

Coagulation factor VIII

Name	Coagulation factor VIII
Synonyms	<ul style="list-style-type: none"> • AHF • Antihemophilic factor • F8C • Procoagulant component
Gene Name	F8
Organism	Human
Amino acid sequence	<pre>>1c1 BSEQ0000032 Coagulation factor VIII MQIELSTCFFLCLLRFCFSATRRYYLGAVELSWDYMQSDLGELPVDARFPPRPVKSFFPN TSVVYKKTLFVEFTDHLFNIAPRPPWMGLLGPTIQAEVYDVTVVITLKNMASHPVSLHAV GVSYWKASEGAEYDDQTSQREKEDDKVFPGGSHTYVWQVLKENGPMASDPLCLTYSYLSH VDLVKDLNSGLIGALLVCREGLAKEKTQTLHKFILLFAVFDEGKSWHSETKNSLMQDRD AASARAWPKMHTVNGYVNRSLPGLIGCHRKSVYWHVIGMGTTPEVHSIFLEGHTFLVRNH RQASLEISPIITFLTAQTLMLDLGQFLLFCHISSHQHDGMEAYVKVDSCPEEPQLRMKNNE EAEDYDDDLTDSEMDVVRFDDDNSPSFIQIRSVAKKHPKTWVHYIAAEEEDWDYAPLVLA PDDRSYKSQYLNNGPQRIGRKYKVRFMAYTDETFKTREAIQHESGILGPLYGEVGDTL LIIFKNQASRPYNIYPHGITDVRPLYSRRLPKGVKHLKDFPILPGEIFKYKWTVTVEDGP TKSDPRCLTRYYSFVNMRDLASGLIGPLLICYKESVDQRGNQIMSDKRNVILFSVFDE NRSWYLTENIQRFLPNPAGVQLEDPEFQASNIMHSINGYVFDSLQLSVCLHEVAYWYILS IGAQTDFLSVFFSGYTFKHKMVYEDTLTFPFSGETVFMSMENPGLWILGCHNSDFRNRG MTALLKVSSCDKNTGDYYEDSYEDISAYLLSKNNAIEPRFSQNSRHPSTRQKQFNATTI PENDIEKTDWPFAHRTMPKIQNVSSDLLMLLRQSPTPHGLSLSDLQEAKYETFSDDPS PGAIDSNNLSLSEMTFRPQLHHSMDMVFTPESEGLQLRLNEKLGTTAATELKKLDFKVSST</pre>

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 EVLGCEAQDLY

Number of residues

2351

Molecular Weight

267007.42

Theoretical pI

7.37

GO Classification	<p>Functions</p> <ul style="list-style-type: none">• oxidoreductase activity• copper ion binding <p>Processes</p> <ul style="list-style-type: none">• blood coagulation• COPII vesicle coating• ER to Golgi vesicle-mediated transport• membrane organization• platelet degranulation• platelet activation• post-translational protein modification• protein N-linked glycosylation via asparagine• cellular protein metabolic process• acute-phase response• blood coagulation, intrinsic pathway <p>Components</p> <ul style="list-style-type: none">• extracellular region• plasma membrane• extracellular space• endoplasmic reticulum lumen• endoplasmic reticulum-Golgi intermediate compartment membrane• ER to Golgi transport vesicle• Golgi membrane• platelet alpha granule lumen
General Function	Oxidoreductase activity

Specific Function	Factor VIII, along with calcium and phospholipid, acts as a cofactor for factor IXa when it converts factor X to the activated form, factor Xa.
Pfam Domain Function	<ul style="list-style-type: none"> • Cu-oxidase (PF00394 🔗 (http://pfam.sanger.ac.uk/family?acc=PF00394)) • Cu-oxidase_2 (PF07731 🔗 (http://pfam.sanger.ac.uk/family?acc=PF07731)) • Cu-oxidase_3 (PF07732 🔗 (http://pfam.sanger.ac.uk/family?acc=PF07732)) • F5_F8_type_C (PF00754 🔗 (http://pfam.sanger.ac.uk/family?acc=PF00754))
Transmembrane Regions	Not Available
GenBank Protein ID	182818 🔗 (http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?val=182818)
UniProtKB ID	P00451 🔗 (http://www.uniprot.org/uniprot/P00451)
UniProtKB Entry Name	FA8_HUMAN 🔗 (http://www.uniprot.org/uniprot/FA8_HUMAN)
Cellular Location	Secreted
Gene sequence	<pre>>1c1 BSEQ0015996 Coagulation factor VIII (F8) ATGCAAATAGAGCTCTCCACCTGCTTCTTTCTGTGCCTTTTGCATTCTGCTTTAGTGCC ACCAGAAAGATACTACCTGGGTGCAGTGGAACTGTCATGGGACTATATGCAAAGTGATCTC GGTGAGCTGCCTGTGGACGCAAGATTTCTCCTAGAGTGCCAAAATCTTTTCCATTCAAC ACCTCAGTCGTGTACAAAAAGACTCTGTTTGTAGAATTCACGGATCACCTTTTCAACATC GCTAAGCCAAGGCCACCCTGGATGGTCTGCTAGGTCTACCATCCAGGCTGAGGTTTAT GATACAGTGGTCATTACACTTAAGAACATGGCTTCCCATCCTGTCAGTCTTCATGCTGTT GGTGTATCCTACTGGAAAGCTTCTGAGGGAGCTGAATATGATGATCAGACCAGTCAAAGG GAGAAAGAAGATGATAAAGTCTTCCCTGGTGGAAAGCCATACATATGTCTGGCAGGTCTCG AAAGAGAATGGTCCAATGGCCTCTGACCCACTGTGCCTTACCTACTCATATCTTCTCAT GTGGACCTGGTAAAAGACTTGAATTCAGGCCTCATTGGAGCCCTACTAGTATGTAGAGAA GGGAGTCTGGCCAAGGAAAAGACACAGACCTTGACAAAATTTATACTACTTTTTGCTGTA TTTGATGAAGGGAAAAGTTGGCACTCAGAAACAAAGAACTCCTTGATGCAGGATAGGGAT GCTGCATCTGCTCGGCCTGGCCTAAAATGCACACAGTCAATGGTTATGTAACAGGTCT CTGCCAGTCTGATTGGATGCCACAGGAAATCAGTCTATTGGCATGTGATTGGAATGGGC ACCACTCCTGAAGTGAATCAATATTCCTCGAAGGTCACACATTTCTTGAGGAACCAT CGCCAGGCGTCCTTGGAAATCTCGCAATAAATTTCTTACTGCTCAAACACTCTTGATG GACCTTGGACAGTTTCTACTGTTTTGTCATATCTTCCCACCAACATGATGGCATGGAA GCTTATGTCAAAGTAGACAGCTGTCCAGAGGAACCCCAACTACGAATGAAAAATAATGAA GAAGCGGAAGACTATGATGATGATCTTACTGATTCTGAAATGGATGTGGTCAGGTTTGAT GATGACAACTCTCCTTCTTTATCCAAATTCGCTCAGTTGCCAAGAAGCATCCTAAAAT</pre>

