

The Italian AICE-Genetics hemophilia A database: results and correlation with clinical phenotype

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Citation: Margaglione M, Castaman G, Morfini M, Rocino A, Santagostino E, Tagariello G, Tagliaferri AR, Zanon E, Bicocchi MP, Castaldo G, Peyvandi F, Santacroce R, Torricelli F, Grandone E, Mannuccio PM, and the AICE-Genetics Study Group. The Italian AICE-Genetics hemophilia A database: results and correlation with clinical phenotype. Haematologica 2008 May; 93(5): 722-728. doi: 10.3324/haematol.12427

Supplementary Table S1.

Location	Nucleotide	Protein	Occurrence	Effect	Reported/New	Clinical Phenotype
Exon 13						
Exon 2	c.200dupA	p.Thr49AspfsX15	2	insertion	reported	severe
Exon 4	c.450_51 insT	p.Gly132TrpfsX10	1	insertion	new	severe
Exon 5	c.648_49 ins10bp	p.Lys198fsX6	1	insertion	new	severe
Exon 8	c.1073_74 insT	p.Asn339fsX	1	insertion	reported	severe
Exon 8	c.1189_1190insC	p.His378ProfsX3	1	insertion	new	severe
Exon 10	c.1464_1465 insCCAGAC	p.Ser470_Arg471insProAsp	1	insertion	new	moderate
Exon 10	c.1473_1474 insCCAGAC	p.Tyr473_Asn474insProAsp	1	insertion	new	moderate
Exon 14	c.2729_30 insC	p.Asp892ArgfsX8	1	insertion	new	severe
Exon 14	c.2777dupC	p.Ser908LysfsX7	1	insertion	new	severe
Exon 14	c.2945dupA	p.Val964CysfsX22	3	insertion	reported	severe/moderate
Exon 14	c.3637dupA	p.Ile1194AsnfsX26	6	insertion	reported	severe
Exon 14	c.3870dupA	p.Gly1272ArgfsX27	3	insertion	reported	severe
Exon 14	c.3911dupA	p.Gln1286AlafsX15	1	insertion	new	severe
Exon 14	c.3984dupA	p.Arg1310SerfsX2	1	insertion	new	severe
Exon 14	c.4329dupA	p.Asp1425ArgfsX18	3	insertion	reported	severe
Exon 14	c.4379dupA	p.Asn1442fsX	17	insertion	reported	severe/moderate
Exon 14	c.4774_75 insA	p.Thr1573AsnfsX20	1	insertion	new	severe
Exon 14	c.4788dupA	p.Glu1578ArgfsX15	1	insertion	new	severe
Exon 14	c.4825dupA	p.Thr1590AsnfsX3	2	insertion	reported	severe
Exon 14	c.4895_96 insT	p.Asn1614LysfsX1	1	insertion	reported	severe
Exon 14	c.4921_22 insT	p.Glu1623ArgfsX8	2	insertion	reported	severe/moderate
Exon 16	c.5501dupA	p.Thr1815AsnfsX12	3	insertion	new	severe
Exon 18	c.5961dupA	p.Glu1969ArgfsX4	1	insertion	reported	severe
Exon 22	c.6312_13 insG	p.Thr2086AspfsX20	1	insertion	new	severe
Exon 25	c.6894_95 insAAC	p.Val2280_Lys2281insAsn	1	insertion	new	severe
Exon 26	c.6966_67 insGACT	p.Arg2304AspfsX30	1	insertion	new	severe
Exon 1	c.172_188 del15bp	p.Tyr10del	1	small deletion	new	moderate
Exon 2	c.199_200 delAA	p.Lys48AspfsX14	1	small deletion	reported	severe
Exon 2	c.205_206 delCT	p.Leu50ValfsX13	3	small deletion	reported	severe
Exon 2	c.209_212 delTTGT	p.Phe51fsX	3	small deletion	reported	severe

Exon 2	c.224 delA	p.Asp56ValfsX17	1	small deletion	reported	severe
Exon 3	c.288_293 del GGCTGA	p.Gln77_Ala78del	1	small deletion	new	severe
Exon 3	c.347_353 del GTCTCA	p.Ser97MetfsX42	1	small deletion	new	severe
Exon 4	c.427_428 delGA	p.Glu124ArgfsX2	1	small deletion	new	severe
Exon 4	c.486_87 delGA	p.Asn144TrpfsX6	1	small deletion	new	severe
Intron 4	c.602_13+c.602_615del		1	small deletion	new	severe
Exon 5	c.654 delT	p.Phe199LeufsX17	1	small deletion	new	severe
Exon 5	c.665_670 +5del		1	small deletion	reported	severe
Exon 5	c.668_670 delaAG	p.Glu204del	1	small deletion	new	severe
Exon 6	c.677_9delGTT	p.S207fsX8	1	small deletion	new	severe
