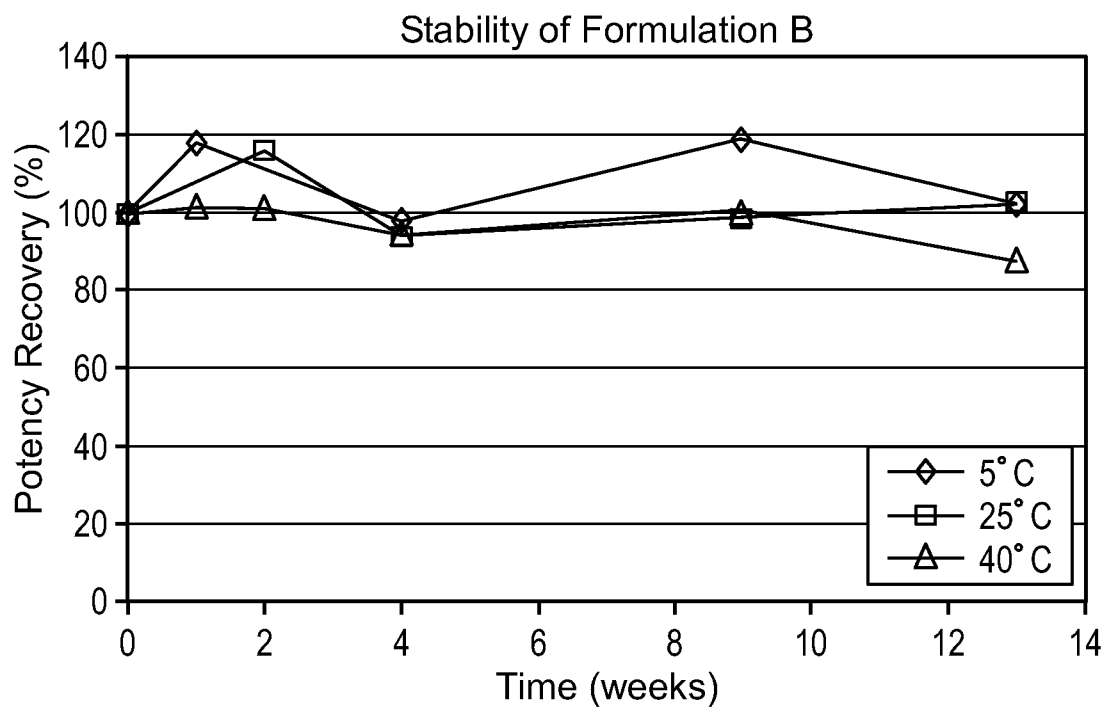


FIG. 17



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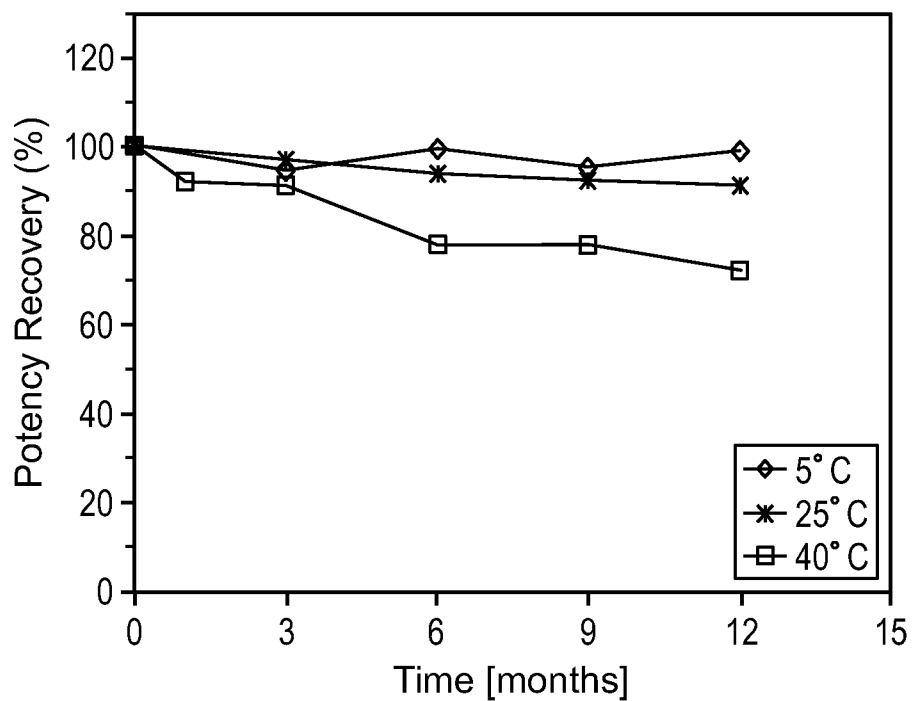


FIG. 18

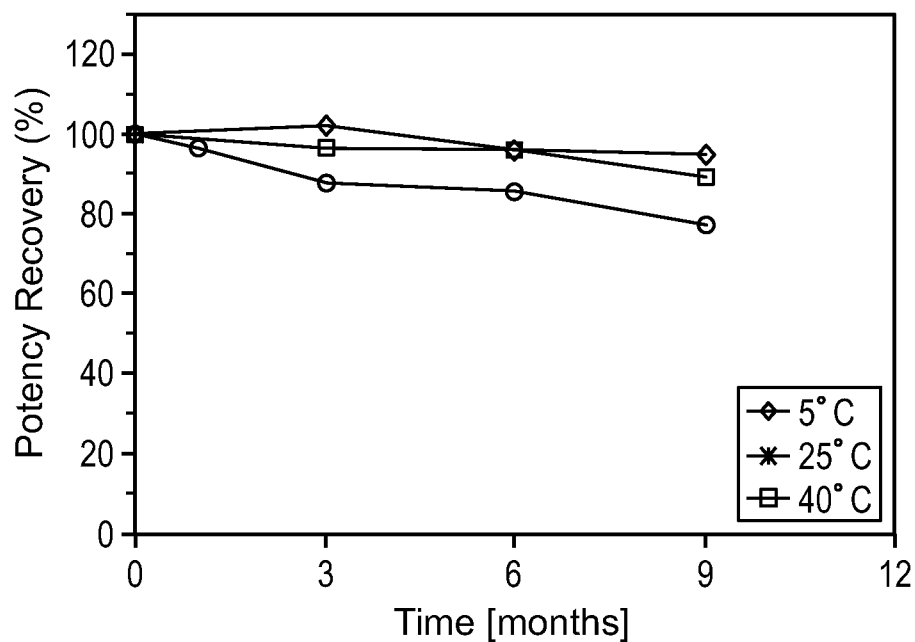


FIG. 19

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## FIG. 20

B-Domain Deleted SQ Human Factor VIII

Amino Acid Sequence SEQ ID NO: 3

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121 gvsywkaseg aeyddqtsqr ekeddkvfpg gshtyvwqvl kengpmasdp lcltysylsh
181 vdlvkdlngs ligallvcre gslakektqt lhkfillfav fdegkswhse tknslmqdrd
241 aasarawpkm htvngyvnrslpgligchrk svywhvigmg tpevhsifl eghtflvrnh
301 rqaaleispi tfltaqtllm dlqqfl1fch issqhhdgme aykvdscepepqlrmknne
361 eaedydddltdsemdvvrfd ddnspsf1qi rsvakkhpkt wvhyaeeee dwdyapl1vla
421 pddrsyksqy lnngprigr kykkvrfmay tdetftrea iqhesgilgp llygevgdtl
481 liifknqasr pyniypghit dvrplysrrl pkgvkh1kdf pilpgeifky kwtvtvedgp
541 tksdprcltr ysssfvnmerr dlasgligpl licykesvdq rgnqimsdkr nvilfsvfde
601 nrswylteni qrflpnpagv qledpefqas nimhsingyv fds1qlsvcl hevaywyils
661 igaqtdflsv ffsqytfkhk mvyedtl1tf pfsgetvfms menpg1wilg chnsdfrnrg
721 mtallkvssc dkntgdyyed syedisayll sknnaieprs fsqnpplv1kr hqreitr1tl
781 qsdqeeidyd dtisvemk dfdiydeden qsprsfqkt rhyfiaaver lwdygmsssp
841 hvlnraqsg svpqfkkvfv qeftdgsftq plyrgelneh lgllgpyira evednimvtf
901 rnqasrpysf ysslisyeed qrqgaep1rkn fvkpnetkty fwkvqhhmap tkdefdc1kaw
961 ayfsdvdlek dvhsgligpl lvchtntlnp ahgrqvtvqe falfftifde tkswyf1tenm
1021 erncrapcni qmedptfken yrfhaingyi mdtlpglvma qdqrirwyll smgsnenihs
1081 ihfsghvftv rkkeykmal ynlypgvfet vemlpskagi wrvecligeh lhagmst1fl
1141 vysnkcqtp1l gmasghirdf qitasgqygq wapklar1lhy sgsinawstk epfswikvdl
1201 lapmiihgik tqgarqkfss lyisqfiimy sldgkkwqty rgnstgtl1mv ffgnvdssgi
1261 khnifnppii aryirlhpth ysirstl1rme lmgcdlnscs mplgmeskai sdaqitassy
1321 ftnmfatwsp skar1hlqgr snawrpqvn1n pkewlqvdfq ktmkvtgvt1t qgvks1ltsm
1381 yvkefliss qdghqwtl1ff qngkvkvfqg nqdsftpvvn sl1dpp1ltry lrihpqswvh
1441 qialrmevlg cea1qdly

```

# FIG. 21A

Human Factor VIII Amino Acid Sequence

SEQ ID NO: 1

```

1  mqielsstcfff  lclllrfcfsa  trryylgave  lswdymqsdl  gelpvdarfp  prvpksfpfn
61  tsvvykktlff  veftvhlfnl  akprppwmg1  lgptiaevy  dtvvitlknm  ashpvs1hav
121  gvsywkaseg  aeyddqtsqr  ekeddkvfpg  gshtyvwqv1  kengpmasdp  lcltysylsh
181  vdlvkdlngs  ligallvcre  gslakektqt  lkhfillfav  fdegkshwse  tknslmqdrd
241  aasarawpkm  htvngyvnr  lpgligchrk  svywhvigmg  ttpevhsifl  eghtflvrnh
301  rqaaleispi  tfltaqtllm  dlqgflfch  isshqhdgme  aykvdscepe  epqlrmknne
361  eaedydddl  dsemdvvrfd  ddnspsfiqi  rsvakkhpkt  wvhyaaeae  dwdyaplvla
421  pddrsyksqy  lnngpqrigr  kykkvrfmay  tdetfktrea  iqhesgilgp  llygevgdt1
481  liifknqasr  pyniypghit  dvrplysrr1  pkgvkhlkdf  pilpgeifky  kwtvtvedgp
541  tksdprcltr  yyssfvnmer  dlasgligp1  licykesvdq  rgnqimsdkr  nvilfsvfde
601  nrswylteni  qrflpnpagv  qledpefqas  nimhsingyv  fdsllqlsvcl  hevaywyils
661  igaqtdflsv  ffsgytfkhh  mvyedtltlf  pfsgetvfms  menpglwilg  chnsdfrnrg
721  mtallkvssc  dkntgdyyed  syedisayll  sknaieprs  fsqnsrhpst  rqkqfnatti
781  pendiektdp  wfahrtpmpk  iqnvsssdll  mllrqspth  glslsdlqea  kyetfsddps
841  pgaidsnns1  semthfrpql  hhsgdmvftp  esglqlrlne  klgttaatel  kkldfkvsst
901  snnlistips  dnlaagtdnt  sslgppsmpv  hydsqldttl  fgkssplte  sggpls1see
961  ndskllesg  lmnsqesswg  knvssstesgr  lfkgrahgp  alltkdnalf  kvsisl1ktn
1021  ktsnnsatnr  kthidgpsll  ienspsvwqn  ilesdtefkk  vtplihdrml  mdknata1rl
1081  nhmsnktts  knmemvqqk  egpipdaqn  pdmsffkmlf  lpesarwiqr  thgkns1nsg

```

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## FIG. 21B

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 1201 lhennthqe kkiqeeiekk etliqenvvl pqihtvtgtk nfmknlflls trqnvegsye  
 1261 gayapvlqdf rslndstnrt kkhtahfskk geeenleglg nqtqgiveky acttrispnt  
 1321 sqqnfvqtqs kralqftrlp leetelekri ivddtstqws knmkhltpst ltqidyneke  
 1381 kgaitqspls dcltrshsip qanrslpia kvssfpsirp iyltrvlfqd nsshlpaaasy  
 1441 rkkdsgvqes shflqgakk nlsalitle mtgdqrevgs lgtsatnsvt ykkventvlp  
 1501 kpdlpktskg vellpkvhiy qkd1fptets ngspghldlv egsl1qgteg aikwneanrp  
 1561 gkvpflrvat essaktpsk1 ldplawdnhy gtqipkeewk sqekspekta fkkdt1sl  
 1621 nacesnhaia ainegqnkpe ievtwakqgr terlcsqnpp vlkrhquireit rt1lqsdqee  
 1681 idyddtisve mkkedfdi1d edenqsprsf qkktrhyfia aver1wdygm sssphv1rn1r  
 1741 aqsgsvpqfk kvvfqeftdg sftqplyrge lnehlgllgp yiraevedni mvtfrnqasr  
 1801 pysfysslis yeedqrqgae prknfvkpne tktyfwkvqh hmaptkdefd ckawayfsdv  
 1861 dlekdvhsg1 igp1lvcht1n t1npahgrqv tvqefalfft ifdetkswyf tenmerncra  
 1921 pcniqmedpt fkenyrfhai ngyimdt1pg lvmaqqdrir wyllsmgsne nihsihfsg1h  
 1981 vftvrkkey kmalynlypg vfetvem1ps kagiwrvecl ige1lhagms t1flvysnkc  
 2041 qtplgmasgh irdfqitasg qygqwapkla rlhysgsina wstkepfswi kvdl1lapmii  
 2101 hgiktqgarq kfsslyisqf iimysldgk1 wqtyrgnstg tlmvffgnvd ssgikh1nifn  
 2161 ppiariyir1 hpthysir1st lrmelmgcd1 nscsmplgme skaisdaqit assyftnmfa  
 2221 twspskarlh lqgrsnawrp qvnnpkewlq vdfqktmkvt gvt1tqgvks1 ltsmyvkef1  
