

Table S1: Multiple sequence alignment for amino acid sequences of HEV pORF1. Alignment is performed by using ClustalW program in Bioedit software. It represents the conserved cleavage site for thrombin and factor Xa on pORF1.

Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd	Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd
BAA011172.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAD95603.1	DADGRLTCK	TLTEHPIIH	SVIRHSTVI
AAA654882.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAE02700.1	DAEGRRLTCK	TLTEHPIIH	SVVHSTIS
CAA669336.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDT	BAE02703.1	DAEGRRLTCK	TLTEHPIIH	SVVHSTIS
BAA018665.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAE48647.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
CAA678009.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAE48650.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
AAC971826.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE48653.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
AAD15812.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE75896.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
AAD15815.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDC	BAE79643.1	DADGRLTCK	TLTEHPIIH	SVIRHSTVL
AAB820029.2	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAE79646.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
AAC279334.2	DADGRLTCK	TLTEHPIIH	SVIRGTIDN	BAE79649.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
CAB832009.1	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	BAE79652.1	DADGRLTCK	TLTEHPIIH	SVIRHSTIL
AAG16764.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE79655.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
CAC501800.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE79658.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
CAC501820.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDR	BAE79661.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
BAB633939.2	DADGRLTCK	TLTEHPIIH	SVIRGTVDS	BAE79664.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
BAB933533.2	DTDGRRLTCK	TLTEHPIIH	SVIRGTINN	BAE79667.1	DTDGRRLTCK	TLTEHPIIH	SVIRHSTIN
BAB933533.3	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE86998.1	DADGRLTCK	TLTEHPIIH	SVIRHSTIL
BAB933533.4	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAE92012.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAB933533.5	CVDGRLDCE	TLTEHPIIH	SVIRGNPDA	ABE27147.1	DADGRLTCK	TLTEHPIIH	SVIRHSTIL
BAB933533.6	DADGRLTCK	TLTEHPIIH	SVIRGNPDA	ABE73522.1	CADGRLDCE	TLTEHPIIH	SVIRHGNPL
AAM663329.1	ASPIRLECR	TLTEHPIIH	SVIRGTVDS	BAE98084.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
BAC056699.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDC	BAE98087.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
BAC448399.7	DADGRLTCK	TLTEHPIIH	SVIRGNPDC	BAE98090.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
BAC652550.1	ASPIRLECR	TLTEHPIIH	SVIRGTVDS	ABK80468.1	DAEGRRLTCK	TLTEHPIIH	SVVHSTIS
BAC652553.1	CAEGRDCE	TLTEHPIIH	SVIRGNPDR	BAF65205.1	ASPIHLECF	TLTEHPIIH	SVIRHSSPL
AAO673358.1	DADGRLTCK	TLTEHPIIH	SVIRGNPDA	BAF65208.1	ASPGHLECF	TLTEHPIIH	STIRHSTPL
AAO729391.1	DVDGRLTCK	TLTEHPIIH	SVIRGTVDS	ABU98319.1	DADGRLTCK	TLTEHPIIH	SVIRHSTIL
BAC756330.1	DADGRLTCK	TLTEHPIIH	SVIRGTVDS	BAF92625.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAC77167.1	ASPIRLECR	TLTEHPIIH	SVIRGTVDS	ABY66135.1	DAEGRRLTCK	TLTEHPIIH	SVVHSTIS
BAC79251.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	BAG09237.1	CVDGRLDCE	TLTEHPIIH	SVIRHGNPL
AAT409994.1	ASPIRLECR	TLTEHPIIH	SVIRGNPDC	ACA83737.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
AAT409997.1	DTDGRRLTCK	TLTEHPIIH	SVIRGNPDC	ACB15468.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
AAT410000.1	DTDGRRLTCK	TLTEHPIIH	SVIRGTINN	BAG32124.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
BAD34958.1	DTDGRRLTCK	TLTEHPIIH	SVIRGTINN	BAG32127.1	DADGRLTCK	TLTEHPIIH	SVVHSTVI
BAD34961.1	AGPIRLECR	TLTEHPIIH	SVIRGTINN	BAG32130.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD34964.1	AGPIRLECR	TLTEHPIIH	SVIRGNPDC	BAG32133.1	DADGRLTCK	TLTEHPIIH	SVIRHSTIL
BAD74166.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDC	BAG32136.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD74169.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDC	ACD62590.1	ASPGHLECF	TLTEHPIIH	STIRHGNPL
BAD74172.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDC	BAG67067.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD74175.1	AGPIRLECF	TLTEHPIIH	SVIRGNPDC	BAG67064.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD74178.1	DADGRLTCK	TLTEHPIIH	SVIRGNPDC	BAG67061.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD74181.1	DADGRLTCK	TLTEHPIIH	SVIRGTIDS	BAG67058.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
BAD74184.1	DADGRLTCK	TLTEHPIIH	SVIRGTIDS	BAG67055.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
AAW58001.1	DADGRLTCK	TLTEHPIIH	SVIRGTIDS	BAG67073.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL
			SAIRHSTVDS	BAG67070.1	AGPIHLECF	TLTEHPIIH	SVIRHGNPL

Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd
ACJ70098.1	CVDGRIDCF	TLTHRRPIIH	SVIERRGNPD
BAH03559.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH03562.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH03565.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH03568.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH03571.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08575.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08578.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08581.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08584.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08587.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08590.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08593.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08596.1	ASPGRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08599.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH08602.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08605.1	ASPGRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08608.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08611.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH08614.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH08617.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH08620.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH08623.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH08626.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ACL54976.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH10868.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
ACN76575.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAH59608.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH59611.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
ACR56297.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAH69078.1	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ACT52380.1	VGPGRLECF	TLTHRRPIIH	SVIERRGNPD
ACT52383.1	TGPGRLECF	TLTHRRPIIH	SVIERRGNPD
ACU12604.1	ASPGRLECF	TLTHRRPIIH	SVIERRGNPD
ADA61126.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ADG36436.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAJ09463.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
BAJ09466.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ADK88897.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ADO17767.1	DADGRFLTCK	TLTHRRPIIH	SVIERRGNPD
ADV71352.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAJ61825.1	DTAGRFLACF	TLTHRRPIIH	SVIERRGNPD
ADV92628.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
ADV92631.1	AGPIRLECF	TLTHRRPIIH	SVIERRGNPD
BAJ22676.2	DTDGRFLTCK	TLTHRRPIIH	SVIERRGNPD

Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd
ADZ57215.1	DAEGRILTCK	LTHRRPIIHA	PSVIERRGNPD
BAK08878.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAK08881.1	ASPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AEA48881.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
BAK19084.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
BAK19087.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
BAK52233.1	ASPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAK52236.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AEO16235.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AEV52326.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AEW87504.1	ASPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AEX32827.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAL61209.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFD33683.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFD33686.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFH34996.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AFH34999.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AFH35002.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AFJ93211.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFJ93214.1	CVDGRIDCF	LTHRRPIIHA	PSVIERRGNPD
AFJ93217.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFJ93220.1	DTDGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFK65670.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFK65673.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFO10118.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFP73624.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFP82888.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAM36431.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AFR53968.1	ASPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AFR53971.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AFR53974.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AFV92497.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAM68889.1	VGPGRLECF	LTHRRPIIHA	PSVIERRGNPD
BAM68892.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAM68895.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
BAM78544.1	AGPIRLECF	LTHRRPIIHA	PSVIERRGNPD
AGE91705.1	DADGRILTCK	LTHRRPIIHA	PSVIERRGNPD
BAN57590.1	CTDGRIDCF	LTHRRPIIHA	PSVIERRGNPD
BAN57593.1	CTDGRIDCF	LTHRRPIIHA	PSVIERRGNPD
BAN63764.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
BAN63767.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
BAN63770.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
BAN63773.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD
AGT38396.1	DAEGRILTCK	LTHRRPIIHA	PSVIERRGNPD
AGT96613.1	ASPGRLECF	LTHRRPIIHA	PSVIERRGNPD

Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd site	Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd site
AGT96616.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	APG39844.1	TSGRLDCC	TLTERRPIIHS	SVIIRGNPDC
AGT96619.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	BAX25173.1	GPIRLECR	TLTERRPIIHS	SVIIRGNPDC
AGT96622.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	BAX25176.1	GPIRLECR	TLTERRPIIHS	SVIIRGNPDC
AGT96625.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	ARJ36906.1	GPGRLECR	TLSERRPIIHS	SVIIRGNPDC
AHC54587.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGYPDC	BBA54710.1	ADGRIDCE	TLTERRPIIHS	SVIIRGNPDC
AHC54590.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGYPDC	BBA54718.1	SPIRLECR	TLTERRPIIHS	SVIIRGNPDC
AHC55358.1	AGPGRLLECR	YTLTERRPIIHS	STIIRGNPDC	BBA80360.1	ADGRIDCE	TLTERRPIIHS	SVIIRGNPDC
BAO31621.1	DTNRLVCR	YSLTERRPIIHS	SVVIRGAVEC	BBA80363.1	ADGRIDCE	TLTERRPIIHS	SVIIRGNPDC
AHF50467.1	DAEGRLETCR	YTLTERRPIIHS	SVVIRSTIS	BBA80366.1	ADGRIDCE	TLTERRPIIHS	SVIIRGNPDC
AHL67385.1	AGPGRLLECR	YTLLERRPIIHS	SVIIRGNPDC	BBA80369.1	ADGRIDCE	TLTERRPIIHS	SVIIRGNPDC
AHL67388.1	AGPGRLLECR	YTLLERRPIIHS	SVIIRGNPDC	ATN40272.1	SPGRLECR	TLSERRPIIHS	STIIRGDPDC
AHY61295.1	SYPGRIEICK	YTLTERRPIIHS	SVIIRGTPDC	ATN40275.1	SPGRLECR	TLSERRPIIHS	STIIRGDPDC
AHY61298.1	SHPGRIEICK	YTLTERRPIIHS	SVIIRGSPDC	sp P29324.1	VDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
AID70919.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGSPDC	sp O03495.1	TSGRLDCC	TLTERRPIIHS	SVIIRGNPDC
AI120308.1	AGPIRLECR	YTLSERRPIIHS	SVIIRGNPDC	sp O04610.1	VDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
AIJ27593.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	sp P33424.2	VDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
AIP98381.1	---EPEVY	YTVTERRPIIHS	AALEKARTI	sp O9IVZ9.1	ADGRLETCR	TLTERRPIIHS	SVIIRGTDI
BAP86929.1	DADGRLETCR	YTLSERRPIIHS	SVIIRGTIDC	sp O9WC28.1	VDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
BAP86932.1	DADGRLETCR	YTLSERRPIIHS	SVIIRGTIDC	sp O81862.1	VDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
BAC95549.1	ASPGRLLECR	YTLSERRPIIHS	SVIIRGNPDC	ADM35748.1	---EPEVY	TVTERRPIIHS	AAVEKARTI
AHC70111.2	AGMIFAAER	YTLSERRPIIHS	AEAIRIPDI	ADM35754.1	---EPEVY	TVTERRPIIHS	AAVEKARTI
AKO44212.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGSPDC	sp O6J8G2.1	SPGRLECR	TLTERRPIIHS	SVIIRGNPDC
BAS29763.1	DTDGRLETCR	YTLTERRPIIHS	SVIIRGTPDC	BAS66972.1	TDCRIDCE	TLTERRPIIHS	SVIIRGNPDC
ALH42675.1	ASPGRLLECR	YTLTERRPIIHS	STIIRGNPDC	BAL04585.1	GPIRLECR	TLTERRPIIHS	SVIIRGNPDC
ALM55661.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNLDC	BAT32882.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
ALN69915.1	SHPGRIEICK	YTLAERRPIIHS	SVIIRGTPDC	BAT32885.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
BAT70058.1	---LADVY	YTECHRRPIIHS	AAVEKARTI	AOG18209.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
BAT70062.1	---LADVY	YTECHRRPIIHS	AAVEKARTI	AOG18211.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
ALY05712.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AOG18214.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
ALY05715.1	AGPIRLECR	YTLXERRPIIHS	SVIIRGNPDC	AOG18217.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
ALY05718.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AOG18220.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
ALY05721.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AOG18223.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
AMA22617.1	AGPIRLECR	YTLVERRPIIHS	SVIIRGNPDC	AOG18226.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
AMD16213.1	ANPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	BAS32609.1	AEGRLVCK	TPTERRPIIHS	SVIIRGNPDC
AMD16216.1	ASPGRLLECR	YTLSERRPIIHS	STIIRGNPDC	sp O9YLR1.1	SPGRLECR	TLSERRPIIHS	SVIIRGNPDC
AMD16219.1	ANPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AFP19142.1	YDSKTRRY	CRGAVNVII	VCPVAP---
BAU50952.1	DADGRLETCR	YTLTERRPIIHS	SVVIRGTIG	YP00657650	YDSKTRRY	CRGAVNVII	VCPVAP---
ANQ45732.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AAC97208.1	SPGRLECR	TLTERRPIIHS	SVIIRGNPDC
ANW09725.1	AGPGRLLECR	YTLTERRPIIHS	SVIIRGNPDC	BAB793004.1	SPGRLECR	TLTERRPIIHS	SVIIRGNPDC
BAV53288.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AAM92182.1	SPGRLECR	TLTERRPIIHS	SVIIRGNPDC
BAV53291.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	BAC75627.1	ADGRLETCR	TLTERRPIIHS	SVIIRGNPDC
BAV53291.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	AAT02424.1	ADGRLETCR	TLTERRPIIHS	SVIIRGNPDC
BAV57605.1	AGPIRLECR	YTLTERRPIIHS	SVIIRGNPDC	ABB88699.2	ADGRLETCR	TLTERRPIIHS	SVIIRGNPDC
AOR52320.1	VDSNRLEDCR	YTLAERRPIIHS	SVIIRGTPDC	ACH879553.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
AOR52323.1	VDSNRLEDCR	YTLAERRPIIHS	SVIIRGTPDC	ACH879556.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC
AOR52326.1	VDSNRLEDCR	YTLAERRPIIHS	SVIIRGTPDC	ACH879559.1	SPGRLECR	TLSERRPIIHS	STIIRGNPDC

Accession no.	Factor Xa site	Thrombin 1 st site	Thrombin 2 nd site
BAH22710.1	ASPGRLLECR	TLTERRPIIHS	STIIRGNPDC
BAH22713.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
BAH22716.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC
BAH22719.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC
ACV66469.1	AGPIRLECR	TLTERRPIIHS	SVIIRGNPDC
ACV66480.1	ASPGRLLECR	TLTERRPIIHS	STIIRGNLDC
ACZ67487.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
ADC32804.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
AFY63501.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
AGV22365.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
AHZ44444.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC
AHZ44447.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC
AKE33870.1	DADGRLETCR	TLTERRPIIHS	SVIIRSTVD
AMH40418.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC
ART85723.1	ASPIRLECR	TLTERRPIIHS	SVIIRGNPDC