Exon 6	c.785 delC	p.Pro243GlnfsX2	1	small deletion	new	severe
Moderate						
Exon 7	c.919 delA	p.Ile288SerfsX3	1	small deletion	new	severe
Exon 7	c.957_59 delaGAT	p.Leu300del	1	small deletion	reported	severe
Exon 7	c.958_60 delaTG	p.Met301del	1	small deletion	reported	severe
Exon 9	c.1332 delA	p.Lys425SerfsX37	1	small deletion	reported	
Moderate						
Exon 11	c.1582_88 delTTCAAAT	p.Phe509_Tyr511IlefsX5	1	small deletion	new	severe
Exon 11	c.1748 delA	p.Asn1564ThrfsX2	1	small deletion	new	severe
Exon 12	c.1845 delC	p.Asn597IlefsX43	1	small deletion	new	severe
Exon 12	c.1874_75 delaCA	p.Pro606ArgfsX16	1	small deletion	new	severe
Exon 13	c.2015_17 delTCT	p.Phe653del	1	small deletion	reported	severe
Exon 13	c.2058_2059 delaAC	p.Leu669SerfsX40	1	small deletion	new	severe
Exon 13	c.2203_28 delaAACAC	p.Lys659_His660del	1	small deletion	new	severe
Exon 14	c.2202 delT	p.Gly716ValfsX15	1	small deletion	new	severe
Exon 14	c.2216 delA	p.Glu720GlyfsX11	1	small deletion	new	severe
Exon 14	c.2521_30 delCCTGGAGCAA	p.Pro822fsX	1	small deletion	new	severe
Exon 14	c.2534_37 delaCAG	p.Asp826ValfsX7	1	small deletion	new	severe
Exon 14	c.2810 delA	p.Asp918ValfsX18	1	small deletion	new	severe
Exon 14	c.2842 delC	p.Thr930LeufsX7	1	small deletion	reported	severe
Exon 14	c.3060_3064 delaCAA	p.Lys1002LeufsX19	1	small deletion	reported	severe
Exon 14	c.3090 delaAGA	p.Lys1012LeufsX8	1	small deletion	reported	severe
Exon 14	c.3224 delC	p.Ala1056ValfsX3	1	small deletion	new	severe
Exon 14	c.3430 delA	p.Ser1125ValfsX4	1	small deletion	new	severe
Exon 14	c.3548_49 delaAA	p.Lys1164ArgfsX13	1	small deletion	reported	severe
Exon 14	c.3618_3621 delaCAC	p.Asn1188IlefsX9	1	small deletion	reported	severe
Exon 14	c.3620_21 delaC giusta	p.His1188GlnfsX32	1	small deletion	reported	severe
Exon 14	c.3637 delA	p.Ile1194PhefsX4	15	small deletion	reported	
Severe/Moderate						
Exon 14	c.3652 delG	p.Glu1199LysfsX5	3	small deletion	new	severe
Exon 14	c.3835 delA	p.Arg1260GlnfsX28	2	small deletion	new	severe
Exon 14	c.3904 delaCAC	p.Thr1283AlafsX14	1	small deletion	new	severe
Exon 14	c.4097 delT	p.Lys1347fsX	1	small deletion	new	severe
Exon 14	c.4121_24 delaTAGA	p.Ile1355ThrfsX48	1	small deletion	reported	severe
Exon 14	c.4296_4300 delTTCTC	p.His1415SerfsX5	1	small deletion	reported	severe
Exon 14	c.4379 delA	p.Asn1441IlefsX5	6	small deletion	reported	severe
Exon 14	c.4380 delT	p.Asn1441LysfsX5	1	small deletion	reported	mild
Exon 14	c.4388_91 delaCTT	p.Ser1444fsX	1	small deletion	new	severe
Exon 14	c.4477_80 delaAAG	p.Lys1474LeufsX74	1	small deletion	new	severe
Exon 14	c.4483 delG	p.Glu1476ArgfsX72	1	small deletion	new	severe
Exon 14	c.4687 delG	p.Val1154PhefsX4	1	small deletion	new	severe
Exon 14	c.5125 delA	p.Ser1690AlafsX23	1	small deletion	reported	severe
Exon 14	c.5213_16 delaGAAA	p.Arg1719ThrfsX20	1	small deletion	reported	severe
Exon 16	c.5473_5478 delaTTTGTC	p.Phe1806_Val1807del	1	small deletion	new	severe
Exon 16	c.5563 delG	p.Ala1836LeufsX16	1	small deletion	new	severe
Exon 17	c.5688_89 delaCT	p.Leu1878ValfsX5	1	small deletion	new	
Moderate						
Exon 18	c.5961 delA	p.Glu1969ArgfsX41	4	small deletion	reported	
Severe/Moderate						
Exon 18	c.5973_76 delaAATG	p.Ala1973HisfsX36	2	small deletion	reported	severe