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GenBank Gene ID	M14113 🔗 (http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?val=M14113)
GeneCard ID	Not Available
GenAtlas ID	F8 🔗 (http://www.dsi.univ-paris5.fr/genatlas/fiche1.php?symbol=F8)
HGNC ID	HGNC:3546 🔗 (http://www.genenames.org/data/hgnc_data.php?hgnc_id=HGNC:3546)
Chromosome Location	X
Locus	Xq28
References	<ol style="list-style-type: none"> 1. Truett MA, Blacher R, Burke RL, Caput D, Chu C, Dina D, Hartog K, Kuo CH, Masiarz FR, Merryweather JP, et al.: Characterization of the polypeptide composition of human factor VIII:C and the nucleotide sequence and expression of the human kidney cDNA. <i>DNA</i>. 1985 Oct;4(5):333-49. 3935400 🔗 (http://www.ncbi.nlm.nih.gov/pubmed/3935400) 2. Wood WI, Capon DJ, Simonsen CC, Eaton DL, Gitschier J, Keyt B, Seeburg PH, Smith DH, Hollingshead P, Wion KL, Delwart E, Tuddenham EG, Vehar GA, Lawn RM: Expression of active human factor VIII from recombinant DNA clones. <i>Nature</i>. 1984 Nov 22-28;312(5992):330-7. 6438526 🔗 (http://www.ncbi.nlm.nih.gov/pubmed/6438526) 3. Levinson B, Kenwick S, Gamel P, Fisher K, Gitschier J: Evidence for a third transcript from the human factor VIII gene. <i>Genomics</i>. 1992 Nov;14(3):585-9. 1427887 🔗 (http://www.ncbi.nlm.nih.gov/pubmed/1427887)

4. Toole JJ, Knopf JL, Wozney JM, Sultzman LA, Buecker JL, Pittman DD, Kaufman RJ, Brown E, Shoemaker C, Orr EC, et al.: Molecular cloning of a cDNA encoding human antihemophilic factor. *Nature*. 1984 Nov 22-28;312(5992):342-7. 6438528  (<http://www.ncbi.nlm.nih.gov/pubmed/6438528>)
5. Gitschier J, Wood WI: Sequence of the exon-containing regions of the human factor VIII gene. *Hum Mol Genet*. 1992 Jun;1(3):199-200. 1303178  (<http://www.ncbi.nlm.nih.gov/pubmed/1303178>)
6. Ota T, Suzuki Y, Nishikawa T, Otsuki T, Sugiyama T, Irie R, Wakamatsu A, Hayashi K, Sato H, Nagai K, Kimura K, Makita H, Sekine M, Obayashi M, Nishi T, Shibahara T, Tanaka T, Ishii S, Yamamoto J, Saito K, Kawai Y, Isono Y, Nakamura Y, Nagahari K, Murakami K, Yasuda T, Iwayanagi T, Wagatsuma M, Shiratori A, Sudo H, Hosoi T, Kaku Y, Kodaira H, Kondo H, Sugawara M, Takahashi M, Kanda K, Yokoi T, Furuya T, Kikkawa E, Omura Y, Abe K, Kamihara K, Katsuta N, Sato K, Tanikawa M, Yamazaki M, Ninomiya K, Ishibashi T, Yamashita H, Murakawa K, Fujimori K, Tanai H, Kimata M, Watanabe M, Hiraoka S, Chiba Y, Ishida S, Ono Y, Takiguchi S, Watanabe S, Yosida M, Hotuta T, Kusano J, Kanehori K, Takahashi-Fujii A, Hara H, Tanase TO, Nomura Y, Togiya S, Komai F, Hara R, Takeuchi K, Arita M, Imose N, Musashino K, Yuuki H, Oshima A, Sasaki N, Aotsuka S, Yoshikawa Y, Matsunawa H, Ichihara T, Shiohata N, Sano S, Moriya S, Momiyama H, Satoh N, Takami S, Terashima Y, Suzuki O, Nakagawa S, Senoh A, Mizoguchi H, Goto Y, Shimizu F, Wakebe H, Hishigaki H, Watanabe T, Sugiyama A, Takemoto M, Kawakami B, Yamazaki M, Watanabe K, Kumagai A, Itakura S, Fukuzumi Y, Fujimori Y, Komiyama M, Tashiro H, Tanigami A, Fujiwara T, Ono T, Yamada K, Fujii Y, Ozaki K, Hirao M, Ohmori Y, Kawabata A, Hikiji T, Kobatake N, Inagaki H, Ikema Y, Okamoto S, Okitani R, Kawakami T, Noguchi S, Itoh T, Shigeta K, Senba T, Matsumura K, Nakajima Y, Mizuno T, Morinaga M, Sasaki M, Togashi T, Oyama M, Hata H, Watanabe M, Komatsu T, Mizushima-Sugano J, Satoh T, Shirai Y, Takahashi Y, Nakagawa K, Okumura K, Nagase T, Nomura N, Kikuchi H, Masuho Y, Yamashita R, Nakai K, Yada T, Nakamura Y, Ohara O, Isogai T, Sugano S: Complete sequencing and characterization of 21,243 full-length human cDNAs. *Nat Genet*. 2004 Jan;36(1):40-5. Epub 2003 Dec 21. 14702039  (<http://www.ncbi.nlm.nih.gov/pubmed/14702039>)
7. Ross MT, Grafham DV, Coffey AJ, Scherer S, McLay K, Muzny D, Platzer M, Howell GR, Burrows C, Bird CP, Frankish A, Lovell FL, Howe KL, Ashurst JL, Fulton RS, Sudbrak R, Wen G, Jones MC, Hurler ME, Andrews TD, Scott CE, Searle S, Ramser J, Whittaker A, Deadman R, Carter NP, Hunt SE, Chen R, Cree A, Gunaratne P, Havlak P, Hodgson A, Metzker ML, Richards S, Scott G, Steffen D, Sodergren E, Wheeler DA, Worley KC, Ainscough R, Ambrose KD, Ansari-Lari MA, Aradhya S, Ashwell RI, Babbage AK, Bagguley CL, Ballabio A, Banerjee R, Barker GE, Barlow KF, Barrett IP, Bates KN, Beare DM, Beasley H, Beasley O, Beck A, Bethel G, Blechschmidt K, Brady N, Bray-Allen S, Bridgeman AM, Brown AJ, Brown MJ, Bonnin D, Bruford EA, Buhay C, Burch P, Burford D, Burgess J, Burrill W, Burton J, Bye JM, Carder C, Carrel L, Chako J, Chapman JC, Chavez D, Chen E, Chen G, Chen Y, Chen Z, Chinault C, Ciccodicola A, Clark SY, Clarke G, Clee CM, Clegg S, Clerc-Blankenburg K, Clifford K, Copley V, Cole CG, Conquer JS, Corby N, Connor RE, David R, Davies J, Davis C, Davis J, Delgado O, Deshazo D, Dhami P, Ding Y, Dinh H, Dodsworth S, Draper H, Dugan-Rocha S, Dunham A, Dunn M, Durbin KJ, Dutta I, Eades T, Ellwood M, Emery-Cohen A, Errington H, Evans KL, Faulkner L, Francis F, Frankland J, Fraser AE, Galgoczy P, Gilbert J, Gill R, Glockner G, Gregory SG, Gribble S, Griffiths C, Grocock R, Gu Y, Gwilliam R, Hamilton C, Hart EA, Hawes A, Heath PD, Heitmann K, Hennig S, Hernandez J, Hinzmann B, Ho S, Hoffs M, Howden PJ, Huckle EJ, Hume J, Hunt PJ, Hunt AR, Isherwood J, Jacob L, Johnson D, Jones S, de Jong PJ, Joseph SS, Keenan S, Kelly S, Kershaw JK, Khan Z, Kioschis P, Klages S, Knights AJ, Kosiura A, Kovar-Smith C, Laird GK, Langford C, Lawlor S, Leversha M, Lewis L, Liu W, Lloyd C, Lloyd DM, Louseged H, Loveland JE, Lovell JD, Lozado R, Lu J, Lyne R, Ma J, Maheshwari M, Matthews LH, McDowall J, McLaren S, McMurray A, Meidl P, Meitinger T, Milne S, Miner G, Mistry SL, Morgan M, Morris S, Muller I, Mullikin JC, Nguyen N, Nordsiek G, Nyakatura G, O'Dell CN, Okwuonu G, Palmer S, Pandian R, Parker D, Parrish J, Pasternak S, Patel D, Pearce AV, Pearson DM, Pelan SE, Perez L, Porter KM, Ramsey Y, Reichwald K, Rhodes S, Ridler KA, Schlessinger D, Schueler MG, Sehra HK, Shaw-Smith C, Shen H, Sheridan EM, Shownkeen R, Skuce CD, Smith ML, Sotheran EC, Steingruber HE, Steward CA, Storey R, Swann RM, Swarbreck D, Tabor PE, Taudien S, Taylor T, Teague B, Thomas K, Thorpe A, Timms K, Tracey A, Trevanion S, Tromans AC, d'Urso M, Verduzco D, Villasana D, Waldron L, Wall M, Wang Q, Warren J, Warry GL, Wei X, West A, Whitehead SL, Whiteley MN, Wilkinson JE, Willey DL, Williams G, Williams L, Williamson A, Williamson H, Wilming L, Woodmansey RL, Wray PW, Yen J, Zhang J, Zhou J, Zoghbi H, Zorilla S, Buck D, Reinhardt R, Poustka A, Rosenthal A, Lehrach H, Meindl A, Minx PJ, Hillier LW, Willard HF, Wilson RK, Waterston RH, Rice CM, Vaudin M, Coulson A, Nelson DL, Weinstock G, Sulston JE, Durbin R, Hubbard T, Gibbs RA, Beck S, Rogers J, Bentley DR: The DNA sequence of the human X chromosome. *Nature*. 2005 Mar 17;434(7031):325-37. 15772651  (<http://www.ncbi.nlm.nih.gov/pubmed/15772651>)