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Sequence Listing  
SEQUENCE LISTING

<110> Bayer HealthCare LLC  
Wang, De Qian  
Ma, Xinghang  
Tsvetkova, Nelly

<120> RECOMBINANT FACTOR VIII FORMULATIONS

<130> BHC 135014 PCT (0081563-000121)

<150> US 61/869,191

<151> 2013-08-23

<150> US 61/779,495

<151> 2013-03-15

<160> 5

<170> PatentIn version 3.5

<210> 1

<211> 2351

<212> PRT

<213> Homo sapiens

<400> 1

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

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val  
50 55 60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile  
65 70 75 80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln  
85 90 95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser  
100 105 110

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser  
115 120 125

Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp  
130 135 140

Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu  
145 150 155 160

# Sequence\_Listing

Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser  
 165 170 175  
 Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile  
 180 185 190  
 Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr  
 195 200 205  
 Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly  
 210 215 220  
 Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp  
 225 230 235 240  
 Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr  
 245 250 255  
 Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val  
 260 265 270  
 Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile  
 275 280 285  
 Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser  
 290 295 300  
 Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met  
 305 310 315 320  
 Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His  
 325 330 335  
 Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro  
 340 345 350  
 Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp  
 355 360 365  
 Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser  
 370 375 380  
 Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr  
 385 390 395 400  
 Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro  
 405 410 415  
 Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn  
 420 425 430

# Sequence Listing

Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met  
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 Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu  
 450 455 460  
 Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu  
 465 470 475 480  
 Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro  
 485 490 495  
 His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys  
 500 505 510  
 Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe  
 515 520 525  
 Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp  
 530 535 540  
 Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg  
 545 550 555 560  
 Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu  
 565 570 575  
 Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val  
 580 585 590  
 Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu  
 595 600 605  
 Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp  
 610 615 620  
 Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val  
 625 630 635 640  
 Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp  
 645 650 655  
 Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe  
 660 665 670  
 Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr  
 675 680 685  
 Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro  
 690 695 700

# Sequence Listing

Gly 705 Leu Trp Ile Leu Gly 710 Cys His Asn Ser Asp 715 Phe Arg Asn Arg Gly 720  
 Met Thr Ala Leu 725 Leu Lys Val Ser Ser Cys 730 Asp Lys Asn Thr Gly 735 Asp  
 Tyr Tyr Glu 740 Asp Ser Tyr Glu Asp Ile 745 Ser Ala Tyr Leu Leu 750 Ser Lys  
 Asn Asn Ala 755 Ile Glu Pro Arg Ser 760 Phe Ser Gln Asn Ser 765 Arg His Pro  
 Ser Thr Arg Gln Lys Gln Phe 775 Asn Ala Thr Thr Ile 780 Pro Glu Asn Asp  
 Ile 785 Glu Lys Thr Asp Pro 790 Trp Phe Ala His Arg 795 Thr Pro Met Pro Lys 800  
 Ile Gln Asn Val Ser 805 Ser Ser Asp Leu Leu 810 Met Leu Leu Arg Gln Ser 815  
 Pro Thr Pro His 820 Gly Leu Ser Leu Ser 825 Asp Leu Gln Glu Ala Lys Tyr  
 Glu Thr Phe 835 Ser Asp Asp Pro Ser 840 Pro Gly Ala Ile Asp 845 Ser Asn Asn  
 Ser Leu 850 Ser Glu Met Thr His 855 Phe Arg Pro Gln Leu His His Ser Gly  
 Asp 865 Met Val Phe Thr Pro 870 Glu Ser Gly Leu Gln 875 Leu Arg Leu Asn Glu 880  
 Lys Leu Gly Thr Thr 885 Ala Ala Thr Glu Leu 890 Lys Lys Leu Asp Phe Lys 895  
 Val Ser Ser Thr 900 Ser Asn Asn Leu Ile 905 Ser Thr Ile Pro Ser 910 Asp Asn  
 Leu Ala Ala 915 Gly Thr Asp Asn Thr Ser Ser Leu Gly Pro 925 Pro Ser Met  
 Pro Val His Tyr Asp Ser Gln 935 Leu Asp Thr Thr Leu Phe Gly Lys Lys 940  
 Ser 945 Ser Pro Leu Thr Glu 950 Ser Gly Gly Pro Leu 955 Ser Leu Ser Glu Glu 960  
 Asn Asn Asp Ser Lys 965 Leu Leu Glu Ser Gly 970 Leu Met Asn Ser Gln Glu 975