Intron 22	c.6430_21+c.6430_6439 del		1	small deletion	new	severe
Exon 24	c.6650 delA	p.Asn2198IlefsX25	1	small deletion	new	severe
Exon 25	c.6875_76 delaTC	p.Phe2274SerfsX90	1	small deletion	reported	severe
Exon 25	c.6881_82 delaTT	p.Phe2275SerfsX90	1	small deletion	reported	severe
Exon 26	c.6916 delC	p.Gln2287LysfsX7	1	small deletion	reported	severe
Promoter Exon 6			1	large deletion	reported	severe
Exon 1			1	large deletion	reported	severe
Exons 2-6			1	large deletion	new	severe
Exons 2-25			2	large deletion	reported	severe
Exons 5-10			1	large deletion	new	severe
Exons 7-22			1	large deletion	reported	severe
Exons 14-22			1	large deletion	reported	severe
Exons 14-26			2	large deletion	new	severe
Exons 21-22			1	large deletion	reported	severe
Exon 26			1	large deletion	reported	severe

F8 gene						
Exon 1	c.3 G>C	p.Met19Ile	1	large deletion	reported	severe
Exon 1	c.38 T>G	p.Leu7Arg	1	missense	reported	severe
Exon 1	c.63 A>G	p.Arg3Gly	2	missense	new	severe
Exon 1	c.102 C>G	p.Asp15Glu	1	missense	new	severe
Exon 1	c.103 T>C	p.Tyr16His	1	missense	new	mild
Exon 1	c.104 A>G	p.Tyr16Cys	1	missense	reported	severe
Exon 1	c.143 G>A	p.Arg29Lys	2	missense	new	severe
Moderate						
Exon 2	c.201 A>G	p.Thr49Ala	1	missense	reported	severe
Exon 2	c.214 G>A	p.Glu53Lys	1	missense	reported	severe
Moderate						
Exon 2	c.214 G>C	p.Glu53Asp	1	missense	new	mild
Exon 2	c.223 G>T	p.Asp56Tyr	1	missense	reported	severe
Exon 2	c.265 G>A	p.Gly70Ser	1	missense/splicing	reported	severe
Exon 3	c.362G>T	p.Gly102Val	1	missense	reported	
Moderate						
Exon 3	c.275 G>A	p.Gly73Asp	1	missense	new	severe
Exon 3	c.276 C>T	p.Pro74Ser	1	missense	new	severe
Moderate						
Exon 3	c.293 A>T	p.Glu79Val	1	missense	reported	
Moderate						
Exon 3	c.308 T>A	p.Val84Glu	1	missense	new	severe
Exon 3	c.339 T>A	p.His94Gln	1	missense	new	severe
Exon 3	c.341 C>T	p.Pro95Leu	1	missense	new	
Moderate						
Exon 3	c.343 T>G	p.Val96Gly	1	missense	new	
Moderate						
Exon 3	c.350 T>C	p.Leu98Pro	1	missense	new	severe
Exon 3	c.353 A>G	p.His99Arg	1	missense	new	
Moderate						
Exon 3	c.362 G>C	p.Gly102Ala	1	missense	new	severe
Exon 3	c.392 A>G	p.Ile76Val	1	missense	new	mild
Exon 3	c.440 G>A	p.Ala92Thr	1	missense	reported	severe
Exon 4	c.403 G>T	p.Asp116Tyr	1	missense	reported	severe
Exon 4	c.446 C>T	p.Pro130Leu	1	missense	new	severe
Exon 4	c.518 T>G	p.Leu154Arg	1	missense	new	mild
Exon 4	c.523 A>G	p.Tyr156Cys	1	missense	new	
Moderate						
Exon 4	c.555 G>A	p.Asp167Lys	1	missense	new	
Moderate						
Exon 4	c.575 T>C	p.Ile173Thr	1	missense	new	
Moderate						
Exon 4	c.578 G>A	p.Gly174Glu	1	missense	new	severe
Exon 4	c.584 T>G	p.Leu176Arg	1	missense	new	severe
Exon 4	c.591 T>C	p.Cys179Gly	1	missense	reported	
Moderate						
Exon 5	c.642 T>G	p.Phe195Leu	1	missense	reported	mild
Exon 5	c.650 T>C	p.Lys198Pro	1	missense	new	severe
Exon 4	c.651 T>A	p.Val162Glu	1	missense	reported	mild
Exon 6	c.758 T>G	p.Val234Gly	1	missense	new	severe
Exon 6	c.761 A>G	p.Asn235Ser	1	missense	new	severe
Exon 6	c.761 A>T	p.Asn235Ile	2	missense	reported	severe
Exon 6	c.763 G>T	p.Gly236Cys	1	missense	new	severe
Exon 6	c.776 G>C	p.Arg240Thr	1	missense	new	mild
Exon 7	c.800 G>A	p.Cys248Tyr	1	missense	reported	
Moderate						
Exon 7	c.883T>C	p.Phe276Leu	1	missense	reported	
Moderate						
Exon 7	c.886 C>T	p.Leu277Phe	1	missense	new	severe
