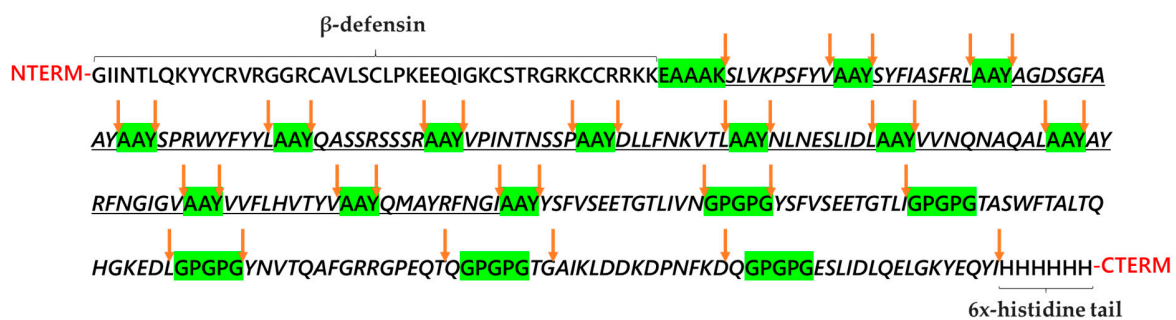


**Supplementary Figure S1. Phylogenetic analysis between human coronaviruses:** ML tree was constructed using A. envelope and B. membrane nucleotide sequences from human coronaviruses.



**Supplementary Figure S2. Multisubunit vaccine construct:** Figure shows different components of our vaccine construct, namely N-terminus (NTERM),  $\beta$ -defensin amino acid sequence, followed by EAAK, AAY and GPGPG linkers, epitopes and C-terminus (CTERM) 6x Histidine sequence. In between linkers, CD8 T-cell (Italics and underlined) and B-cells (italics) epitopes are shown. The arrows indicate proteasomal sites within the construct that may give rise to mature epitopes.

## A. Conservation of Envelope Protein Sequence in HCoV

	1	10	20	30	40	50																																															
COV2	MY	S..FV	SEET	GTL	IVN	S	VL	LF	LA	F	VV	F	LLV	T	LA	I	L	T	ALR	LC	AYC	CNIV	N	VS	LV	K	P	SF	YV																								
SARS	MY	S..FV	SEET	GTL	IVN	S	VL	LF	LA	F	VV	F	LLV	T	LA	I	L	T	ALR	LC	AYC	CNIV	N	VS	LV	K	P	TV	YV																								
MERS	ML	P..FV	Q	ERIGLF	IVN	FF	I	FT	VV	CA	I	T	LLV	C	MA	F	L	T	ATR	LC	VQC	MTGF	N	TL	LV	Q	P	AL	YI																								
NL63	MF	L..RL	I	DDNG.I	VLN	S	IL	WL	LV	M	I	FF	VL	A	M	F	I	K	LI	Q	LC	F	T	C	H	Y	F	S	R	L.Q	P	VYKI																					
229E	MF	L..KL	V	DDHA.L	VVN	V	LL	WC	VV	L	I	V	I	LLV	C	I	T	I	K	LI	K	LC	F	T	C	H	M	F	C	N	R	T	VY	G	P	I	K	N	V														
OC43	MF	MADAYL	A	D	T	VWY	V	G	Q	I	I	F	I	V	A	I	C	L	L	V	T	I	V	V	A	F	L	A	T	F	K	LC	I	Q	L	C	G	M	C	N	T	L	V	L	S	P	S	I	YV				
HKU1	MV	D..V	F	F	T	D	T	A	WY	I	G	Q	I	F	F	L	V	L	S	C	V	I	F	L	I	F	V	V	A	L	L	A	T	I	K	LC	I	Q	I	C	G	F	C	N	I	F	I	I	S	P	S	A	YV