# Sequence Listing

Ser Ser Trp Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe  
 980 985 990  
 Lys Gly Lys Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala  
 995 1000 1005  
 Leu Phe Lys Val Ser Ile Ser Leu Leu Lys Thr Asn Lys Thr Ser  
 1010 1015 1020  
 Asn Asn Ser Ala Thr Asn Arg Lys Thr His Ile Asp Gly Pro Ser  
 1025 1030 1035  
 Leu Leu Ile Glu Asn Ser Pro Ser Val Trp Gln Asn Ile Leu Glu  
 1040 1045 1050  
 Ser Asp Thr Glu Phe Lys Lys Val Thr Pro Leu Ile His Asp Arg  
 1055 1060 1065  
 Met Leu Met Asp Lys Asn Ala Thr Ala Leu Arg Leu Asn His Met  
 1070 1075 1080  
 Ser Asn Lys Thr Thr Ser Ser Lys Asn Met Glu Met Val Gln Gln  
 1085 1090 1095  
 Lys Lys Glu Gly Pro Ile Pro Pro Asp Ala Gln Asn Pro Asp Met  
 1100 1105 1110  
 Ser Phe Phe Lys Met Leu Phe Leu Pro Glu Ser Ala Arg Trp Ile  
 1115 1120 1125  
 Gln Arg Thr His Gly Lys Asn Ser Leu Asn Ser Gly Gln Gly Pro  
 1130 1135 1140  
 Ser Pro Lys Gln Leu Val Ser Leu Gly Pro Glu Lys Ser Val Glu  
 1145 1150 1155  
 Gly Gln Asn Phe Leu Ser Glu Lys Asn Lys Val Val Val Gly Lys  
 1160 1165 1170  
 Gly Glu Phe Thr Lys Asp Val Gly Leu Lys Glu Met Val Phe Pro  
 1175 1180 1185  
 Ser Ser Arg Asn Leu Phe Leu Thr Asn Leu Asp Asn Leu His Glu  
 1190 1195 1200  
 Asn Asn Thr His Asn Gln Glu Lys Lys Ile Gln Glu Glu Ile Glu  
 1205 1210 1215  
 Lys Lys Glu Thr Leu Ile Gln Glu Asn Val Val Leu Pro Gln Ile  
 1220 1225 1230

# Sequence Listing

His	Thr	Val	Thr	Gly	Thr	Lys	Asn	Phe	Met	Lys	Asn	Leu	Phe	Leu
	1235					1240					1245			
Leu	Ser	Thr	Arg	Gln	Asn	Val	Glu	Gly	Ser	Tyr	Glu	Gly	Ala	Tyr
	1250					1255					1260			
Ala	Pro	Val	Leu	Gln	Asp	Phe	Arg	Ser	Leu	Asn	Asp	Ser	Thr	Asn
	1265					1270					1275			
Arg	Thr	Lys	Lys	His	Thr	Ala	His	Phe	Ser	Lys	Lys	Gly	Glu	Glu
	1280					1285					1290			
Glu	Asn	Leu	Glu	Gly	Leu	Gly	Asn	Gln	Thr	Lys	Gln	Ile	Val	Glu
	1295					1300					1305			
Lys	Tyr	Ala	Cys	Thr	Thr	Arg	Ile	Ser	Pro	Asn	Thr	Ser	Gln	Gln
	1310					1315					1320			
Asn	Phe	Val	Thr	Gln	Arg	Ser	Lys	Arg	Ala	Leu	Lys	Gln	Phe	Arg
	1325					1330					1335			
Leu	Pro	Leu	Glu	Glu	Thr	Glu	Leu	Glu	Lys	Arg	Ile	Ile	Val	Asp
	1340					1345					1350			
Asp	Thr	Ser	Thr	Gln	Trp	Ser	Lys	Asn	Met	Lys	His	Leu	Thr	Pro
	1355					1360					1365			
Ser	Thr	Leu	Thr	Gln	Ile	Asp	Tyr	Asn	Glu	Lys	Glu	Lys	Gly	Ala
	1370					1375					1380			
Ile	Thr	Gln	Ser	Pro	Leu	Ser	Asp	Cys	Leu	Thr	Arg	Ser	His	Ser
	1385					1390					1395			
Ile	Pro	Gln	Ala	Asn	Arg	Ser	Pro	Leu	Pro	Ile	Ala	Lys	Val	Ser
	1400					1405					1410			
Ser	Phe	Pro	Ser	Ile	Arg	Pro	Ile	Tyr	Leu	Thr	Arg	Val	Leu	Phe
	1415					1420					1425			
Gln	Asp	Asn	Ser	Ser	His	Leu	Pro	Ala	Ala	Ser	Tyr	Arg	Lys	Lys
	1430					1435					1440			
Asp	Ser	Gly	Val	Gln	Glu	Ser	Ser	His	Phe	Leu	Gln	Gly	Ala	Lys
	1445					1450					1455			
Lys	Asn	Asn	Leu	Ser	Leu	Ala	Ile	Leu	Thr	Leu	Glu	Met	Thr	Gly
	1460					1465					1470			
Asp	Gln	Arg	Glu	Val	Gly	Ser	Leu	Gly	Thr	Ser	Ala	Thr	Asn	Ser
	1475					1480					1485			

# Sequence\_Listing

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Leu	Pro	Lys	Thr	Ser	Gly	Lys	Val	Glu	Leu	Leu	Pro	Lys	Val	His
	1505					1510					1515			
Ile	Tyr	Gln	Lys	Asp	Leu	Phe	Pro	Thr	Glu	Thr	Ser	Asn	Gly	Ser
	1520					1525					1530			
Pro	Gly	His	Leu	Asp	Leu	Val	Glu	Gly	Ser	Leu	Leu	Gln	Gly	Thr
	1535					1540					1545			
Glu	Gly	Ala	Ile	Lys	Trp	Asn	Glu	Ala	Asn	Arg	Pro	Gly	Lys	Val
	1550					1555					1560			
Pro	Phe	Leu	Arg	Val	Ala	Thr	Glu	Ser	Ser	Ala	Lys	Thr	Pro	Ser
	1565					1570					1575			
Lys	Leu	Leu	Asp	Pro	Leu	Ala	Trp	Asp	Asn	His	Tyr	Gly	Thr	Gln
	1580					1585					1590			
Ile	Pro	Lys	Glu	Glu	Trp	Lys	Ser	Gln	Glu	Lys	Ser	Pro	Glu	Lys
	1595					1600					1605			
Thr	Ala	Phe	Lys	Lys	Lys	Asp	Thr	Ile	Leu	Ser	Leu	Asn	Ala	Cys
	1610					1615					1620			
Glu	Ser	Asn	His	Ala	Ile	Ala	Ala	Ile	Asn	Glu	Gly	Gln	Asn	Lys
	1625					1630					1635			
Pro	Glu	Ile	Glu	Val	Thr	Trp	Ala	Lys	Gln	Gly	Arg	Thr	Glu	Arg
	1640					1645					1650			
Leu	Cys	Ser	Gln	Asn	Pro	Pro	Val	Leu	Lys	Arg	His	Gln	Arg	Glu
	1655					1660					1665			
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	1670					1675					1680			
Asp	Asp	Thr	Ile	Ser	Val	Glu	Met	Lys	Lys	Glu	Asp	Phe	Asp	Ile
	1685					1690					1695			
Tyr	Asp	Glu	Asp	Glu	Asn	Gln	Ser	Pro	Arg	Ser	Phe	Gln	Lys	Lys
	1700					1705					1710			
Thr	Arg	His	Tyr	Phe	Ile	Ala	Ala	Val	Glu	Arg	Leu	Trp	Asp	Tyr
	1715					1720					1725			
Gly	Met	Ser	Ser	Ser	Pro	His	Val	Leu	Arg	Asn	Arg	Ala	Gln	Ser
	1730					1735					1740			

# Sequence\_Listing

Gly	Ser	Val	Pro	Gln	Phe	Lys	Lys	Val	Val	Phe	Gln	Glu	Phe	Thr
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Asp	Gly	Ser	Phe	Thr	Gln	Pro	Leu	Tyr	Arg	Gly	Glu	Leu	Asn	Glu
1760						1765					1770			
His	Leu	Gly	Leu	Leu	Gly	Pro	Tyr	Ile	Arg	Ala	Glu	Val	Glu	Asp
1775						1780					1785			
Asn	Ile	Met	Val	Thr	Phe	Arg	Asn	Gln	Ala	Ser	Arg	Pro	Tyr	Ser
1790						1795					1800			
Phe	Tyr	Ser	Ser	Leu	Ile	Ser	Tyr	Glu	Glu	Asp	Gln	Arg	Gln	Gly
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Ala	Glu	Pro	Arg	Lys	Asn	Phe	Val	Lys	Pro	Asn	Glu	Thr	Lys	Thr