Exon 7	c.901 C>T	p.Arg282Cys	2	missense	reported	severe
Exon 7	c.902 G>A	p.Arg282His	2	missense	reported	
Severe/Moderate						
Exon 7	c.920 T>G	p.Ile288Ser	4	missense	reported	severe
Exon 7	c.938 T>G	p.Leu294Arg	1	missense	reported	severe
Exon 7	c.967 G>A	p.Gly304Arg	1	missense	new	severe
Exon 7	c.980 T>G	p.Leu308Arg	1	missense	new	severe
Exon 7	c.986 G>A	p.Cys310Tyr	1	missense	new	severe
Exon 8	c.1025 A>G	p.Tyr323Cys	1	missense	reported	
Moderate						
Exon 8	c.1030 A>C	p.Lys325Glu	1	missense	new	mild
Exon 8	c.1033 G>T	p.Val326Leu	2	missense	reported	
Moderate						
Exon 8	c.1171 C>T	p.Arg372Cys	3	missense	reported	
Moderate						

Exon 8	c.1171 C>G	p.Arg372Gly	1	missense	reported	
Moderate						
Exon 8	c.1172 G>A	p.Arg372His	1	missense	reported	
Moderate						
Exon 8	c.1173 T>C	p.Ser373Leu	3	missense	reported	
Moderate						
Exon 8	c.1202 G>C	p.Trp382Cys	1	missense	new	mild
Exon 9	c.1313 T>C	p.Ile419Tyr	1	missense	new	mild
Exon 9	c.1315 G>T	p.Gly420Cys	1	missense	new	severe
Exon 9	c.1316 G>C	p.Gly420Ala	1	missense	new	
Moderate						
Exon 9	c.1322 T>C	p.Tyr423Asn	1	missense	new	severe
Exon 9	c.1332 A>C	p.Lys425Asn	3	missense	reported	severe
Exon 9	c.1364 T>C	p.Phe436Ser	1	missense	new	severe
Exon 9	c.1372 C>T	p.Arg439Cys	3	missense	reported	mild
Exon 9	c.1396 G>A	p.Gly447Arg	2	missense	new	
Severe/Moderate						
Exon 9	c.1414 T>A	p.Leu453His	1	missense	new	
Moderate						
Exon 9	c.1424 G>A	p.Gly420Ser	1	missense	reported	
Moderate						
Exon 9	c.1430 G>T	p.Gly458Val	7	missense	reported	
Moderate						
Exon 10	c.1475 A>G	p.Tyr473Cys	2	missense	reported	
Exon 10	c.1492 G>A	p.Gly479Arg	3	missense	reported	mild
Moderate/Mild						
Exon 10	c.1537 G>A	p.Gly494Ser	1	missense	reported	severe
Exon 11	c.1630 G>A	p.Asp525Asn	1	missense	reported	mild
Exon 11	c.1636 C>T	p.Arg527Trp	4	missense	reported	mild
Exon 11	c.1640 G>A	p.Cys528Tyr	1	missense	reported	
Moderate						
Exon 11	c.1648 C>T	p.Arg531Cys	3	missense	reported	mild
Exon 11	c.1649 G>A	p.Arg531His	1	missense	reported	mild
Exon 11	c.1682 A>C	p.Asp542Ala	1	missense	reported	severe
Exon 11	c.1694 A>G	p.Lys510Glu	1	missense	reported	
Moderate						
Exon 11	c.1700 T>C	p.Ile548Thr	6	missense	reported	mild
Exon 12	c.1754 T>C	p.Ile566Thr	1	missense	reported	severe
Exon 12	c.1810 T>C	p.Trp585Arg	1	missense	new	severe
Exon 12	c.1813 T>C	p.Tyr586His	1	missense	reported	severe
Exon 12	c.1814 A>G	p.Tyr586Cys	1	missense	new	severe
Exon 12	c.1834 C>T	p.Arg593Cys	2	missense	reported	
Moderate/Mild						
Exon 12	c.1897 A>G	p.Met614Val	1	missense	new	mild
Exon 12	c.1909 G>T	p.Glu581Asp	1	missense	new	severe
Exon 13	c.1913 G>A	p.Gly619Asp	1	missense	new	severe
Exon 13	c.1945 T>C	p.Cys630Arg	1	missense	new	severe
Exon 13	c.1966 T>C	p.Trp637Arg	1	missense	new	severe
Exon 13	c.1966 T>G	p.Trp637Gly	1	missense	new	
Moderate						
Exon 13	c.1991 A>G	p.Gln645Arg	1	missense	new	
Moderate						
Exon 13	c.2029 T>G	p.Phe658Val	1	missense	new	
Moderate						
Exon 13	c.2043 G>T	p.Met662Ile	1	missense	reported	
Moderate						
Exon 13	c.2061 C>T	p.Pro672Ser	1	missense	new	
Exon 13	c.2081 G>A	p.Gly675Glu	2	missense	new	severe
Severe/Moderate						
Exon 13	c.2090 T>C	p.Val678Ala	1	missense	reported	mild
Exon 13	c.2102 T>C	p.Met682Thr	1	missense	new	mild
Exon 14	c.2149 C>T	p.Arg698Trp	1	missense	reported	mild