8. Gerhard DS, Wagner L, Feingold EA, Shenmen CM, Grouse LH, Schuler G, Klein SL, Old S, Rasooly R, Good P, Guyer M, Peck AM, Derge JG, Lipman D, Collins FS, Jang W, Sherry S, Feolo M, Misquitta L, Lee E, Rotmistrovsky K, Greenhut SF, Schaefer CF, Buetow K, Bonner TI, Haussler D, Kent J, Kiekhaus M, Furey T, Brent M, Prange C, Schreiber K, Shapiro N, Bhat NK, Hopkins RF, Hsie F, Driscoll T, Soares MB, Casavant TL, Scheetz TE, Brown-stein MJ, Usdin TB, Toshiyuki S, Carninci P, Piao Y, Dudekula DB, Ko MS, Kawakami K, Suzuki Y, Sugano S, Gruber CE, Smith MR, Simmons B, Moore T, Waterman R, Johnson SL, Ruan Y, Wei CL, Mathavan S, Gunaratne PH, Wu J, Garcia AM, Hulyk SW, Fuh E, Yuan Y, Sneed A, Kowis C, Hodgson A, Muzny DM, McPherson J, Gibbs RA, Fahey J, Helton E, Kettelman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Madari A, Young AC, Wetherby KD, Granite SJ, Kwong PN, Brinkley CP, Pearson RL, Bouffard GG, Blakesly RW, Green ED, Dickson MC, Rodriguez AC, Grimwood J, Schmutz J, Myers RM, Butterfield YS, Griffith M, Griffith OL, Krzywinski MI, Liao N, Morin R, Palmquist D, Petrescu AS, Skalska U, Smailus DE, Stott JM, Schnerch A, Schein JE, Jones SJ, Holt RA, Baross A, Marra MA, Clifton S, Makowski KA, Bosak S, Malek J: The status, quality, and expansion of the NIH full-length cDNA project: the Mammalian Gene Collection (MGC). *Genome Res.* 2004 Oct;14(10B):2121-7. 15489334 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/15489334) (<http://www.ncbi.nlm.nih.gov/pubmed/15489334>)
9. Severs JC, Carnine M, Eguizabal H, Mock KK: Characterization of tyrosine sulfate residues in antihemophilic recombinant factor VIII by liquid chromatography electrospray ionization tandem mass spectrometry and amino acid analysis. *Rapid Commun Mass Spectrom.* 1999;13(11):1016-23. 10368977 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10368977) (<http://www.ncbi.nlm.nih.gov/pubmed/10368977>)
10. Leyte A, van Schijndel HB, Niehrs C, Huttner WB, Verbeet MP, Mertens K, van Mourik JA: Sulfation of Tyr1680 of human blood coagulation factor VIII is essential for the interaction of factor VIII with von Willebrand factor. *J Biol Chem.* 1991 Jan 15;266(2):740-6. 1898735 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1898735) (<http://www.ncbi.nlm.nih.gov/pubmed/1898735>)
11. Pittman DD, Wang JH, Kaufman RJ: Identification and functional importance of tyrosine sulfate residues within recombinant factor VIII. *Biochemistry.* 1992 Apr 7;31(13):3315-25. 1554716 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1554716) (<http://www.ncbi.nlm.nih.gov/pubmed/1554716>)
12. Saenko EL, Scandella D: The acidic region of the factor VIII light chain and the C2 domain together form the high affinity binding site for von willebrand factor. *J Biol Chem.* 1997 Jul 18;272(29):18007-14. 9218428 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9218428) (<http://www.ncbi.nlm.nih.gov/pubmed/9218428>)
13. McMullen BA, Fujikawa K, Davie EW, Hedner U, Ezban M: Locations of disulfide bonds and free cysteines in the heavy and light chains of recombinant human factor VIII (antihemophilic factor A). *Protein Sci.* 1995 Apr;4(4):740-6. 7613471 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7613471) (<http://www.ncbi.nlm.nih.gov/pubmed/7613471>)
14. Liu T, Qian WJ, Gritsenko MA, Camp DG 2nd, Monroe ME, Moore RJ, Smith RD: Human plasma N-glycoproteome analysis by immunoaffinity subtraction, hydrazide chemistry, and mass spectrometry. *J Proteome Res.* 2005 Nov-Dec;4(6):2070-80. 16335952 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/16335952) (<http://www.ncbi.nlm.nih.gov/pubmed/16335952>)
15. Gilbert GE, Baleja JD: Membrane-binding peptide from the C2 domain of factor VIII forms an amphipathic structure as determined by NMR spectroscopy. *Biochemistry.* 1995 Mar 7;34(9):3022-31. 7893714 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7893714) (<http://www.ncbi.nlm.nih.gov/pubmed/7893714>)
16. Gitschier J: The molecular basis of hemophilia A. *Ann N Y Acad Sci.* 1991;614:89-96. 1902642 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1902642) (<http://www.ncbi.nlm.nih.gov/pubmed/1902642>)
17. White GC 2nd, Shoemaker CB: Factor VIII gene and hemophilia A. *Blood.* 1989 Jan;73(1):1-12. 2491949 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2491949) (<http://www.ncbi.nlm.nih.gov/pubmed/2491949>)
18. Antonarakis SE, Kazazian HH, Tuddenham EG: Molecular etiology of factor VIII deficiency in hemophilia A. *Hum Mutat.* 1995;5(1):1-22. 7728145 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7728145) (<http://www.ncbi.nlm.nih.gov/pubmed/7728145>)
19. Gitschier J, Wood WI, Shuman MA, Lawn RM: Identification of a missense mutation in the factor VIII gene of a mild hemophiliac. *Science.* 1986 Jun 13;232(4756):1415-6. 3012775 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/3012775) (<http://www.ncbi.nlm.nih.gov/pubmed/3012775>)
20. Levinson B, Janco R, Phillips J 3rd, Gitschier J: A novel missense mutation in the factor VIII gene identified by analysis of amplified hemophilia DNA sequences. *Nucleic Acids Res.* 1987 Dec 10;15(23):9797-805. 3122181 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/3122181) (<http://www.ncbi.nlm.nih.gov/pubmed/3122181>)
21. Youssoufian H, Antonarakis SE, Bell W, Griffin AM, Kazazian HH Jr: Nonsense and missense mutations in hemophilia A: estimate of the relative mutation rate at CG dinucleotides. *Am J Hum Genet.* 1988 May;42(5):718-25. 2833855 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2833855) (<http://www.ncbi.nlm.nih.gov/pubmed/2833855>)
22. Youssoufian H, Wong C, Aronis S, Platokoukis H, Kazazian HH Jr, Antonarakis SE: Moderately severe hemophilia A resulting from Glu----Gly substitution in exon 7 of the factor VIII gene. *Am J Hum Genet.* 1988 Jun;42(6):867-71. 2835904 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2835904) (<http://www.ncbi.nlm.nih.gov/pubmed/2835904>)