1820						1825					1830			
Tyr	Phe	Trp	Lys	Val	Gln	His	His	Met	Ala	Pro	Thr	Lys	Asp	Glu
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Phe	Asp	Cys	Lys	Ala	Trp	Ala	Tyr	Phe	Ser	Asp	Val	Asp	Leu	Glu
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Lys	Asp	Val	His	Ser	Gly	Leu	Ile	Gly	Pro	Leu	Leu	Val	Cys	His
1865						1870					1875			
Thr	Asn	Thr	Leu	Asn	Pro	Ala	His	Gly	Arg	Gln	Val	Thr	Val	Gln
1880						1885					1890			
Glu	Phe	Ala	Leu	Phe	Phe	Thr	Ile	Phe	Asp	Glu	Thr	Lys	Ser	Trp
1895						1900					1905			
Tyr	Phe	Thr	Glu	Asn	Met	Glu	Arg	Asn	Cys	Arg	Ala	Pro	Cys	Asn
1910						1915					1920			
Ile	Gln	Met	Glu	Asp	Pro	Thr	Phe	Lys	Glu	Asn	Tyr	Arg	Phe	His
1925						1930					1935			
Ala	Ile	Asn	Gly	Tyr	Ile	Met	Asp	Thr	Leu	Pro	Gly	Leu	Val	Met
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Ala	Gln	Asp	Gln	Arg	Ile	Arg	Trp	Tyr	Leu	Leu	Ser	Met	Gly	Ser
1955						1960					1965			
Asn	Glu	Asn	Ile	His	Ser	Ile	His	Phe	Ser	Gly	His	Val	Phe	Thr
1970						1975					1980			
Val	Arg	Lys	Lys	Glu	Glu	Tyr	Lys	Met	Ala	Leu	Tyr	Asn	Leu	Tyr
1985						1990					1995			

Sequence\_Listing

Pro	Gly	Val	Phe	Glu	Thr	Val	Glu	Met	Leu	Pro	Ser	Lys	Ala	Gly
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Ile	Trp	Arg	Val	Glu	Cys	Leu	Ile	Gly	Glu	His	Leu	His	Ala	Gly
2015						2020					2025			
Met	Ser	Thr	Leu	Phe	Leu	Val	Tyr	Ser	Asn	Lys	Cys	Gln	Thr	Pro
2030						2035					2040			
Leu	Gly	Met	Ala	Ser	Gly	His	Ile	Arg	Asp	Phe	Gln	Ile	Thr	Ala
2045						2050					2055			
Ser	Gly	Gln	Tyr	Gly	Gln	Trp	Ala	Pro	Lys	Leu	Ala	Arg	Leu	His
2060						2065					2070			
Tyr	Ser	Gly	Ser	Ile	Asn	Ala	Trp	Ser	Thr	Lys	Glu	Pro	Phe	Ser
2075						2080					2085			
Trp	Ile	Lys	Val	Asp	Leu	Leu	Ala	Pro	Met	Ile	Ile	His	Gly	Ile
2090						2095					2100			
Lys	Thr	Gln	Gly	Ala	Arg	Gln	Lys	Phe	Ser	Ser	Leu	Tyr	Ile	Ser
2105						2110					2115			
Gln	Phe	Ile	Ile	Met	Tyr	Ser	Leu	Asp	Gly	Lys	Lys	Trp	Gln	Thr
2120						2125					2130			
Tyr	Arg	Gly	Asn	Ser	Thr	Gly	Thr	Leu	Met	Val	Phe	Phe	Gly	Asn
2135						2140					2145			
Val	Asp	Ser	Ser	Gly	Ile	Lys	His	Asn	Ile	Phe	Asn	Pro	Pro	Ile
2150						2155					2160			
Ile	Ala	Arg	Tyr	Ile	Arg	Leu	His	Pro	Thr	His	Tyr	Ser	Ile	Arg
2165						2170					2175			
Ser	Thr	Leu	Arg	Met	Glu	Leu	Met	Gly	Cys	Asp	Leu	Asn	Ser	Cys
2180						2185					2190			
Ser	Met	Pro	Leu	Gly	Met	Glu	Ser	Lys	Ala	Ile	Ser	Asp	Ala	Gln
2195						2200					2205			
Ile	Thr	Ala	Ser	Ser	Tyr	Phe	Thr	Asn	Met	Phe	Ala	Thr	Trp	Ser
2210						2215					2220			
Pro	Ser	Lys	Ala	Arg	Leu	His	Leu	Gln	Gly	Arg	Ser	Asn	Ala	Trp
2225						2230					2235			
Arg	Pro	Gln	Val	Asn	Asn	Pro	Lys	Glu	Trp	Leu	Gln	Val	Asp	Phe
2240						2245					2250			

# Sequence\_Listing

Gln Lys Thr Met Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys  
2255 2260 2265

Ser Leu Leu Thr Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser  
2270 2275 2280

Ser Gln Asp Gly His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys  
2285 2290 2295

Val Lys Val Phe Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val  
2300 2305 2310

Asn Ser Leu Asp Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His  
2315 2320 2325

Pro Gln Ser Trp Val His Gln Ile Ala Leu Arg Met Glu Val Leu  
2330 2335 2340

Gly Cys Glu Ala Gln Asp Leu Tyr  
2345 2350

<210> 2  
<211> 2332  
<212> PRT  
<213> Homo sapiens  
<400> 2

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

Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg Phe Pro Pro  
20 25 30

Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val Tyr Lys Lys  
35 40 45

Thr Leu Phe Val Glu Phe Thr Val His Leu Phe Asn Ile Ala Lys Pro  
50 55 60

Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val  
65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val  
85 90 95

Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser Glu Gly Ala  
100 105 110

Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp Asp Lys Val  
115 120 125

Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu Lys Glu Asn  
Page 10

## Sequence\_Listing

130                                      135                                      140  
 Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser Tyr Leu Ser  
 145                                      150                                      155                                      160  
 His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile Gly Ala Leu  
                                     165                                      170                                      175  
 Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr Gln Thr Leu  
                                     180                                      185                                      190  
 His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly Lys Ser Trp  
                                     195                                      200                                      205  
 His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp Ala Ala Ser  