Exon 14	c.2158 G>A	p.Gly701Ser	1	missense	reported	severe
Exon 14	c.2161 A>G	p.Met702Val	1	missense	new	severe
Exon 14	c.2163 G>C	p.Met702Ile	1	missense	reported	severe
Exon 14	c.2167 G>A	p.Ala704Thr	8	missense	reported	
Severe/Moderate/Mild						
Exon 14	c.2225 A>G	p.Tyr723Cys	1	missense	reported	mild
Exon 14	c.4114 A>G	p.Thr1353Ala	1	missense	new	severe
Exon 14	c.5063 C>T	p.Ser1669Leu	1	missense	new	
Moderate						
Exon 14	c.5122 C>T	p.Arg1689Cys	4	missense	reported	
Severe/Moderate						
Exon 14	c.5140 A>T	p.Thr1695Ser	2	missense	reported	mild
Exon 14	c.5176 T>C	p.Trp1707Arg	1	missense	new	severe

Exon 15	c.5253A>T	p.Lys1732Asn	1	missense	new	severe
Exon 15	c.5275 G>A	p.Asp1740Asn	1	missense	new	mild
Exon 15	c.5302 C>T	p.Arg1749Cys	2	missense	reported	mild
Exon 15	c.5320 C>T	p.His1755Tyr	1	missense	reported	mild
Exon 15	c.5321 A>G	p.His1755Arg	1	missense	reported	severe
Exon 15	c.5323 T>G	p.Leu1756Val	1	missense	reported	
Moderate						
Exon 15	c.5330 T>C	p.Leu1758Pro	3	missense	reported	severe
Exon 15	c.5335 G>A	p.Gly1760Arg	1	missense	reported	severe
Exon 16	c.5387 A>G	p.Asn1777 Ser	1	missense	new	severe
Exon 16	c.5398 C>A	p.Arg1781Ser	1	missense	new	severe
Exon 16	c.5398 C>T	p.Arg1781Cys	2	missense	reported	
Moderate						
Exon 16	c.5399 G>A	p.Arg1781His	10	missense	reported	
Severe/Moderate/Mild						
Exon 16	c.5405 A>C	p.Tyr1783Arg	1	missense	new	severe
Exon 16	c.5405 A>G	p.Tyr1783Cys	1	missense	reported	
Moderate						
Exon 16	c.5411 T>C	p.Phe1785Ser	2	missense	reported	severe
Exon 16	c.5413 T>A	p.Tyr1786Asn	1	missense	new	severe
Exon 16	c.5419 A>G	p.Ser1788Gly	1	missense	new	mild
Exon 16	c.5422 C>T	p.Leu1789Phe	1	missense	reported	mild
Exon 16	c.5522 A>T	p.His1822Leu	1	missense	new	mild
Exon 16	c.5527 G>A	p.Ala1824Thr	1	missense	reported	mild
Exon 16	c.5533 A>C	p.Thr1826Pro	1	missense	reported	severe
Exon 16	c.5558 C>T	p.Ala1834Val	2	missense	reported	mild
Exon 16	c.5561 T>G	p.Trp1835Gly	1	missense	new	severe
Exon 17	c.5674 G>A	p.Val1873Ile	1	missense	new	
Moderate						
Exon 17	c.5715 G>A	p.Gly1850Asp	1	missense	reported	severe
Exon 17	c.5726 A>G	p.Tyr1890Cys	3	missense	reported	
Severe/Moderate						
Exon 17	c.5813 A>G	p.His1919Arg	2	missense	reported	mild
Exon 18	c.5824 G>A	p.Gly1923Asp	1	missense	new	severe
Exon 18	c.5837 A>T	p.Asp1927Val	1	missense	new	mild
Exon 18	c.5879 G>A	p.Arg1941Gln	2	missense	reported	
Severe/Mild						
Exon 18	c.5918 A>G	p.His1954 Arg	1	missense	new	severe
Exon 18	c.5954 G>A	p.Arg1966Gln	2	missense	reported	mild
Exon 18	c.5981 T>C	p.Leu1975Pro	1	missense	reported	
Moderate						
Exon 18	c.5998 G>A	p.Gly1981Ser	2	missense	new	severe
Exon 19	c.6043 T>C	p.Trp1996Arg	3	missense	new	severe
Exon 19	c.6045 G>T	p.Trp1996Cys	1	missense	reported	
Exon 19	c.6046 T>C	p.Arg1997Trp	19	missense	reported	
Severe/Moderate/Mild						
Exon 19	c.6092 C>T	p.Thr2012Ile	1	missense	reported	mild
Exon 19	c.6104 T>G	p.Val2016Gly	1	missense	new	severe
Exon 20	c.6151 A>T	p.Ile2032Phe	1	missense	new	mild
Exon 20	c.6151 T>G	p.Ile2032Ser	1	missense	reported	severe
Exon 20	c.6172 G>C	p.Ala2039Pro	7	missense	reported	
Severe/Moderate						
Exon 21	c.6267 G>T	p.Trp2070Cys	2	missense	new	mild
Exon 21	c.6275 C>T	p.Ser2069Phe	1	missense	reported	severe
Exon 22	c.6320 G>A	p.Gly2088Asp	1	missense	new	severe