23. O'Brien DP, Tuddenham EG: Purification and characterization of factor VIII 1,689-Cys: a nonfunctional cofactor occurring in a patient with severe hemophilia A. *Blood*. 1989 Jun;73(8):2117-22. 2499363 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2499363) (http://www.ncbi.nlm.nih.gov/pubmed/2499363)
24. Shima M, Ware J, Yoshioka A, Fukui H, Fulcher CA: An arginine to cysteine amino acid substitution at a critical thrombin cleavage site in a dysfunctional factor VIII molecule. *Blood*. 1989 Oct;74(5):1612-7. 2506948 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2506948) (http://www.ncbi.nlm.nih.gov/pubmed/2506948)
25. Chan V, Chan TK, Tong TM, Todd D: A novel missense mutation in exon 4 of the factor VIII:C gene resulting in moderately severe hemophilia A. *Blood*. 1989 Dec;74(8):2688-91. 2510835 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2510835) (http://www.ncbi.nlm.nih.gov/pubmed/2510835)
26. Inaba H, Fujimaki M, Kazazian HH Jr, Antonarakis SE: Mild hemophilia A resulting from Arg-to-Leu substitution in exon 26 of the factor VIII gene. *Hum Genet*. 1989 Mar;81(4):335-8. 2495245 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2495245) (http://www.ncbi.nlm.nih.gov/pubmed/2495245)
27. Arai M, Inaba H, Higuchi M, Antonarakis SE, Kazazian HH Jr, Fujimaki M, Hoyer LW: Direct characterization of factor VIII in plasma: detection of a mutation altering a thrombin cleavage site (arginine-372---histidine). *Proc Natl Acad Sci U S A*. 1989 Jun;86(11):4277-81. 2498882 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2498882) (http://www.ncbi.nlm.nih.gov/pubmed/2498882)
28. Arai M, Higuchi M, Antonarakis SE, Kazazian HH Jr, Phillips JA 3rd, Janco RL, Hoyer LW: Characterization of a thrombin cleavage site mutation (Arg 1689 to Cys) in the factor VIII gene of two unrelated patients with cross-reacting material-positive hemophilia A. *Blood*. 1990 Jan 15;75(2):384-9. 2104766 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2104766) (http://www.ncbi.nlm.nih.gov/pubmed/2104766)
29. Casula L, Murru S, Pecorara M, Ristaldi MS, Restagno G, Mancuso G, Morfini M, De Biasi R, Baudo F, Carbonara A, et al.: Recurrent mutations and three novel rearrangements in the factor VIII gene of hemophilia A patients of Italian descent. *Blood*. 1990 Feb 1;75(3):662-70. 2105106 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2105106) (http://www.ncbi.nlm.nih.gov/pubmed/2105106)
30. Pattinson JK, McVey JH, Boon M, Ajani A, Tuddenham EG: CRM+ haemophilia A due to a missense mutation (372---Cys) at the internal heavy chain thrombin cleavage site. *Br J Haematol*. 1990 May;75(1):73-7. 1973901 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1973901) (http://www.ncbi.nlm.nih.gov/pubmed/1973901)
31. Higuchi M, Wong C, Kochhan L, Olek K, Aronis S, Kasper CK, Kazazian HH Jr, Antonarakis SE: Characterization of mutations in the factor VIII gene by direct sequencing of amplified genomic DNA. *Genomics*. 1990 Jan;6(1):65-71. 2105906 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2105906) (http://www.ncbi.nlm.nih.gov/pubmed/2105906)
32. Traystman MD, Higuchi M, Kasper CK, Antonarakis SE, Kazazian HH Jr: Use of denaturing gradient gel electrophoresis to detect point mutations in the factor VIII gene. *Genomics*. 1990 Feb;6(2):293-301. 2106480 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2106480) (http://www.ncbi.nlm.nih.gov/pubmed/2106480)
33. Kogan S, Gitschier J: Mutations and a polymorphism in the factor VIII gene discovered by denaturing gradient gel electrophoresis. *Proc Natl Acad Sci U S A*. 1990 Mar;87(6):2092-6. 2107542 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/2107542) (http://www.ncbi.nlm.nih.gov/pubmed/2107542)
34. Paynton C, Sarkar G, Sommer SS: Identification of mutations in two families with sporadic hemophilia A. *Hum Genet*. 1991 Aug;87(4):397-400. 1908817 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1908817) (http://www.ncbi.nlm.nih.gov/pubmed/1908817)
35. Higuchi M, Kazazian HH Jr, Kasch L, Warren TC, McGinniss MJ, Phillips JA 3rd, Kasper C, Janco R, Antonarakis SE: Molecular characterization of severe hemophilia A suggests that about half the mutations are not within the coding regions and splice junctions of the factor VIII gene. *Proc Natl Acad Sci U S A*. 1991 Aug 15;88(16):7405-9. 1908096 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1908096) (http://www.ncbi.nlm.nih.gov/pubmed/1908096)
36. Higuchi M, Antonarakis SE, Kasch L, Oldenburg J, Economou-Petersen E, Olek K, Arai M, Inaba H, Kazazian HH Jr: Molecular characterization of mild-to-moderate hemophilia A: detection of the mutation in 25 of 29 patients by denaturing gradient gel electrophoresis. *Proc Natl Acad Sci U S A*. 1991 Oct 1;88(19):8307-11. 1924291 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1924291) (http://www.ncbi.nlm.nih.gov/pubmed/1924291)
37. Schwaab R, Ludwig M, Kochhan L, Oldenburg J, McVey JH, Egli H, Brackmann HH, Olek K: Detection and characterisation of two missense mutations at a cleavage site in the factor VIII light chain. *Thromb Res*. 1991 Feb 1;61(3):225-34. 1851341 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1851341) (http://www.ncbi.nlm.nih.gov/pubmed/1851341)
38. Krepelova A, Vorlova Z, Aquila M, Mori P: GAA(Glu)272---AAA(Lys) and CGA(Arg)1941---CAA(Gln) in the factor VIII gene in two haemophilia A patients of Czech origin. *Br J Haematol*. 1992 Jul;81(3):458. 1356412 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1356412) (http://www.ncbi.nlm.nih.gov/pubmed/1356412)