                                     210                                      215                                      220  
 Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr Val Asn Arg  
 225                                      230                                      235                                      240  
 Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val Tyr Trp His  
                                     245                                      250                                      255  
 Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile Phe Leu Glu  
                                     260                                      265                                      270  
 Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser Leu Glu Ile  
                                     275                                      280                                      285  
 Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met Asp Leu Gly  
                                     290                                      295                                      300  
 Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His Asp Gly Met  
 305                                      310                                      315                                      320  
 Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro Gln Leu Arg  
                                     325                                      330                                      335  
 Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp Leu Thr Asp  
                                     340                                      345                                      350  
 Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser Pro Ser Phe  
                                     355                                      360                                      365  
 Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr Trp Val His  
                                     370                                      375                                      380  
 Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro Leu Val Leu  
 385                                      390                                      395                                      400  
 Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn Asn Gly Pro

# Sequence\_Listing

405

410

415

Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met Ala Tyr Thr  
420 425 430

Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu Ser Gly Ile  
435 440 445

Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu Leu Ile Ile  
450 455 460

Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro His Gly Ile  
465 470 475 480

Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys Gly Val Lys  
485 490 495

His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe Lys Tyr Lys  
500 505 510

Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp Pro Arg Cys  
515 520 525

Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg Asp Leu Ala  
530 535 540

Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu Ser Val Asp  
545 550 555 560

Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val Ile Leu Phe  
565 570 575

Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu Asn Ile Gln  
580 585 590

Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp Pro Glu Phe  
595 600 605

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

Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp Tyr Ile Leu  
625 630 635 640

Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe Ser Gly Tyr  
645 650 655

Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr Leu Phe Pro  
660 665 670

Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro Gly Leu Trp



685

Ser Lys Leu Leu Glu Ser Gly Leu Met Asn Ser Gln Glu Ser Ser Trp

955

945

950

960

Gly Lys Asn Val Ser Ser Thr Glu Ser Gly Arg Leu Phe Lys Gly Lys  
965 970 975

Arg Ala His Gly Pro Ala Leu Leu Thr Lys Asp Asn Ala Leu Phe Lys  
980 985 990

Val Ser Ile Ser Leu Leu Lys Thr Asn Lys Thr Ser Asn Asn Ser Ala  
995 1000 1005

Thr Asn Arg Lys Thr His Ile Asp Gly Pro Ser Leu Leu Ile Glu  
1010 1015 1020

Asn Ser Pro Ser Val Trp Gln Asn Ile Leu Glu Ser Asp Thr Glu  
1025 1030 1035

Phe Lys Lys Val Thr Pro Leu Ile His Asp Arg Met Leu Met Asp  
1040 1045 1050

Lys Asn Ala Thr Ala Leu Arg Leu Asn His Met Ser Asn Lys Thr  
1055 1060 1065

Thr Ser Ser Lys Asn Met Glu Met Val Gl n Gl n Lys Lys Glu Gly  
1070 1075 1080

Pro Ile Pro Pro Asp Ala Gln Asn Pro Asp Met Ser Phe Phe Lys  
1085 1090 1095

Met Leu Phe Leu Pro Glu Ser Ala Arg Trp Ile Gln Arg Thr His  
1100 1105 1110

Gly Lys Asn Ser Leu Asn Ser Gly Gl n Gly Pro Ser Pro Lys Gl n  
1115 1120 1125

Leu Val Ser Leu Gly Pro Glu Lys Ser Val Glu Gly Gln Asn Phe  
1130 1135 1140

Leu Ser Glu Lys Asn Lys Val Val Val Gly Lys Gly Glu Phe Thr  
1145 1150 1155