Exon 22	c.6371 A>G	p.Tyr2105Cys	6	missense	reported	mild
Exon 22	c.6374 G>C	p.Ser2106Thr	1	missense	new	severe
Exon 22	c.6531 C>A	p.Thr2122Asn	1	missense	reported	severe
Exon 23	c.6506 G>T	p.Arg2150Leu	1	missense	new	mild
Exon 23	c.6506 G>A	p.Arg2150His	13	missense	reported	
Severe/Moderate/Mild						
Exon 23	c.6532 C>T	p.Arg2159Qs	3	missense	reported	mild
Exon 23	c.6533 G>A	p.Arg2159His	1	missense	reported	
Moderate						
Exon 23	c.6545 G>A	p.Arg2163His	14	missense	reported	
Severe/Moderate						
Exon 23	c.6546 T>C	p.Phe2127Ser	1	missense	new	mild
Exon 23	c.6560 G>A	p.Gly2168Asp	2	missense	new	severe
Exon 23	c.6563 G>A	p.Cys2169Tyr	1	missense	new	
Moderate						
Exon 24	c.6593 G>A	p.Gly2179Glu	1	missense	new	severe
Exon 24	c.6593 G>T	p.Gly2179Val	2	missense	reported	severe
Exon 24	c.6637 T>C	p.Ser2194Pro	1	missense	new	severe
Exon 24	c.6638 C>G	p.Ser2194Cys	3	missense	new	
Moderate						

Exon 24	c.6652 A>G	p.Met2199Val	1	missense	new	severe
Exon 24	c.6683 G>A	p.Arg2209Gln	5	missense	reported	severe
Exon 24	c.6685 C>T	p.Leu2210Phe	2	missense	reported	severe
Exon 25	c.6738 G>A	p.Glu2228Lys	1	missense	new	mild
Moderate						
Exon 25	c.6742 T>C	p.Trp2229Arg	1	missense	new	mild
Exon 25	c.6744 G>T	p.Trp2229Cys	1	missense	reported	
Moderate						
Exon 25	c.6746 T>C	p.Leu2230Pro	1	missense	new	severe
Exon 25	c.6751 T>C	p.Val2232Ala	3	missense	reported	mild
Exon 25	c.6769 A>G	p.Met2238Arg	1	missense	reported	mild
Exon 25	c.6776 T>C	p.Val2240Asp	1	missense	new	severe
Exon 25	c.6785 T>C	p.Val2243Ala	1	missense	new	
Moderate						
Exon 26	c.6956 C>G	p.Pro2300Arg	1	missense	new	severe
Exon 26	c.6956 C>T	p.Pro2300Leu	4	missense	reported	mild
Exon 26	c.6967 C>T	p.Arg2304Cys	7	missense	reported	
Severe/Moderate						
Exon 26	c.6977 G>A	p.Arg2307Gln	5	missense	reported	
Severe/Moderate/Mild						
Exon 26	c.6982 C>G	p.His2309Asp	1	missense	new	severe
Exon 26	c.6989 A>C	p.Gln2311Pro	1	missense	reported	severe
Exon 26	c.6994 T>C	p.Trp2313Arg	2	missense	reported	severe
Exon 26	c.6995 G>C	p.Trp2313Ser	1	missense	new	
Moderate						
Exon 26	c.7015 A>T	p.Arg2320Trp	1	missense	new	severe
Exon 26	c.7031 G>A	p.Gly2325Asp	1	missense	reported	severe
Exon 26	c.*1 T>C		1	missense	new	mild
Exon 1	c.43 C>T	p.Arg-5Stop	2	nonsense	reported	severe
Exon 1	c.207 G>A	p.Trp14Stop	1	nonsense	reported	severe
Exon 3	c.368 C>G	p.Ser104Stop	1	nonsense	new	severe
Exon 4	c.406 C>T	p.Gln117Stop	1	nonsense	reported	severe
Exon 4	c.436 A>T	p.Lys127Stop	1	nonsense	new	severe
Exon 6	c.686 C>G	p.Ser210Stop	2	nonsense	new	severe
Exon 6	c.709 C>T	p.Gln218Stop	1	nonsense	new	severe
Exon 6	c.741 G>A	p.Trp228Stop	1	nonsense	reported	severe
Exon 7	c.821 G>A	p.Trp255Stop	1	nonsense	reported	severe
Exon 8	c.1063 C>T	p.Arg336Stop	2	nonsense	reported	severe
Exon 9	c.1336 C>T	p.Arg427Stop	5	nonsense	reported	severe
Exon 10	c.1594 C>G	p.Tyr476Stop	1	nonsense	reported	
Moderate						
Exon 10	c.1629C>G	p.Ser488Stop	1	nonsense	reported	severe
Exon 11	c.1738 C>T	p.Gln561Stop	1	nonsense	new	severe
Exon 12	c.1804 C>T	p.Arg583Stop	5	nonsense	reported	
Severe/Moderate						
Exon 12	c.1811 G>A	p.Trp585Stop	1	nonsense	new	severe
Exon 12	c.1831 C>T	p.Gln592Stop	2	nonsense	reported	severe
Exon 13	c.1933 C>T	p.Gln626Stop	1	nonsense	reported	severe
Exon 13	c.2078 C>G	p.Ser674Stop	1	nonsense	new	severe