39. Economou EP, Kazazian HH Jr, Antonarakis SE: Detection of mutations in the factor VIII gene using single-stranded conformational polymorphism (SSCP). *Genomics*. 1992 Jul;13(3):909-11. 1639429 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1639429) (<http://www.ncbi.nlm.nih.gov/pubmed/1639429>)
40. Reiner AP, Thompson AR: Screening for nonsense mutations in patients with severe hemophilia A can provide rapid, direct carrier detection. *Hum Genet*. 1992 Apr;89(1):88-94. 1349567 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1349567) (<http://www.ncbi.nlm.nih.gov/pubmed/1349567>)
41. Nafa K, Baudis M, Deburgrave N, Bardin JM, Sultan Y, Kaplan JC, Delpech M: A novel mutation (Arg-->Leu in exon 18) in factor VIII gene responsible for moderate hemophilia A. *Hum Mutat*. 1992;1(1):77-8. 1301194 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1301194) (<http://www.ncbi.nlm.nih.gov/pubmed/1301194>)
42. Diamond C, Kogan S, Levinson B, Gitschier J: Amino acid substitutions in conserved domains of factor VIII and related proteins: study of patients with mild and moderately severe hemophilia A. *Hum Mutat*. 1992;1(3):248-57. 1301932 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1301932) (<http://www.ncbi.nlm.nih.gov/pubmed/1301932>)
43. Jonsdottir S, Diamond C, Levinson B, Magnusson S, Jensson O, Gitschier J: Missense mutations causing mild hemophilia A in Iceland detected by denaturing gradient gel electrophoresis. *Hum Mutat*. 1992;1(6):506-8. 1301960 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/1301960) (<http://www.ncbi.nlm.nih.gov/pubmed/1301960>)
44. McGinniss MJ, Kazazian HH Jr, Hoyer LW, Bi L, Inaba H, Antonarakis SE: Spectrum of mutations in CRM-positive and CRM-reduced hemophilia A. *Genomics*. 1993 Feb;15(2):392-8. 8449505 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/8449505) (<http://www.ncbi.nlm.nih.gov/pubmed/8449505>)
45. Pieneman WC, Reitsma PH, Briet E: Double strand conformation polymorphism (DSCP) detects two point mutations at codon 280 (AAC-->ATC) and at codon 431 (TAC-->AAC) of the blood coagulation factor VIII gene. *Thromb Haemost*. 1993 May 3;69(5):473-5. 8322269 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/8322269) (<http://www.ncbi.nlm.nih.gov/pubmed/8322269>)
46. Arruda VR, Pieneman WC, Reitsma PH, Deutz-Terlouw PP, Annichino-Bizzacchi JM, Briet E, Costa FF: Eleven novel mutations in the factor VIII gene from Brazilian hemophilia A patients. *Blood*. 1995 Oct 15;86(8):3015-20. 7579394 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7579394) (<http://www.ncbi.nlm.nih.gov/pubmed/7579394>)
47. Pieneman WC, Deutz-Terlouw PP, Reitsma PH, Briet E: Screening for mutations in haemophilia A patients by multiplex PCR-SSCP, Southern blotting and RNA analysis: the detection of a genetic abnormality in the factor VIII gene in 30 out of 35 patients. *Br J Haematol*. 1995 Jun;90(2):442-9. 7794769 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7794769) (<http://www.ncbi.nlm.nih.gov/pubmed/7794769>)
48. Bidichandani SI, Lanyon WG, Shiach CR, Lowe GD, Connor JM: Detection of mutations in ectopic factor VIII transcripts from nine haemophilia A patients and the correlation with phenotype. *Hum Genet*. 1995 May;95(5):531-8. 7759074 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/7759074) (<http://www.ncbi.nlm.nih.gov/pubmed/7759074>)
49. Becker J, Schwaab R, Moller-Taube A, Schwaab U, Schmidt W, Brackmann HH, Grimm T, Olek K, Oldenburg J: Characterization of the factor VIII defect in 147 patients with sporadic hemophilia A: family studies indicate a mutation type-dependent sex ratio of mutation frequencies. *Am J Hum Genet*. 1996 Apr;58(4):657-70. 8644728 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/8644728) (<http://www.ncbi.nlm.nih.gov/pubmed/8644728>)
50. Chan V, Pang A, Chan TP, Chan VW, Chan TK: Molecular characterization of haemophilia A in southern Chinese. *Br J Haematol*. 1996 May;93(2):451-6. 8639447 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/8639447) (<http://www.ncbi.nlm.nih.gov/pubmed/8639447>)
51. Rudzki Z, Duncan EM, Casey GJ, Neumann M, Favaloro EJ, Lloyd JV: Mutations in a subgroup of patients with mild haemophilia A and a familial discrepancy between the one-stage and two-stage factor VIII:C methods. *Br J Haematol*. 1996 Aug;94(2):400-6. 8759905 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/8759905) (<http://www.ncbi.nlm.nih.gov/pubmed/8759905>)
52. Mazurier C, Gaucher C, Jorieux S, Parquet-Gernez A: Mutations in the FVIII gene in seven families with mild haemophilia A. *Br J Haematol*. 1997 Feb;96(2):426-7. 9029040 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9029040) (<http://www.ncbi.nlm.nih.gov/pubmed/9029040>)
53. Morichika S, Shima M, Kamisue S, Tanaka I, Imanaka Y, Suzuki H, Shibata H, Pemberton S, Gale K, McVey J, Tuddenham EG, Yoshioka A: Factor VIII gene analysis in Japanese CRM-positive and CRM-reduced haemophilia A patients by single-strand conformation polymorphism. *Br J Haematol*. 1997 Sep;98(4):901-6. 9326186 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9326186) (<http://www.ncbi.nlm.nih.gov/pubmed/9326186>)
54. Tavassoli K, Eigel A, Pollmann H, Horst J: Mutational analysis of ectopic factor VIII transcripts from hemophilia A patients: identification of cryptic splice site, exon skipping and novel point mutations. *Hum Genet*. 1997 Oct;100(5-6):508-11. 9341862 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9341862) (<http://www.ncbi.nlm.nih.gov/pubmed/9341862>)
55. Liu M, Murphy ME, Thompson AR: A domain mutations in 65 haemophilia A families and molecular modelling of dysfunctional factor VIII proteins. *Br J Haematol*. 1998 Dec;103(4):1051-60. 9886318 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9886318) (<http://www.ncbi.nlm.nih.gov/pubmed/9886318>)