	60	70													
COV2	YSRVKN	L.....NSSRV	P	D..LL	V										
SARS	YSRVKN	LN.....SSEGV	P	D..LL	V										
MERS	YNTGRS	VYVKFQDSKP	P	L	P	D..EW	V								
NL63	FLAYQDYM	.....QIAP	V	P	AEVLN	V									
229E	YHIYQS	YM.....HID	P	F	P	KRVIDF									
OC43	FN	RG	RQ	F	YEFYN	DIK	P	P	V	L	D	V	D	D	V
HKU1	YN	RG	RQ	L	YKSYSEHVI	P	S	T	L	D	D	L	I		

## B. Conservation of Membrane Protein Sequence in HCoV

	1	10	20	30	40	50							
COV2	MADSN	GT.....ITVE	ELKKL	LEQWN	LVIGFL	LTWIC	LQFAY	ANRN	RF	LYIIK	LIF	LW	
SARS	MADNGT	.....ITVE	ELKQL	LEQWN	LVIGFL	LAWI	MLQFAYS	NRN	RF	LYIIK	LV	LW	
MERS	MSNM	TQ.....LTEAQ	IIAI	IKDWN	FAWS	LIFLL	ITVLOYGYP	SRS	MT	VYVFK	MF	VW	
NL63	MSNS	S.....VPLS	EVYVH	LRNWN	FSWN	LILTVFI	VVLOYGHY	KYS	R	LYGLK	MS	VW	
229E	MSND	NC.....TG	DIVTH	LKNWN	FGWN	VILTIFI	VVLOFGHY	KYS	R	LYGLK	ML	VW	
OC43	MS	SKTTPAPVYIWTAD	EAIKF	LKEWN	FSLG	II	ILFIT	IILOFGYT	SRS	MF	VYVIK	MI	LW
HKU1	MN	KSEFFP.....QFTS	DQATF	LKEWN	FSLG	II	ILFIT	IILOFGYT	SRS	MF	VYLIK	MI	LW

### Conserved epitope 1

	60	70	80	90	100	110	
COV2	LLWPVTLACFVL	AAV..	YRINWITGGIAIAMAC	LVGLMWLS	YFIASFRLFART	RSMWSEN	
SARS	LLWPVTLACFVL	AAV..	YRINWVTGGIAIAMAC	IVGLMWLS	YFVASFRLFART	RSMWSEN	
MERS	LLWPSSMALSIF	SAV..	YPIDLASQII	SGIVAAV	SAMMWIS	YFVQSIRLFMRTGSWWSEN	
NL63	CLWPLVLALSIF	DCFVN	FNDWVFFGF	SILMSIIT	LCLWVM	YFVNSFRLWRRVKTFWAFN	
229E	LLWPLVLALSIF	DTWAN	WDSNWAFAVAF	SFLMAVST	LVMWVM	YFANSIRLFRARRTFWAWN	
OC43	LMWPLTIILTIF	NCV..	YALNNVYLGLSIVFTI	VAIIMWIV	YFVNSIRLFRIRTGSFWSEN		
HKU1	LMWPLTITLTIF	NCF..	YALNNAFLAF	SIVFTIIS	IVIWI	L	YFVNSIRLFIRTGSWWSEN

	120					130					140					150					160					170																																
COV2	PETN	I	L	N	V	P	L	H	G	T	I	L	T	R	P	L	L	E	S	E	L	V	T	G	A	V	I	L	R	G	H	L	R	T	A	G	H	L	G	R	.	C	D	I	K	D	L	P	K	E	I	T	V	A	K			
SARS	PETN	I	L	N	V	P	L	R	G	T	I	V	T	R	P	L	M	E	S	E	L	V	I	G	A	V	I	I	R	G	H	L	R	M	A	G	H	S	L	G	R	.	C	D	I	K	D	L	P	K	E	I	T	V	A	K		
MERS	PETN	C	L	L	N	V	P	F	G	T	T	V	T	R	P	L	V	E	D	S	T	S	V	T	A	V	V	T	N	G	H	L	K	M	A	G	M	H	S	F	G	R	.	C	D	Y	D	R	L	P	N	E	V	T	V	A	K	
NL63	PETN	A	I	I	S	L	Q	V	Y	G	H	N	Y	Y	L	P	V	M	A	A	P	T	G	V	I	L	T	L	L	S	G	V	L	L	V	D	G	H	K	I	A	T	R	V	Q	V	G	O	L	P	K	Y	V	I	V	A	T	K
229E	PEVN	A	I	T	V	T	T	V	L	G	Q	T	Y	Y	Q	P	I	Q	A	P	T	G	I	T	V	T	L	L	S	G	V	L	L	V	D	G	H	R	L	A	S	G	V	Q	V	H	N	L	P	E	Y	M	T	V	A	K		
OC43	PETN	N	L	M	C	I	D	M	K	G	T	M	Y	V	R	P	I	E	D	Y	H	T	L	T	V	T	I	I	R	G	H	L	Y	I	Q	G	I	K	L	G	T	G	Y	S	L	A	D	L	P	A	Y	M	T	V	A	K		
HKU1	PETN	N	L	M	C	I	D	M	K	G	K	M	F	V	R	P	V	I	E	D	Y	H	T	L	T	A	T	V	I	R	G	H	L	Y	I	Q	G	V	K	L	G	T	G	Y	T	L	S	D	L	P	V	Y	M	T	V	A	K	