Exon 14	c.2372 G>A	p.Trp772Stop	2	nonsense	reported	severe
Exon 14	c.2440 C>T	p.Arg795Stop	3	nonsense	reported	severe
Exon 14	c.2742 T>G	p.Leu859Stop	1	nonsense	new	severe
Exon 14	c.3113 C>G	p.Ser1019Stop	1	nonsense	new	severe
Exon 14	c.3175 A>T	p.Lys1040Stop	1	nonsense	reported	severe
Exon 14	c.3381 G>A	p.Trp1108Stop	3	nonsense	reported	severe
Exon 14	c.3865 A>T	p.Lys1270Stop	1	nonsense	new	severe
Exon 14	c.3967 C>T	p.Gln1304Stop	1	nonsense	reported	
Moderate						
Exon 14	c.4067 C>A	p.Ser1337Stop	1	nonsense	new	severe
Exon 14	c.4077 G>A	p.Trp1340Stop	1	nonsense	new	severe
Exon 14	c.4156 C>T	p.Gln1367Stop	1	nonsense	new	severe
Exon 14	c.4363 C>T	p.Gln1436Stop	1	nonsense	new	severe
Exon 14	c.4483 G>T	p.Glu1476Stop	1	nonsense	new	severe
Exon 14	c.5143 C>T	p.Arg1696Stop	7	nonsense	reported	severe
Exon 14	c.5177 G>A	p.Trp1707Stop	2	nonsense	reported	
Severe/Moderate						
Exon 14	c.5184 T>A	p.Tyr1709Stop	1	nonsense	reported	severe
Exon 16	c.5508 G>A	p.Trp1817Stop	1	nonsense	reported	severe
Exon 16	c.5515 C>T	p.Gln1820Stop	1	nonsense	new	severe
Exon 17	c.5662 A>T	p.Arg1869Stop	1	nonsense	new	severe
Exon 18	c.5878 C>T	p.Arg1941Stop	1	nonsense	reported	severe
Exon 18	c.5953 C>T	p.Arg1966Stop	5	nonsense	reported	
Severe/Moderate						
Exon 18	c.5985 C>A	p.Tyr1976Stop	1	nonsense	reported	severe
Exon 18	c.5992 G>A	p.Trp1942Stop	1	nonsense	reported	severe
Exon 21	c.6335 G>T	p.Gly2057Stop	1	nonsense	reported	severe

Exon 22	c.6310 A>T	p.Lys2085Stop	1	nonsense	new	severe
Exon 22	c.6403 C>T	p.Arg2116Stop	4	nonsense	reported	severe
Exon 23	c.6605 C>T	p.Arg2147Stop	1	nonsense	reported	severe
Exon 24	c.6682 C>T	p.Arg2209Stop	5	nonsense	reported	severe
Exon 25	c.6739 G>T	p.Glu2228Stop	2	nonsense	new	
Severe/Moderate						
Exon 26	c.6907 C>T	p.Gln2284Stop	1	nonsense	new	severe
Exon 1	c.-43 C>T		1	transcription	new	mild
Exon 1	c.-25 A>G		1	transcription	new	severe
Intron 2	c.144 -2 A>T		1	splicing	new	severe
Intron 2	c.144 -3 C>G		1	splicing	new	severe
Intron 3	c.388 +5 G>A		1	splicing	reported	severe
Intron 4	c.601 +5 G>T		1	splicing	new	mild
Intron 4	c.601 +5 G>A		1	splicing	reported	mild
Intron 4	c.602 -1 G>A		1	splicing	new	severe
Intron 5	c.670 +1 G>T		1	splicing	reported	severe
Intron 5	c.670 +6 A>G		1	splicing	new	mild
Intron 6	c.787 +1 G>A		1	splicing	reported	mild
Intron 6	c.787 +5 G>A		1	splicing	new	severe
Intron 6	c.788 -1 G>T		1	splicing	reported	severe
Intron 9	c.1443 +1 G>C		1	splicing	new	severe
Moderate						
Intron 9	c.1443 +1 G>T		1	splicing	new	Severe
Intron 10	c.1537 +1G>C		1	splicing	new	Severe
Intron 10	c.1538 -1 A>G		1	splicing	new	Severe
Intron 10	c.1538 -18 G>A		17	splicing	new	Moderate
Exon 11	c.1678 G>T		7	splicing	reported	Severe/Moderate
Intron 12	c.1903 +2 T>G		2	splicing	new	Severe
Intron 12	c.1903 +5 G>A		2	splicing	reported	Severe/Moderate
Intron 12	c.1904 -1G>A		1	splicing	new	Severe
Intron 12	c.1904 -2A>G		1	splicing	new	Severe
Intron 14	c.5219 +1 G>A		4	splicing	new	Severe
Intron 14	c.5219 +2 T>C		1	splicing	new	Severe
Intron 14	c.5219 +3 A>G		2	splicing	reported	Moderate
Intron 15	c.5220 -2 A>T		2	splicing	new	Severe
Intron 16	c.5373 +1 G>A		1	splicing	new	Severe
Intron 16	c.5373 +2 T>G		1	splicing	new	Moderate
Intron 19	c.6115 +4 A>G		1	splicing	new	Severe
Intron 19	c.6115 delGTG +1_+3		1	splicing	new	Severe