56. Maugard C, Tuffery S, Aguilar-Martinez P, Schved JF, Gris JC, Demaille J, Claustres M: Protein truncation test: detection of severe haemophilia a mutation and analysis of factor VIII transcripts. *Hum Mutat.* 1998;11(1):18-22. 9450898 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9450898) (http://www.ncbi.nlm.nih.gov/pubmed/9450898)
57. Theophilus BD, Enayat MS, Higuchi M, Kazazian HH, Antonarakis SE, Hill FG: Independent occurrence of the novel Arg2163 to His mutation in the factor VIII gene in three unrelated families with haemophilia A with different phenotypes. *Mutations in brief no. 126. Online. Hum Mutat.* 1998;11(4):334. 10215414 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10215414) (http://www.ncbi.nlm.nih.gov/pubmed/10215414)
58. Freson K, Peerlinck K, Aguirre T, Arnout J, Vermylen J, Cassiman JJ, Matthijs G: Fluorescent chemical cleavage of mismatches for efficient screening of the factor VIII gene. *Hum Mutat.* 1998;11(6):470-9. 9603440 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9603440) (http://www.ncbi.nlm.nih.gov/pubmed/9603440)
59. Tavassoli K, Eigel A, Dworniczak B, Valtseva E, Horst J: Identification of four novel mutations in the factor VIII gene: three missense mutations (E1875G, G2088S, I2185T) and a 2-bp deletion (1780delTC). *Hum Mutat.* 1998;Suppl 1:S260-2. 9452104 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9452104) (http://www.ncbi.nlm.nih.gov/pubmed/9452104)
60. Tavassoli K, Eigel A, Wilke K, Pollmann H, Horst J: Molecular diagnostics of 15 hemophilia A patients: characterization of eight novel mutations in the factor VIII gene, two of which result in exon skipping. *Hum Mutat.* 1998;12(5):301-3. 9792405 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9792405) (http://www.ncbi.nlm.nih.gov/pubmed/9792405)
61. Laprise SL, Mak EK, Killoran KA, Layman LC, Gray MR: Use of denaturing gradient gel blots to screen for point mutations in the factor VIII gene. *Hum Mutat.* 1998;12(6):393-402. 9829908 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9829908) (http://www.ncbi.nlm.nih.gov/pubmed/9829908)
62. Williams IJ, Abuzenadah A, Winship PR, Preston FE, Dolan G, Wright J, Peake IR, Goodeve AC: Precise carrier diagnosis in families with haemophilia A: use of conformation sensitive gel electrophoresis for mutation screening and polymorphism analysis. *Thromb Haemost.* 1998 Apr;79(4):723-6. 9569180 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9569180) (http://www.ncbi.nlm.nih.gov/pubmed/9569180)
63. Hay CR, Ludlam CA, Colvin BT, Hill FG, Preston FE, Waseem N, Bagnall R, Peake IR, Berntorp E, Mauser Bunschoten EP, Fijnvandraat K, Kasper CK, White G, Santagostino E: Factor VIII inhibitors in mild and moderate-severity haemophilia A. UK Haemophilia Centre Directors Organisation. *Thromb Haemost.* 1998 Apr;79(4):762-6. 9569189 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/9569189) (http://www.ncbi.nlm.nih.gov/pubmed/9569189)
64. Keeling DM, Sukhu K, Kemball-Cook G, Waseem N, Bagnall R, Lloyd JV: Diagnostic importance of the two-stage factor VIII:C assay demonstrated by a case of mild haemophilia associated with His1954-->Leu substitution in the factor VIII A3 domain. *Br J Haematol.* 1999 Jun;105(4):1123-6. 10554831 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10554831) (http://www.ncbi.nlm.nih.gov/pubmed/10554831)
65. Strmecki L, Benedik-Dolnicar M, Vouk K, Komel R: Screen of 55 Slovenian haemophilia A patients: identification of 2 novel mutations (S-1R and IVS23+1G-->A) and discussion of mutation spectrum. *Mutation in brief no. 241. Online. Hum Mutat.* 1999;13(5):413. 10338101 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10338101) (http://www.ncbi.nlm.nih.gov/pubmed/10338101)
66. Moller-Morlang K, Tavassoli K, Eigel A, Pollmann H, Horst J: Mutational-screening in the factor VIII gene resulting in the identification of three novel mutations, one of which is a donor splice mutation. *Mutations in brief no. 245. Online. Hum Mutat.* 1999;13(6):504. 10408784 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10408784) (http://www.ncbi.nlm.nih.gov/pubmed/10408784)
67. Waseem NH, Bagnall R, Green PM, Giannelli F: Start of UK confidential haemophilia A database: analysis of 142 patients by solid phase fluorescent chemical cleavage of mismatch. *Haemophilia Centres. Thromb Haemost.* 1999 Jun;81(6):900-5. 10404764 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10404764) (http://www.ncbi.nlm.nih.gov/pubmed/10404764)
68. Jacquemin M, Lavend'homme R, Benhida A, Vanzielegem B, d'Oiron R, Lavergne JM, Brackmann HH, Schwaab R, VandenDriessche T, Chuah MK, Hoylaerts M, Gilles JG, Peerlinck K, Vermylen J, Saint-Remy JM: A novel cause of mild/moderate hemophilia A: mutations scattered in the factor VIII C1 domain reduce factor VIII binding to von Willebrand factor. *Blood.* 2000 Aug 1;96(3):958-65. 10910910 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10910910) (http://www.ncbi.nlm.nih.gov/pubmed/10910910)
69. Liu ML, Shen BW, Nakaya S, Pratt KP, Fujikawa K, Davie EW, Stoddard BL, Thompson AR: Hemophilic factor VIII C1- and C2-domain missense mutations and their modeling to the 1.5-angstrom human C2-domain crystal structure. *Blood.* 2000 Aug 1;96(3):979-87. 10910913 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10910913) (http://www.ncbi.nlm.nih.gov/pubmed/10910913)
70. Roelse JC, De Laaf RT, Timmermans SM, Peters M, Van Mourik JA, Voorberg J: Intracellular accumulation of factor VIII induced by missense mutations Arg593-->Cys and Asn618-->Ser explains cross-reacting material-reduced haemophilia A. *Br J Haematol.* 2000 Feb;108(2):241-6. 10691849 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10691849) (http://www.ncbi.nlm.nih.gov/pubmed/10691849)

