

Numbering is for the alignment, not for FVIII sequence

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          10          20          30          40          50          60          70          80          90         100
FVIII C1 2037-2192  ....|....|....|....|....|....|....|....|....|....|....|....|....|....|....|
FV C1 1904-2065   DRDCRMPMGLSTGIISDSQIKASEFLG-----YWEPRLARLNNGGSYNAWSVEKLAAEFAS-KPWIQVDMQKEVIITGIQTQGA--KHYLKSCYTTEF
Lact C1 70-229   ---CVEPLGMENGNIAANSQIAASSVRVTFGLGQ-HWVPELARLNRAAGMVNAWT----PSSNDD-NPWIQVNLRRMWVTGVVTQGA--SRLASHEYLKAF
Neuro C1 275-430 ---CMEALGMESGEIHSQITASSQYST-----NWSAERSRLNYPE--NGWT----PGEDSY-REWIQVDLGLLRVFTAVGTQGAISKETKKKYVVKTY
FVIII C2 2193-2351 ---CSMPLGMESKAISDAQITASSYFTNMFA--TWSPSKARLHLQGRSNAWR---PQVNNP-KEWLQVDFQKTMKVTGVTTQGV--KSLLTSMYVKEF
FV C2 2066-2224  ---CSTPLGMENKGIENKQITASSFKKSWWGD--YWEPRFRARLNAQGRVNAWQ---AKANNN-KOWLEIDLLKIKKITAIIITQGC--KSLSSEMYVKS
Lact C2 230-387  ---CANPLGLKNNIPDKQITASSSYKTWGLHLFSWNPYSYARLDKQGNFNAWV---AGSYGN-DQWLQVDLGSSEKVTGIITQGA--RNFGSVQFVASY
Neuro C2 431-583 ---CSGMLGMVSGLISDSQITSSNQGDR-----NWMPENIRLVTSRSGWALP---PAPHSYINEWLQIDLGEEKIVRGIITQGG--KHRENKVFMRKF
Disc SM 2-152    ---STQGLVQLLANAQCHLRTSTNYNGVHTQFNSALNYKNNGTNTIDGSEAWC----SSIVDT-NQYIVAGCEVPRTFMCVALQGR----GDADQWVTSY

          110         120         130         140         150         160         170
FVIII C1 2037-2192  ....|....|....|....|....|....|....|....|....|....|....|....|....|....|.
FV C1 1904-2065   YVAYSSNQINWQIFKGNSTRNVMYFNGNSDASTIKENQFDPPIVARYIRISPT-RAYNRPTLRLELQCEVNG
Lact C1 70-229   KVAYSLNGHEFDFIHDVNKK-HKEFVGNWNKNAVHVNLFETPVEAQYVRLYPT-SCHTACTLRFELLGCELNG
Neuro C1 275-430  KIDVSSNGEDWITIKEGNKP--VLFQGNTPDVVVAVFPKPLITRFVRIKPA-TWETGISMRFEVYGCKITDYP
FVIII C2 2193-2351 LISSSQDGHQWTLFFQNGKV--KVFQGNQDSFTPVVNSLDPPLLTRYLRHPQ-SWVHQIALRMEVLGCEAQDLY
FV C2 2066-2224  TIHYSEQGVEWKPYRLKSSMVDKIFEGNTNTKGHVKNFFNPPIISRFIRVIPK-TWNQSIARLELFGCDIY
Lact C2 230-387  KVAYSNDSANWTEYQDPRTGSSKIFPGNWDNHSHKKNLFETPILARYVRILPV-AWHNRIALRLELLGC
Neuro C2 431-583  KIGYSNNGSDWKMIMDDSKRKAKEFEGNNYDTPRELRTF-PALSTRFIRIYPERATHGGLGLRMELLGC
Disc SM 2-152    KIRYSLDNVSWFEYRNG-----AAVTGVTDNRNTVVNHFFDTPIRARSIAIHPL-TWNGHISLRCEFYTQ
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