Intron 21	c.6187 -1 G>A		1	splicing	new	Severe
Intron 21	c.6188 +1 G>T		2	splicing	reported	Sever
Intron 25	c.6723 +1 G>T		1	splicing	reported	Severe
Exon 15	del 3bp inser22bp	p.Val1727delins22bp	1	insertion/small deletion	new	severe
Exon 24	c.6623_25 delAGA Gln2189p.Gln2189del + p.Thr2191del		1	small deletions	new	severe
	+c.6628 delA Thr2191					

Supplementary Table S2. F8 mutations occurring at a CpG site

Exon	Mutation	Type	Phenotype (number)
1c.43C>T	p.Arg>5Stop	nonsense	severe(2)
7c.901C>T	p.Arg282Cys	missense	severe(1)
7c.902G>A	p.Arg282His	missense	severe(1)/moderate(2)
8c.1063C>T	p.Arg336Stop	nonsense	severe(2)
8c.1171C>T	p.Arg372Cys	missense	moderate(3)
8c.1172G>A	p.Arg372His	missense	moderate(1)
9c.1336C>T	p.Arg427Stop	nonsense	severe(5)
10c.1492 G>A	p.Gly479Arg	missense	moderate(2)/mild(1)
11c.1636C>T	p.Arg527Trp	missense	mild(3)
11c.1648C>T	p.Arg531Cys	missense	mild(3)
11c.1649G>A	p.Arg531His	missense	mild(1)
12c.1804C>T	p.Arg583Stop	nonsense	severe(3)/moderate(2)
12c.1834C>T	p.Arg593Cys	missense	moderate(1)/mild(1)
14c.2149C>T	p.Arg698Trp	missense	mild(1)
14c.2167G>A	p.Ala704Thr	missense	severe(1)/moderate(2)/mild(7)
14c.2440C>T	p.Arg795Stop	nonsense	severe(3)
14c.5122C>T	p.Arg1689Cys	missense	severe(1)/moderate(1)/mild(1)
14c.5123G>A	p.Arg1689His	missense	mild(1)
14c.5252C>T	p.Arg1696Stop	nonsense	severe(7)
15 c.5302C>T	p.Arg1749Cys	missense	mild(1)
16c.5398C>T	p.Arg1781Cys	missense	moderate(1)
16c.5399G>A	p.Arg1781His	missense	severe(2)/moderate(6)/mild(3)
18c.5878C>T	p.Arg1941Stop	nonsense	severe(1)
18c.5879G>A	p.Arg1941Gln	missense	severe(1)/mild(1)
18c.5953C>T	p.Arg1966Stop	nonsense	severe(4)/moderate(1)
18c.5954G>A	p.Arg1966Gln	missense	mild(1)
19c.6046C>T	p.Arg1997Trp	missense	severe(13)/moderate(5)/mild(1)
22c.6403C>T	p.Arg2116Stop	nonsense	severe(4)
23c.6496C>T	p.Arg2147Stop	nonsense	severe(1)
23c.6506G>A	p.Arg2150His	missense	severe(1)/moderate(7)/mild(5)
23c.6532C>T	p.Arg2159Cys	missense	mild(2)
23c.6533G>A	p.Arg2159His	missense	moderate(1)
23c.6544C>T	p.Arg2163Cys	missense	severe(2)
23c.6545G>A	p.Arg2163His	missense	severe(6)/moderate(5)
24c.6682C>T	p.Arg2209Stop	nonsense	severe(5)
24c.6683G>A	p.Arg2209Gln	missense	severe(5)
26c.6956C>T	p.Pro2300Leu	missense	mild(4)
26c.6967C>T	p.Arg2304Cys	missense	severe(5)/moderate(2)
26c.6977G>A	p.Arg2307Gln	missense	severe(1)/moderate(1)/mild(3)

Supplementary Table S3. F8 mutations and clinical phenotype in patients with moderate and mild HA and inhibitors.

Location	Nucleotide	Protein	Effect	Clinical Phenotype
Exon 2	c.214 G>A	p.Glu53Lys	missense	moderate
Exon 3	c.353 A>G	p.His99Arg	missense	moderate
Exon 9	c.1372 C>T	p.Arg439Cys	missense	mild
Exon 9	c.1430 G>T	p.Gly458Val	missense	mild
Intron 9	c.1443 +1 G>C		splicing	moderate
Exon 10	c.1594 C>G	p.Tyr476Stop	nonsense	moderate
Intron 10	c.1538 -18 G>A		splicing	mild
Intron 10	c.1538 -18 G>A		splicing	moderate
Exon 14	c.3967 C>T	p.Gln1304Stop	nonsense	moderate
Exon 16	c.5398 C>T	p.Arg1781Cys	missense	moderate
Exon 16	c.5399 G>A	p.Arg1781His	missense	moderate
Exon 16	IVS16 +2 t>g			moderate
Exon 17	c.5726 A>G	p.Tyr1890Cys	missense	moderate
Exon 22	c.6371 A>G	p.Tyr2105Cys	missense	mild
Exon 23	c.6506 G>A	p.Arg2150His	missense	mild
Exon 23	6506G>A	p.Arg2150His	missense	moderate
Exon 23	6506G>A	p.Arg2150His	missense	moderate