71. Schwaab R, Oldenburg J, Kemball-Cook G, Albert T, Juhler C, Hanfland P, Ingerslev J: Assay discrepancy in mild haemophilia A due to a factor VIII missense mutation (Asn694Ile) in a large Danish family. *Br J Haematol.* 2000 Jun;109(3):523-8. 10886198 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10886198) (http://www.ncbi.nlm.nih.gov/pubmed/10886198)
72. Tagariello G, Belvini D, Salviato R, Are A, De Biasi E, Goodeve A, Davoli P: Experience of a single Italian center in genetic counseling for hemophilia: from linkage analysis to molecular diagnosis. *Haematologica.* 2000 May;85(5):525-9. 10800171 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10800171) (http://www.ncbi.nlm.nih.gov/pubmed/10800171)
73. Goodeve AC, Williams I, Bray GL, Peake IR: Relationship between factor VIII mutation type and inhibitor development in a cohort of previously untreated patients treated with recombinant factor VIII (Recombinate). *Recombinate PUP Study Group. Thromb Haemost.* 2000 Jun;83(6):844-8. 10896236 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10896236) (http://www.ncbi.nlm.nih.gov/pubmed/10896236)
74. Akkarapatumwong V, Oranwiroon S, Pung-amritt P, Treesucon A, Thanootarakul P, Veerakul G, Mahasandana C, Panyim S, Yenchitsomanus P: Mutations of the factor VIII gene in thai hemophilia A patients. *Hum Mutat.* 2000 Jan;15(1):117-8. 10612839 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/10612839) (http://www.ncbi.nlm.nih.gov/pubmed/10612839)
75. Leuer M, Oldenburg J, Lavergne JM, Ludwig M, Fregin A, Eigel A, Ljung R, Goodeve A, Peake I, Olek K: Somatic mosaicism in hemophilia A: a fairly common event. *Am J Hum Genet.* 2001 Jul;69(1):75-87. Epub 2001 Jun 14. 11410838 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11410838) (http://www.ncbi.nlm.nih.gov/pubmed/11410838)
76. Ivaskevicius V, Jurgutis R, Rost S, Muller A, Schmitt C, Wulff K, Herrmann FH, Muller CR, Schwaab R, Oldenburg J: Lithuanian haemophilia A and B registry comprising phenotypic and genotypic data. *Br J Haematol.* 2001 Mar;112(4):1062-70. 11298607 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11298607) (http://www.ncbi.nlm.nih.gov/pubmed/11298607)
77. Theophilus BD, Enayat MS, Williams MD, Hill FG: Site and type of mutations in the factor VIII gene in patients and carriers of haemophilia A. *Haemophilia.* 2001 Jul;7(4):381-91. 11442643 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11442643) (http://www.ncbi.nlm.nih.gov/pubmed/11442643)
78. Bauduer F, Ducout L, Bendriss P, Falaises B, Lavergne JM: Mild haemophilia A discovered in a previously multi-operated 73-year-old man: characterization of a new mutation. *Haemophilia.* 2001 Jul;7(4):419-21. 11442647 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11442647) (http://www.ncbi.nlm.nih.gov/pubmed/11442647)
79. Timur AA, Gurgey A, Aktuglu G, Kavakli K, Canatan D, Olek K, Caglayan SH: Molecular pathology of haemophilia A in Turkish patients: identification of 36 independent mutations. *Haemophilia.* 2001 Sep;7(5):475-81. 11554935 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11554935) (http://www.ncbi.nlm.nih.gov/pubmed/11554935)
80. Bogdanova N, Lemcke B, Markoff A, Pollmann H, Dworniczak B, Eigel A, Horst J: Seven novel and four recurrent point mutations in the factor VIII (F8C) gene. *Hum Mutat.* 2001 Dec;18(6):546. 11748850 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11748850) (http://www.ncbi.nlm.nih.gov/pubmed/11748850)
81. Vidal F, Farssac E, Altisent C, Puig L, Gallardo D: Rapid hemophilia A molecular diagnosis by a simple DNA sequencing procedure: identification of 14 novel mutations. *Thromb Haemost.* 2001 Apr;85(4):580-3. 11341489 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11341489) (http://www.ncbi.nlm.nih.gov/pubmed/11341489)
82. Valleix S, Vinciguerra C, Lavergne JM, Leuer M, Delpech M, Negrier C: Skewed X-chromosome inactivation in monozygotic twin sisters results in severe and mild hemophilia A. *Blood.* 2002 Oct 15;100(8):3034-6. 12351418 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12351418) (http://www.ncbi.nlm.nih.gov/pubmed/12351418)
83. Mazurier C, Parquet-Gernez A, Gaucher C, Lavergne JM, Goudemand J: Factor VIII deficiency not induced by FVIII gene mutation in a female first cousin of two brothers with haemophilia A. *Br J Haematol.* 2002 Nov;119(2):390-2. 12406074 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12406074) (http://www.ncbi.nlm.nih.gov/pubmed/12406074)
84. Sukarova-Stefanovska E, Zisovski N, Muratovska O, Kostova S, Efremov GD: Three novel point mutations causing haemophilia A. *Haemophilia.* 2002 Sep;8(5):715-8. 12199686 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12199686) (http://www.ncbi.nlm.nih.gov/pubmed/12199686)
85. Cutler JA, Mitchell MJ, Smith MP, Savidge GF: The identification and classification of 41 novel mutations in the factor VIII gene (F8C). *Hum Mutat.* 2002 Mar;19(3):274-8. 11857744 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11857744) (http://www.ncbi.nlm.nih.gov/pubmed/11857744)
86. Frusconi S, Passerini I, Girolami F, Masieri M, Linari S, Longo G, Morfini M, Torricelli F: Identification of seven novel mutations of F8C by DHPLC. *Hum Mutat.* 2002 Sep;20(3):231-2. 12203998 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12203998) (http://www.ncbi.nlm.nih.gov/pubmed/12203998)
87. Citron M, Godmilow L, Ganguly T, Ganguly A: High throughput mutation screening of the factor VIII gene (F8C) in hemophilia A: 37 novel mutations and genotype-phenotype correlation. *Hum Mutat.* 2002 Oct;20(4):267-74. 12325022 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12325022) (http://www.ncbi.nlm.nih.gov/pubmed/12325022)
88. Liu ML, Nakaya S, Thompson AR: Non-inversion factor VIII mutations in 80 hemophilia A families including 24 with alloimmune responses. *Thromb Haemost.* 2002 Feb;87(2):273-6. 11858487 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/11858487) (http://www.ncbi.nlm.nih.gov/pubmed/11858487)