		180	190	200	210	220
COV2	S	RTLSYYKLGLASQ	RVAGDSGFAAYSRYRIGNYK	LNTDHSSSSDNIALLVQ..		
SARS	S	RTLSYYKLGLASQ	RVGTDSDGFAAYNRYRIGNYK	LNTDHAGSNDNIALLVQ..		
MERS	P	NVLIALKMKRQ	SYGTNSGVAIYHRYKAGNYRSP	.PITADIELALLRA..		
NL63	PS	TTIIVCDRVGRSVNETS	QTGWAFYVRAKHGDFSGVASQEGVLSEREKLLHLI			
229E	PS	TTIIVCDRVGRSVNETS	QTGWAFYVRAKHGDFSAVSPSPMSNMENERLLHFF			
OC43	V	THLCTYKRGFLDRISD	TSGFAYVVKSKVGNRYLPS	TQKGSGMDTALLRNNI		
HKU1	V	QVLCTYKRAFLDKLDV	NSGFAYVVKSKVGNRYLPS	.SKPSGMDTALLRA..		



### C. Conservation of Nucleocapsid Protein Sequence in HCoV

	1	10	20	30	40	
COV2	.MSDNGPQ.N	QRNAPR.IT	.....	FGGPSTG	SNQNGERSGARS	KQRPPQGLP..
SARS	.MSDNGPQSN	QRSAPR.IT	.....	FGGPTDSTD	NNQNGGRNGARP	KQRPPQGLP..
MERS	.....MA	SFAAPRAVS	.....	FADNNDITN	TNLSRGRG....	RNPKPRAAP..
NL63	.....	MASVN	.....	WADD	.....	RAARKKFP..
229E	.....	MATVK	.....	WADASE	.....	PQRGRQGRIP..
OC43	MSFTPGKQSS	SRASSGNRS	SGNGILK..	WADQSDQFR	NVQTRGRR.AQP	KQTATSQQPSG
HKU1	MSYTPGHHAG	SRSSSGNRS	.GILKKT	WVDQSE	RSHQTYNRGR	RK.PQP.KFTVSTQPPQ..

	50	60	70	80	90	100				
COV2	..NNTASWFTALTQ	HGKE.DLKF	PRGQGVPI	INTNSSPD	DQIGYYRRATRR	.IRGGDGKM				
SARS	..NNTASWFTALTQ	HGKE.ELRF	PRGQGVPI	INTNSGPD	DQIGYYRRATRR	.VRGGDGKM				
MERS	..NNTVSWYTGLTQ	HGKV.PLTF	PPGQGVPL	NANSTPAQ	NAGYWRQDRK	.INTGNG.I				
NL63	.....PSFYMP	LLVSSDKAPY	RVIPRNLVPI	GKG.NKDE	QIGYWNVQER	.WRMRRGQR				
229E	.....YSLYSPLLV	DSEQ.PWKV	IPRNLVPI	NKK.DKNKL	IGYWNVQKR	.FRTRKGKR				
OC43	GNVVPYY	SWFSGITQ	FQKGKEFE	FAEGQGVPI	APGVPAT	EAKGYWYRHNRRSF	KTADGNQ			
HKU1	GNTIPHY	SWFSGITQ	FQKGRDF	KFPD	GQGVPI	AYGIPPS	EAKGYWYKHNR	RRSF	KTAD	GQQ