Lys Asp Val Gly Leu Lys Glu Met Val Phe Pro Ser Ser Arg Asn  
1160 1165 1170

Leu Phe    Leu Thr Asn Leu Asp    Asn Leu His Glu Asn    Asn Thr His  
1175                      1180                      1185

Asn Gln Glu Lys Lys Ile Gln Glu Glu Ile Glu Lys Lys Glu Thr  
1190 1195 1200

Leu Ile Gln Glu Asn Val Val Leu Pro Gln Ile His Thr Val Thr

## Sequence\_Listing

1205						1210						1215			
Gly	Thr	Lys	Asn	Phe	Met	Lys	Asn	Leu	Phe	Leu	Leu	Ser	Thr	Arg	
1220						1225					1230				
Gln	Asn	Val	Glu	Gly	Ser	Tyr	Glu	Gly	Ala	Tyr	Ala	Pro	Val	Leu	
1235						1240					1245				
Gln	Asp	Phe	Arg	Ser	Leu	Asn	Asp	Ser	Thr	Asn	Arg	Thr	Lys	Lys	
1250						1255					1260				
His	Thr	Ala	His	Phe	Ser	Lys	Lys	Gly	Glu	Glu	Glu	Asn	Leu	Glu	
1265						1270					1275				
Gly	Leu	Gly	Asn	Gln	Thr	Lys	Gln	Ile	Val	Glu	Lys	Tyr	Ala	Cys	
1280						1285					1290				
Thr	Thr	Arg	Ile	Ser	Pro	Asn	Thr	Ser	Gln	Gln	Asn	Phe	Val	Thr	
1295						1300					1305				
Gln	Arg	Ser	Lys	Arg	Ala	Leu	Lys	Gln	Phe	Arg	Leu	Pro	Leu	Glu	
1310						1315					1320				
Glu	Thr	Glu	Leu	Glu	Lys	Arg	Ile	Ile	Val	Asp	Asp	Thr	Ser	Thr	
1325						1330					1335				
Gln	Trp	Ser	Lys	Asn	Met	Lys	His	Leu	Thr	Pro	Ser	Thr	Leu	Thr	
1340						1345					1350				
Gln	Ile	Asp	Tyr	Asn	Glu	Lys	Glu	Lys	Gly	Ala	Ile	Thr	Gln	Ser	
1355						1360					1365				
Pro	Leu	Ser	Asp	Cys	Leu	Thr	Arg	Ser	His	Ser	Ile	Pro	Gln	Ala	
1370						1375					1380				
Asn	Arg	Ser	Pro	Leu	Pro	Ile	Ala	Lys	Val	Ser	Ser	Phe	Pro	Ser	
1385						1390					1395				
Ile	Arg	Pro	Ile	Tyr	Leu	Thr	Arg	Val	Leu	Phe	Gln	Asp	Asn	Ser	
1400						1405					1410				
Ser	His	Leu	Pro	Ala	Ala	Ser	Tyr	Arg	Lys	Lys	Asp	Ser	Gly	Val	
1415						1420					1425				
Gln	Glu	Ser	Ser	His	Phe	Leu	Gln	Gly	Ala	Lys	Lys	Asn	Asn	Leu	
1430						1435					1440				
Ser	Leu	Ala	Ile	Leu	Thr	Leu	Glu	Met	Thr	Gly	Asp	Gln	Arg	Glu	
1445						1450					1455				
Val	Gly	Ser	Leu	Gly	Thr	Ser	Ala	Thr	Asn	Ser	Val	Thr	Tyr	Lys	

## Sequence\_Listing

1460						1465						1470			
Lys	Val	Glu	Asn	Thr	Val	Leu	Pro	Lys	Pro	Asp	Leu	Pro	Lys	Thr	
1475						1480					1485				
Ser	Gly	Lys	Val	Glu	Leu	Leu	Pro	Lys	Val	His	Ile	Tyr	Gln	Lys	
1490						1495					1500				
Asp	Leu	Phe	Pro	Thr	Glu	Thr	Ser	Asn	Gly	Ser	Pro	Gly	His	Leu	
1505						1510					1515				
Asp	Leu	Val	Glu	Gly	Ser	Leu	Leu	Gln	Gly	Thr	Glu	Gly	Ala	Ile	
1520						1525					1530				
Lys	Trp	Asn	Glu	Ala	Asn	Arg	Pro	Gly	Lys	Val	Pro	Phe	Leu	Arg	
1535						1540					1545				
Val	Ala	Thr	Glu	Ser	Ser	Ala	Lys	Thr	Pro	Ser	Lys	Leu	Leu	Asp	
1550						1555					1560				
Pro	Leu	Ala	Trp	Asp	Asn	His	Tyr	Gly	Thr	Gln	Ile	Pro	Lys	Glu	
1565						1570					1575				
Glu	Trp	Lys	Ser	Gln	Glu	Lys	Ser	Pro	Glu	Lys	Thr	Ala	Phe	Lys	
1580						1585					1590				
Lys	Lys	Asp	Thr	Ile	Leu	Ser	Leu	Asn	Ala	Cys	Glu	Ser	Asn	His	
1595						1600					1605				
Ala	Ile	Ala	Ala	Ile	Asn	Glu	Gly	Gln	Asn	Lys	Pro	Glu	Ile	Glu	
1610						1615					1620				
Val	Thr	Trp	Ala	Lys	Gln	Gly	Arg	Thr	Glu	Arg	Leu	Cys	Ser	Gln	
1625						1630					1635				
Asn	Pro	Pro	Val	Leu	Lys	Arg	His	Gln	Arg	Glu	Ile	Thr	Arg	Thr	
1640						1645					1650				
Thr	Leu	Gln	Ser	Asp	Gln	Glu	Glu	Ile	Asp	Tyr	Asp	Asp	Thr	Ile	
1655						1660					1665				
Ser	Val	Glu	Met	Lys	Lys	Glu	Asp	Phe	Asp	Ile	Tyr	Asp	Glu	Asp	
1670						1675					1680				
Glu	Asn	Gln	Ser	Pro	Arg	Ser	Phe	Gln	Lys	Lys	Thr	Arg	His	Tyr	
1685						1690					1695				
Phe	Ile	Ala	Ala	Val	Glu	Arg	Leu	Trp	Asp	Tyr	Gly	Met	Ser	Ser	
1700						1705					1710				
Ser	Pro	His	Val	Leu	Arg	Asn	Arg	Ala	Gln	Ser	Gly	Ser	Val	Pro	

## Sequence\_Listing

1715						1720						1725			
Gln	Phe	Lys	Lys	Val	Val	Phe	Gln	Glu	Phe	Thr	Asp	Gly	Ser	Phe	
1730						1735					1740				
Thr	Gln	Pro	Leu	Tyr	Arg	Gly	Glu	Leu	Asn	Glu	His	Leu	Gly	Leu	
1745						1750					1755				
Leu	Gly	Pro	Tyr	Ile	Arg	Ala	Glu	Val	Glu	Asp	Asn	Ile	Met	Val	
1760						1765					1770				
Thr	Phe	Arg	Asn	Gln	Ala	Ser	Arg	Pro	Tyr	Ser	Phe	Tyr	Ser	Ser	
1775						1780					1785				
Leu	Ile	Ser	Tyr	Glu	Glu	Asp	Gln	Arg	Gln	Gly	Ala	Glu	Pro	Arg	
1790						1795					1800				
Lys	Asn	Phe	Val	Lys	Pro	Asn	Glu	Thr	Lys	Thr	Tyr	Phe	Trp	Lys	
1805						1810					1815				
Val	Gln	His	His	Met	Ala	Pro	Thr	Lys	Asp	Glu	Phe	Asp	Cys	Lys	
1820						1825					1830				
Ala	Trp	Ala	Tyr	Phe	Ser	Asp	Val	Asp	Leu	Glu	Lys	Asp	Val	His	
1835						1840					1845				
Ser	Gly	Leu	Ile	Gly	Pro	Leu	Leu	Val	Cys	His	Thr	Asn	Thr	Leu	
1850						1855					1860				
Asn	Pro	Ala	His	Gly	Arg	Gln	Val	Thr	Val	Gln	Glu	Phe	Ala	Leu	
1865						1870					1875				
Phe	Phe	Thr	Ile	Phe	Asp	Glu	Thr	Lys	Ser	Trp	Tyr	Phe	Thr	Glu	
1880						1885					1890				
Asn	Met	Glu	Arg	Asn	Cys	Arg	Ala	Pro	Cys	Asn	Ile	Gln	Met	Glu	
1895						1900					1905				
Asp	Pro	Thr	Phe	Lys	Glu	Asn	Tyr	Arg	Phe	His	Ala	Ile	Asn	Gly	
1910						1915					1920				
Tyr	Ile	Met	Asp	Thr	Leu	Pro	Gly	Leu	Val	Met	Ala	Gln	Asp	Gln	
1925						1930					1935				
Arg	Ile	Arg	Trp	Tyr	Leu	Leu	Ser	Met	Gly	Ser	Asn	Glu	Asn	Ile	
1940						1945					1950				
His	Ser	Ile	His	Phe	Ser	Gly	His	Val	Phe	Thr	Val	Arg	Lys	Lys	
1955						1960					1965				
Glu	Glu	Tyr	Lys	Met	Ala	Leu	Tyr	Asn	Leu	Tyr	Pro	Gly	Val	Phe	