89. Klopp N, Oldenburg J, Uen C, Schneppenheim R, Graw J: 11 hemophilia A patients without mutations in the factor VIII encoding gene. *Thromb Haemost.* 2002 Aug;88(2):357-60. 12195713 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12195713) (http://www.ncbi.nlm.nih.gov/pubmed/12195713)
90. Bicocchi MP, Pasino M, Lanza T, Bottini F, Boeri E, Mori PG, Molinari AC, Rosano C, Acquila M: Analysis of 18 novel mutations in the factor VIII gene. *Br J Haematol.* 2003 Sep;122(5):810-7. 12930394 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12930394) (http://www.ncbi.nlm.nih.gov/pubmed/12930394)
91. Habart D, Kalabova D, Novotny M, Vorlova Z: Thirty-four novel mutations detected in factor VIII gene by multiplex CSGE: modeling of 13 novel amino acid substitutions. *J Thromb Haemost.* 2003 Apr;1(4):773-81. 12871415 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12871415) (http://www.ncbi.nlm.nih.gov/pubmed/12871415)
92. Yenchitsomanus P, Akkarapatumwong V, Pung-Amritt P, Intorasoot S, Thanootarakul P, Oranwiroon S, Veerakul G, Mahasandana C: Genotype and phenotype of haemophilia A in Thai patients. *Haemophilia.* 2003 Mar;9(2):179-86. 12614369 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/12614369) (http://www.ncbi.nlm.nih.gov/pubmed/12614369)
93. Bicocchi MP, Pasino M, Lanza T, Bottini F, Molinari AC, Caprino D, Rosano C, Acquila M: Small FVIII gene rearrangements in 18 hemophilia A patients: five novel mutations. *Am J Hematol.* 2005 Feb;78(2):117-22. 15682412 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/15682412) (http://www.ncbi.nlm.nih.gov/pubmed/15682412)
94. Hill M, Deam S, Gordon B, Dolan G: Mutation analysis in 51 patients with haemophilia A: report of 10 novel mutations and correlations between genotype and clinical phenotype. *Haemophilia.* 2005 Mar;11(2):133-41. 15810915 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/15810915) (http://www.ncbi.nlm.nih.gov/pubmed/15810915)
95. Cai XH, Wang XF, Dai J, Fang Y, Ding QL, Xie F, Wang HL: Female hemophilia A heterozygous for a de novo frameshift and a novel missense mutation of factor VIII. *J Thromb Haemost.* 2006 Sep;4(9):1969-74. Epub 2006 Jun 27. 16805874 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/16805874) (http://www.ncbi.nlm.nih.gov/pubmed/16805874)
96. Vencesla A, Corral-Rodriguez MA, Baena M, Cornet M, Domenech M, Baiget M, Fuentes-Prior P, Tizzano EF: Identification of 31 novel mutations in the F8 gene in Spanish hemophilia A patients: structural analysis of 20 missense mutations suggests new intermolecular binding sites. *Blood.* 2008 Apr 1;111(7):3468-78. doi: 10.1182/blood-2007-08-108068. Epub 2008 Jan 9. 18184865 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/18184865) (http://www.ncbi.nlm.nih.gov/pubmed/18184865)
97. Albanes S, Ruiz-Saez A, Boadas A, de Bosch N, Porco A: Identification of factor VIII gene mutations in patients with severe haemophilia A in Venezuela: identification of seven novel mutations. *Haemophilia.* 2011 Sep;17(5):e913-8. doi: 10.1111/j.1365-2516.2011.02500.x. Epub 2011 Mar 4. 21371196 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/21371196) (http://www.ncbi.nlm.nih.gov/pubmed/21371196)
98. Nair PS, Shetty S, Ghosh K: Factor VIII Antigen, Activity, and Mutations in Hemophilia A. *Clin Appl Thromb Hemost.* 2016 May;22(4):381-5. doi: 10.1177/1076029614562951. Epub 2014 Dec 29. 25550078 [🔗](http://www.ncbi.nlm.nih.gov/pubmed/25550078) (http://www.ncbi.nlm.nih.gov/pubmed/25550078)

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