#### Conserved epitope 1

	110	120	130	140	150
COV2	KDLSPRWYFYLLGTGPEAGLPY	GANKDGIIVWVATEGALNTPKDHIGTRNPANNAIV...			
SARS	KELSPRWYFYLLGTGPEASLPY	GANKBGIIVWVATEGALNTPKDHIGTRNPNNNAIV...			
MERS	KQLAPRWYFYLLGTGPEALLPF	RAVKDGIIVWVHEDEGATDAP.STFGTRNPNNDSAIV...			
NL63	VDLPPKVLHFFYYLLGTGPHKDLKF	RQRSDGVVWVAKEGAKTVN.TSLGNRRKRNRQKPLEPKFS			
229E	VDLSPKLHFFYYLLGTGPHKDAKF	RERVEGVVWVADEGAKTEP.TGYGVRRKNSSEPEIPHFN			
OC43	RQLLPRWYFYLLGTGPHAKDQY	GTDIDGVYWVASNQADVNTPADIVDRDPSSDEAIP...			
HKU1	KQLLPRWYFYLLGTGPHYANASY	GESHBGIFWVASHQADTSIPSDVSARDPTIQEAI			

	160	170	180	190	200	210	
COV2	LQLPQGTTL	LPKGFYAEGSR	GGSSQASSR	SSSR..	SRNSSRNST	PGSSRGTS	SPARMAGNGGD
SARS	LQLPQGTTL	LPKGFYAEGSR	GGSSQASSR	SSSR..	SRGNSRNST	PGSSRGNS	SPARMASGGGE
MERS	TQFAPGTL	LPKNFHIEGT	GGNSQSSSR	ASSV..	SRNSSRSSSS	QGSSRSGN	..STRGTS
NL63	IALPPELS	VVE..FEDRS	NNSSSRASSR	SSSTR	RNNSRDSSRST	SRRQSSRT	SDSNQSSDLV
229E	QKLPGNV	TVEE..EPDS	RAPSRSSQSS	SSSR..	SRGESKSS	SRNPFSSDR	..NHNSQDDIM
OC43	TRFPPTV	LPQGYIEGS	G.....	RSAPN..	SRSTSTSS	RASSAGS	..RSRANS
HKU1	TRFSPGTI	LPQGYVEGS	G.....	RSASN..	SRPGSSRS	SSRGPNNR	S..LSRSNS