# Sequence\_Listing

1970	1975	1980										
Glu Thr 1985	Val Glu Met Leu	Pro 1990	Ser Lys Ala Gly	Ile 1995	Trp Arg Val							
Glu Cys 2000	Leu Ile Gly Glu	His 2005	Leu His Ala Gly	Met 2010	Ser Thr Leu							
Phe Leu 2015	Val Tyr Ser Asn	Lys 2020	Cys Gln Thr Pro	Leu 2025	Gly Met Ala							
Ser Gly 2030	His Ile Arg Asp	Phe 2035	Gln Ile Thr Ala	Ser 2040	Gly Gln Tyr							
Gly Gln 2045	Trp Ala Pro Lys	Leu 2050	Ala Arg Leu His	Tyr 2055	Ser Gly Ser							
Ile Asn 2060	Ala Trp Ser Thr	Lys 2065	Glu Pro Phe Ser	Trp 2070	Ile Lys Val							
Asp Leu 2075	Leu Ala Pro Met	Ile 2080	Ile His Gly Ile	Lys 2085	Thr Gln Gly							
Ala Arg 2090	Gln Lys Phe Ser	Ser 2095	Leu Tyr Ile Ser	Gln 2100	Phe Ile Ile							
Met Tyr 2105	Ser Leu Asp Gly	Lys 2110	Lys Trp Gln Thr	Tyr 2115	Arg Gly Asn							
Ser Thr 2120	Gly Thr Leu Met	Val 2125	Phe Phe Gly Asn	Val 2130	Asp Ser Ser							
Gly Ile 2135	Lys His Asn Ile	Phe 2140	Asn Pro Pro Ile	Ile 2145	Ala Arg Tyr							
Ile Arg 2150	Leu His Pro Thr	His 2155	Tyr Ser Ile Arg	Ser 2160	Thr Leu Arg							
Met Glu 2165	Leu Met Gly Cys	Asp 2170	Leu Asn Ser Cys	Ser 2175	Met Pro Leu							
Gly Met 2180	Glu Ser Lys Ala	Ile 2185	Ser Asp Ala Gln	Ile 2190	Thr Ala Ser							
Ser Tyr 2195	Phe Thr Asn Met	Phe 2200	Ala Thr Trp Ser	Pro 2205	Ser Lys Ala							
Arg Leu 2210	His Leu Gln Gly	Arg 2215	Ser Asn Ala Trp	Arg 2220	Pro Gln Val							
Asn Asn	Pro Lys Glu Trp	Leu	Gln Val Asp Phe	Gln	Lys Thr Met							

# Sequence\_Listing

2225

2230

2235

Lys Val Thr Gly Val Thr Thr Gln Gly Val Lys Ser Leu Leu Thr  
2240 2245 2250

Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly  
2255 2260 2265

His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Phe  
2270 2275 2280

Gln Gly Asn Gln Asp Ser Phe Thr Pro Val Val Asn Ser Leu Asp  
2285 2290 2295

Pro Pro Leu Leu Thr Arg Tyr Leu Arg Ile His Pro Gln Ser Trp  
2300 2305 2310

Val His Gln Ile Ala Leu Arg Met Glu Val Leu Gly Cys Glu Ala  
2315 2320 2325

Gln Asp Leu Tyr  
2330

<210> 3

<211> 1457

<212> PRT

<213> Artificial Sequence

<220>

<223> Deletion variant of human FVIII

<400> 3

Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe  
1 5 10 15

Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser  
20 25 30

Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg  
35 40 45

Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val  
50 55 60

Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile  
65 70 75 80

Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln  
85 90 95

Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser  
100 105 110

# Sequence\_Listing

His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser  
 115 120 125  
 Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp  
 130 135 140  
 Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu  
 145 150 155 160  
 Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser  
 165 170 175  
 Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile  
 180 185 190  
 Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr  
 195 200 205  
 Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly  
 210 215 220  
 Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp  
 225 230 235 240  
 Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr  
 245 250 255  
 Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val  
 260 265 270  
 Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile  
 275 280 285  
 Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser  
 290 295 300  
 Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met  
 305 310 315 320  
 Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His  
 325 330 335  
 Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro  
 340 345 350  
 Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp  
 355 360 365  
 Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser  
 370 375 380



# Sequence\_Li st i ng

Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr  
 385 390 395 400  
 Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro  
 405 410 415  
 Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn  
 420 425 430  
 Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met  
 435 440 445  
 Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu  
 450 455 460  
 Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu  
 465 470 475 480  
 Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro  
 485 490 495  
 His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys  
 500 505 510  
 Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe  
 515 520 525  
 Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp  
 530 535 540  
 Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg  
 545 550 555 560  
 Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu  
 565 570 575  
 Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val  
 580 585 590  
 Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu  
 595 600 605  
 Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp  
 610 615 620  
 Pro Glu Phe Gln Ala Ser Asn Ile Met His Ser Ile Asn Gly Tyr Val  
 625 630 635 640  
 Phe Asp Ser Leu Gln Leu Ser Val Cys Leu His Glu Val Ala Tyr Trp  
 645 650 655

# Sequence\_Li st i ng

Tyr Ile Leu Ser Ile Gly Ala Gln Thr Asp Phe Leu Ser Val Phe Phe  
 660 665 670  
 Ser Gly Tyr Thr Phe Lys His Lys Met Val Tyr Glu Asp Thr Leu Thr  
 675 680 685  
 Leu Phe Pro Phe Ser Gly Glu Thr Val Phe Met Ser Met Glu Asn Pro  
 690 695 700  
 Gly Leu Trp Ile Leu Gly Cys His Asn Ser Asp Phe Arg Asn Arg Gly  
 705 710 715 720  
 Met Thr Ala Leu Leu Lys Val Ser Ser Cys Asp Lys Asn Thr Gly Asp  
 725 730 735  
 Tyr Tyr Glu Asp Ser Tyr Glu Asp Ile Ser Ala Tyr Leu Leu Ser Lys  
 740 745 750  
 Asn Asn Ala Ile Glu Pro Arg Ser Phe Ser Gln Asn Pro Pro Val Leu  
 755 760 765  
 Lys Arg His Gln Arg Glu Ile Thr Arg Thr Thr Leu Gln Ser Asp Gln  
 770 775 780  
 Glu Glu Ile Asp Tyr Asp Asp Thr Ile Ser Val Glu Met Lys Lys Glu  
 785 790 795 800  
 Asp Phe Asp Ile Tyr Asp Glu Asp Glu Asn Gln Ser Pro Arg Ser Phe  
 805 810 815  
 Gln Lys Lys Thr Arg His Tyr Phe Ile Ala Ala Val Glu Arg Leu Trp  
 820 825 830  
 Asp Tyr Gly Met Ser Ser Ser Pro His Val Leu Arg Asn Arg Ala Gln  
 835 840 845  
 Ser Gly Ser Val Pro Gln Phe Lys Lys Val Val Phe Gln Glu Phe Thr  
 850 855 860  
 Asp Gly Ser Phe Thr Gln Pro Leu Tyr Arg Gly Glu Leu Asn Glu His  
 865 870 875 880  
 Leu Gly Leu Leu Gly Pro Tyr Ile Arg Ala Glu Val Glu Asp Asn Ile  
 885 890 895  
 Met Val Thr Phe Arg Asn Gln Ala Ser Arg Pro Tyr Ser Phe Tyr Ser  
 900 905 910  
 Ser Leu Ile Ser Tyr Glu Glu Asp Gln Arg Gln Gly Ala Glu Pro Arg  
 915 920 925

# Sequence\_Listing

Lys Asn Phe Val Lys Pro Asn Glu Thr Lys Thr Tyr Phe Trp Lys Val  
 930 935 940  
 Gln His His Met Ala Pro Thr Lys Asp Glu Phe Asp Cys Lys Ala Trp  
 945 950 955 960  
 Ala Tyr Phe Ser Asp Val Asp Leu Glu Lys Asp Val His Ser Gly Leu  
 965 970 975  
 Ile Gly Pro Leu Leu Val Cys His Thr Asn Thr Leu Asn Pro Ala His  
 980 985 990  
 Gly Arg Gln Val Thr Val Gln Glu Phe Ala Leu Phe Phe Thr Ile Phe  
 995 1000 1005  
 Asp Glu Thr Lys Ser Trp Tyr Phe Thr Glu Asn Met Glu Arg Asn  
 1010 1015 1020  
 Cys Arg Ala Pro Cys Asn Ile Gln Met Glu Asp Pro Thr Phe Lys  
 1025 1030 1035  
 Glu Asn Tyr Arg Phe His Ala Ile Asn Gly Tyr Ile Met Asp Thr  
 1040 1045 1050  
 Leu Pro Gly Leu Val Met Ala Gln Asp Gln Arg Ile Arg Trp Tyr  
 1055 1060 1065  
 Leu Leu Ser Met Gly Ser Asn Glu Asn Ile His Ser Ile His Phe  
 1070 1075 1080  
 Ser Gly His Val Phe Thr Val Arg Lys Lys Glu Glu Tyr Lys Met  
 1085 1090 1095  
 Ala Leu Tyr Asn Leu Tyr Pro Gly Val Phe Glu Thr Val Glu Met  
 1100 1105 1110  
 Leu Pro Ser Lys Ala Gly Ile Trp Arg Val Glu Cys Leu Ile Gly  
 1115 1120 1125  
 Glu His Leu His Ala Gly Met Ser Thr Leu Phe Leu Val Tyr Ser  
 1130 1135 1140  
 Asn Lys Cys Gln Thr Pro Leu Gly Met Ala Ser Gly His Ile Arg  
 1145 1150 1155  
 Asp Phe Gln Ile Thr Ala Ser Gly Gln Tyr Gly Gln Trp Ala Pro  
 1160 1165 1170  
 Lys Leu Ala Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser  
 1175 1180 1185

# Sequence\_Listing

Thr	Lys	Glu	Pro	Phe	Ser	Trp	Ile	Lys	Val	Asp	Leu	Leu	Ala	Pro
	1190					1195					1200			
Met	Ile	Ile	His	Gly	Ile	Lys	Thr	Gln	Gly	Ala	Arg	Gln	Lys	Phe
	1205					1210					1215			
Ser	Ser	Leu	Tyr	Ile	Ser	Gln	Phe	Ile	Ile	Met	Tyr	Ser	Leu	Asp
	1220					1225					1230			
Gly	Lys	Lys	Trp	Gln	Thr	Tyr	Arg	Gly	Asn	Ser	Thr	Gly	Thr	Leu
	1235					1240					1245			
Met	Val	Phe	Phe	Gly	Asn	Val	Asp	Ser	Ser	Gly	Ile	Lys	His	Asn
	1250					1255					1260			
Ile	Phe	Asn	Pro	Pro	Ile	Ile	Ala	Arg	Tyr	Ile	Arg	Leu	His	Pro
	1265					1270					1275			
Thr	His	Tyr	Ser	Ile	Arg	Ser	Thr	Leu	Arg	Met	Glu	Leu	Met	Gly
	1280					1285					1290			
Cys	Asp	Leu	Asn	Ser	Cys	Ser	Met	Pro	Leu	Gly	Met	Glu	Ser	Lys
	1295					1300					1305			
Ala	Ile	Ser	Asp	Ala	Gln	Ile	Thr	Ala	Ser	Ser	Tyr	Phe	Thr	Asn
	1310					1315					1320			
Met	Phe	Ala	Thr	Trp	Ser	Pro	Ser	Lys	Ala	Arg	Leu	His	Leu	Gln
	1325					1330					1335			
Gly	Arg	Ser	Asn	Ala	Trp	Arg	Pro	Gln	Val	Asn	Asn	Pro	Lys	Glu
	1340					1345					1350			
Trp	Leu	Gln	Val	Asp	Phe	Gln	Lys	Thr	Met	Lys	Val	Thr	Gly	Val
	1355					1360					1365			
Thr	Thr	Gln	Gly	Val	Lys	Ser	Leu	Leu	Thr	Ser	Met	Tyr	Val	Lys
	1370					1375					1380			
Glu	Phe	Leu	Ile	Ser	Ser	Ser	Gln	Asp	Gly	His	Gln	Trp	Thr	Leu
	1385					1390					1395			
Phe	Phe	Gln	Asn	Gly	Lys	Val	Lys	Val	Phe	Gln	Gly	Asn	Gln	Asp
	1400					1405					1410			
Ser	Phe	Thr	Pro	Val	Val	Asn	Ser	Leu	Asp	Pro	Pro	Leu	Leu	Thr
	1415					1420					1425			
Arg	Tyr	Leu	Arg	Ile	His	Pro	Gln	Ser	Trp	Val	His	Gln	Ile	Ala
	1430					1435					1440			

# Sequence\_Listing

Leu Arg Met Glu Val Leu Gly Cys Glu Ala Gln Asp Leu Tyr  
 1445 1450 1455

<210> 4  
 <211> 4  
 <212> PRT  
 <213> Homo sapiens

<400> 4

Ser Phe Ser Gln  
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<210> 5  
 <211> 10  
 <212> PRT  
 <213> Homo sapiens

<400> 5

Asn Pro Pro Val Leu Lys Arg His Gln Arg  
 1 5 10