	220	230	240	250
COV2	AA <b>L</b> <b>A</b> <b>L</b> L.....	<b>L</b> D <b>R</b> <b>L</b> <b>N</b> <b>L</b> ESKMSGKG <b>Q</b> <b>Q</b> <b>Q</b> <b>G</b> .....	<b>Q</b> T <b>V</b> <b>T</b> <b>K</b> <b>K</b> <b>S</b> <b>A</b> <b>A</b> <b>E</b> <b>A</b>	
SARS	TA <b>L</b> <b>A</b> <b>L</b> L.....	<b>L</b> D <b>R</b> <b>L</b> <b>N</b> <b>L</b> ESKMSGKG <b>Q</b> <b>Q</b> <b>Q</b> <b>G</b> .....	<b>Q</b> T <b>V</b> <b>T</b> <b>K</b> <b>K</b> <b>S</b> <b>A</b> <b>A</b> <b>E</b> <b>A</b>	
MERS	SG <b>I</b> <b>G</b> <b>A</b> <b>V</b> GGD <b>L</b> <b>L</b> Y.....	<b>L</b> D <b>L</b> <b>L</b> <b>N</b> <b>R</b> LQALESgK <b>V</b> <b>K</b> <b>Q</b> <b>S</b> <b>Q</b> <b>P</b> .....	<b>K</b> <b>V</b> <b>I</b> <b>T</b> <b>K</b> <b>K</b> <b>D</b> <b>A</b> <b>A</b> <b>A</b> <b>A</b>	
NL63	AA <b>V</b> <b>T</b> <b>L</b> <b>A</b> L <b>K</b> <b>N</b> L <b>G</b> <b>F</b> <b>D</b> ...N <b>Q</b> <b>S</b> <b>K</b> <b>S</b> <b>P</b> <b>S</b> <b>S</b> <b>S</b> <b>G</b> <b>T</b> <b>S</b> <b>T</b> <b>P</b> <b>K</b> <b>K</b> <b>F</b> <b>N</b> <b>K</b> <b>P</b> .....	<b>L</b> <b>S</b> <b>Q</b> <b>P</b> <b>R</b> <b>A</b> <b>D</b> <b>K</b> <b>P</b> <b>S</b> <b>Q</b>		
229E	KA <b>V</b> <b>A</b> <b>A</b> <b>A</b> L <b>K</b> <b>S</b> L <b>G</b> <b>F</b> <b>D</b> <b>K</b> <b>P</b> <b>Q</b> E <b>K</b> <b>D</b> <b>K</b> <b>S</b> A <b>K</b> <b>T</b> <b>G</b> <b>T</b> <b>P</b> <b>K</b> <b>P</b> <b>S</b> <b>R</b> <b>N</b> <b>Q</b> <b>S</b> PASSQSA <b>A</b> <b>K</b> <b>I</b> <b>L</b> <b>A</b> <b>R</b> <b>S</b> <b>Q</b> S <b>S</b> <b>E</b> <b>T</b> <b>K</b> <b>E</b> <b>Q</b> <b>K</b> <b>H</b> <b>E</b>			
OC43	T <b>P</b> <b>T</b> <b>S</b> <b>G</b> <b>V</b> <b>T</b> <b>P</b> <b>D</b> <b>M</b> <b>A</b> <b>D</b> <b>Q</b> ... <b>I</b> <b>A</b> <b>S</b> <b>L</b> <b>V</b> <b>L</b> A <b>K</b> <b>L</b> <b>G</b> <b>K</b> <b>D</b> <b>A</b> <b>T</b> <b>K</b> <b>P</b> <b>Q</b> <b>Q</b> <b>V</b> <b>T</b> .....	<b>K</b> <b>H</b> <b>T</b> <b>A</b> <b>K</b> <b>E</b> <b>V</b> <b>R</b> <b>Q</b> <b>K</b> <b>I</b>		
HKU1	HS <b>D</b> <b>S</b> <b>I</b> <b>V</b> K <b>P</b> <b>D</b> <b>M</b> <b>A</b> <b>D</b> <b>E</b> ... <b>I</b> <b>A</b> <b>S</b> <b>L</b> <b>V</b> <b>L</b> A <b>K</b> <b>L</b> <b>G</b> <b>K</b> <b>D</b> <b>S</b> <b>K</b> <b>P</b> <b>Q</b> <b>Q</b> <b>V</b> <b>T</b> .....	<b>K</b> <b>Q</b> <b>N</b> <b>A</b> <b>K</b> <b>E</b> <b>I</b> <b>R</b> <b>H</b> <b>K</b> <b>I</b>		

	260	270	280	290	300	310
COV2	SK <b>K</b> <b>P</b> <b>R</b> <b>Q</b> <b>K</b> <b>R</b> <b>T</b> A <b>T</b> <b>K</b> A... <b>Y</b> <b>N</b> <b>V</b> <b>T</b> <b>Q</b> A <b>F</b> <b>G</b> <b>R</b> <b>R</b> <b>G</b> <b>P</b> E <b>Q</b> <b>T</b> <b>Q</b> <b>G</b> <b>N</b> <b>E</b> <b>G</b> <b>D</b> <b>Q</b> <b>E</b> <b>L</b> <b>I</b> <b>R</b> <b>Q</b> <b>G</b> <b>T</b> DYKH <b>W</b> <b>P</b> <b>Q</b> <b>I</b> <b>A</b> <b>Q</b> <b>F</b> <b>A</b> <b>P</b> <b>S</b> <b>A</b> <b>S</b>					
SARS	SK <b>K</b> <b>P</b> <b>R</b> <b>Q</b> <b>K</b> <b>R</b> <b>T</b> A <b>T</b> <b>K</b> Q... <b>Y</b> <b>N</b> <b>V</b> <b>T</b> <b>Q</b> A <b>F</b> <b>G</b> <b>R</b> <b>R</b> <b>G</b> <b>P</b> E <b>Q</b> <b>T</b> <b>Q</b> <b>G</b> <b>N</b> <b>E</b> <b>G</b> <b>D</b> <b>Q</b> <b>D</b> <b>L</b> <b>I</b> <b>R</b> <b>Q</b> <b>G</b> <b>T</b> DYKH <b>W</b> <b>P</b> <b>Q</b> <b>I</b> <b>A</b> <b>Q</b> <b>F</b> <b>A</b> <b>P</b> <b>S</b> <b>A</b> <b>S</b>					
MERS	KN <b>K</b> <b>M</b> <b>R</b> <b>H</b> <b>K</b> <b>R</b> <b>T</b> <b>S</b> <b>T</b> <b>K</b> S... <b>F</b> <b>N</b> <b>M</b> <b>V</b> <b>Q</b> A <b>F</b> <b>G</b> <b>L</b> <b>R</b> <b>G</b> <b>P</b> G <b>D</b> <b>L</b> <b>Q</b> <b>G</b> <b>N</b> <b>E</b> <b>G</b> <b>D</b> <b>L</b> <b>Q</b> <b>L</b> <b>N</b> <b>K</b> <b>L</b> <b>G</b> <b>T</b> E <b>D</b> <b>P</b> <b>R</b> <b>W</b> <b>P</b> <b>Q</b> <b>I</b> <b>A</b> <b>E</b> <b>L</b> <b>A</b> <b>P</b> <b>T</b> <b>A</b> <b>S</b>					
NL63	L <b>K</b> <b>K</b> <b>F</b> <b>R</b> <b>W</b> <b>K</b> <b>R</b> <b>V</b> <b>P</b> <b>T</b> <b>R</b> E... <b>E</b> <b>N</b> <b>V</b> <b>I</b> <b>Q</b> <b>C</b> <b>F</b> <b>G</b> <b>P</b> <b>R</b> <b>D</b> <b>F</b> <b>N</b> H... <b>N</b> <b>M</b> <b>G</b> <b>D</b> <b>S</b> <b>D</b> <b>L</b> <b>V</b> <b>Q</b> <b>N</b> <b>G</b> <b>V</b> <b>D</b> <b>A</b> <b>K</b> <b>G</b> <b>F</b> <b>P</b> <b>Q</b> <b>L</b> <b>A</b> <b>E</b> <b>L</b> <b>I</b> <b>P</b> <b>N</b> <b>Q</b> <b>A</b>					
229E	M <b>Q</b> <b>K</b> <b>P</b> <b>R</b> <b>W</b> <b>K</b> <b>R</b> <b>P</b> <b>N</b> <b>D</b> <b>D</b> <b>V</b> <b>T</b> S <b>N</b> <b>V</b> <b>T</b> <b>Q</b> <b>C</b> <b>F</b> <b>G</b> <b>P</b> <b>R</b> <b>D</b> <b>L</b> <b>D</b> H... <b>N</b> <b>E</b> <b>G</b> <b>S</b> <b>A</b> <b>G</b> <b>V</b> <b>V</b> <b>A</b> <b>N</b> <b>G</b> <b>V</b> <b>K</b> <b>A</b> <b>K</b> <b>G</b> <b>Y</b> <b>P</b> <b>Q</b> <b>F</b> <b>A</b> <b>E</b> <b>L</b> <b>V</b> <b>P</b> <b>S</b> <b>T</b> <b>A</b>					
OC43	L <b>N</b> <b>K</b> <b>P</b> <b>R</b> <b>Q</b> <b>K</b> <b>R</b> <b>S</b> <b>P</b> <b>N</b> <b>K</b> <b>Q</b> ... <b>C</b> <b>T</b> <b>V</b> <b>Q</b> <b>Q</b> <b>C</b> <b>F</b> <b>G</b> <b>K</b> <b>R</b> <b>G</b> <b>P</b> <b>N</b> <b>Q</b> ... <b>N</b> <b>E</b> <b>G</b> <b>G</b> <b>G</b> <b>E</b> <b>M</b> <b>L</b> <b>K</b> <b>L</b> <b>G</b> <b>T</b> S <b>D</b> <b>P</b> <b>Q</b> <b>F</b> <b>P</b> <b>I</b> <b>L</b> <b>A</b> <b>E</b> <b>L</b> <b>A</b> <b>P</b> <b>T</b> <b>A</b> <b>G</b>					
HKU1	L <b>M</b> <b>K</b> <b>P</b> <b>R</b> <b>Q</b> <b>K</b> <b>R</b> <b>T</b> <b>P</b> <b>N</b> <b>K</b> F... <b>C</b> <b>N</b> <b>V</b> <b>Q</b> <b>Q</b> <b>C</b> <b>F</b> <b>G</b> <b>K</b> <b>R</b> <b>G</b> <b>P</b> <b>L</b> <b>Q</b> ... <b>N</b> <b>E</b> <b>G</b> <b>N</b> <b>E</b> <b>M</b> <b>L</b> <b>K</b> <b>L</b> <b>G</b> <b>T</b> N <b>D</b> <b>P</b> <b>Q</b> <b>F</b> <b>P</b> <b>I</b> <b>L</b> <b>A</b> <b>E</b> <b>L</b> <b>A</b> <b>P</b> <b>T</b> <b>I</b> <b>P</b> <b>G</b>					

	320	330	340	350	360
COV2	<b>A</b> <b>F</b> <b>F</b> <b>G</b> <b>M</b> <b>S</b> <b>R</b> <b>I</b> <b>G</b> <b>M</b> <b>E</b> <b>V</b> <b>T</b> <b>P</b> <b>S</b> ..... <b>G</b> <b>T</b> <b>W</b> <b>L</b> <b>T</b> <b>Y</b> <b>T</b> <b>G</b> <b>A</b> <b>I</b> <b>K</b> <b>L</b> <b>D</b> <b>D</b> <b>K</b> <b>D</b> <b>P</b> <b>N</b> <b>F</b> <b>K</b> <b>D</b> <b>Q</b> <b>V</b> <b>I</b> <b>L</b> <b>L</b> <b>N</b> <b>K</b> <b>H</b> <b>I</b> <b>D</b> <b>A</b> <b>Y</b> <b>K</b>				
SARS	<b>A</b> <b>F</b> <b>F</b> <b>G</b> <b>M</b> <b>S</b> <b>R</b> <b>I</b> <b>G</b> <b>M</b> <b>E</b> <b>V</b> <b>T</b> <b>P</b> <b>S</b> ..... <b>G</b> <b>T</b> <b>W</b> <b>L</b> <b>T</b> <b>Y</b> <b>H</b> <b>G</b> <b>A</b> <b>I</b> <b>K</b> <b>L</b> <b>D</b> <b>D</b> <b>K</b> <b>D</b> <b>P</b> <b>Q</b> <b>F</b> <b>K</b> <b>D</b> <b>N</b> <b>V</b> <b>I</b> <b>L</b> <b>L</b> <b>N</b> <b>K</b> <b>H</b> <b>I</b> <b>D</b> <b>A</b> <b>Y</b> <b>K</b>				
MERS	<b>A</b> <b>F</b> <b>M</b> <b>G</b> <b>M</b> <b>S</b> <b>Q</b> <b>F</b> <b>K</b> <b>L</b> <b>T</b> <b>H</b> <b>Q</b> <b>N</b> <b>N</b> ... <b>D</b> <b>D</b> <b>H</b> <b>G</b> <b>N</b> <b>P</b> <b>V</b> <b>Y</b> <b>F</b> <b>L</b> <b>R</b> <b>Y</b> <b>S</b> <b>G</b> <b>A</b> <b>I</b> <b>K</b> <b>L</b> <b>D</b> <b>P</b> <b>K</b> <b>N</b> <b>P</b> <b>N</b> <b>Y</b> <b>N</b> <b>K</b> <b>W</b> <b>L</b> <b>E</b> <b>L</b> <b>L</b> <b>E</b> <b>Q</b> <b>N</b> <b>I</b> <b>D</b> <b>A</b> <b>Y</b> <b>K</b>				
NL63	<b>A</b> <b>L</b> <b>F</b> <b>F</b> <b>D</b> <b>S</b> <b>E</b> <b>V</b> <b>S</b> <b>T</b> <b>D</b> <b>E</b> <b>V</b> <b>G</b> <b>D</b> ... <b>N</b> <b>V</b> <b>Q</b> <b>I</b> <b>T</b> <b>Y</b> <b>T</b> <b>Y</b> <b>K</b> <b>M</b> <b>L</b> <b>V</b> <b>A</b> <b>K</b> <b>D</b> <b>N</b> <b>K</b> <b>N</b> <b>L</b> <b>P</b> <b>K</b> <b>F</b> <b>I</b> <b>E</b> <b>Q</b> <b>I</b> <b>S</b> .....				
229E	<b>A</b> <b>M</b> <b>L</b> <b>F</b> <b>D</b> <b>S</b> <b>H</b> <b>I</b> <b>V</b> <b>S</b> <b>K</b> <b>E</b> <b>S</b> <b>G</b> <b>N</b> ... <b>T</b> <b>V</b> <b>V</b> <b>L</b> <b>T</b> <b>F</b> <b>T</b> <b>T</b> <b>R</b> <b>V</b> <b>T</b> <b>V</b> <b>P</b> <b>K</b> <b>D</b> <b>H</b> <b>P</b> <b>H</b> <b>L</b> <b>G</b> <b>K</b> <b>F</b> <b>L</b> <b>E</b> <b>E</b> <b>L</b> <b>N</b> .....				
OC43	<b>A</b> <b>F</b> <b>F</b> <b>F</b> <b>G</b> <b>S</b> <b>R</b> <b>L</b> <b>E</b> <b>L</b> <b>A</b> <b>K</b> <b>V</b> <b>Q</b> <b>N</b> <b>L</b> <b>S</b> <b>G</b> <b>N</b> <b>P</b> <b>D</b> <b>E</b> <b>P</b> <b>Q</b> <b>K</b> <b>D</b> <b>V</b> <b>Y</b> <b>E</b> <b>L</b> <b>R</b> <b>Y</b> <b>N</b> <b>G</b> <b>A</b> <b>I</b> <b>R</b> <b>F</b> <b>D</b> <b>S</b> <b>T</b> <b>L</b> <b>S</b> <b>G</b> <b>F</b> <b>E</b> <b>T</b> <b>I</b> <b>M</b> <b>K</b> <b>V</b> <b>L</b> <b>S</b> <b>E</b> <b>N</b> <b>L</b> <b>N</b> <b>A</b> <b>Y</b> <b>Q</b>				
HKU1	<b>A</b> <b>F</b> <b>F</b> <b>F</b> <b>G</b> <b>S</b> <b>K</b> <b>L</b> <b>E</b> <b>L</b> <b>F</b> <b>K</b> <b>R</b> <b>D</b> <b>S</b> ... <b>D</b> <b>A</b> <b>D</b> <b>S</b> <b>P</b> <b>S</b> <b>K</b> <b>D</b> <b>T</b> <b>F</b> <b>E</b> <b>L</b> <b>R</b> <b>Y</b> <b>S</b> <b>G</b> <b>S</b> <b>I</b> <b>R</b> <b>F</b> <b>D</b> <b>S</b> <b>T</b> <b>L</b> <b>P</b> <b>G</b> <b>F</b> <b>E</b> <b>T</b> <b>I</b> <b>M</b> <b>K</b> <b>V</b> <b>L</b> <b>K</b> <b>E</b> <b>N</b> <b>L</b> <b>D</b> <b>A</b> <b>Y</b> <b>V</b>				

	370	380
COV2	..... <b>T</b> <b>F</b> <b>P</b> <b>P</b> <b>T</b> <b>D</b> <b>F</b> <b>S</b> <b>K</b> <b>Q</b> <b>L</b> <b>Q</b> <b>Q</b> <b>S</b> .....	<b>M</b> <b>S</b> <b>S</b> <b>A</b> <b>D</b> <b>S</b> <b>T</b> <b>Q</b> <b>A</b> ...
SARS	..... <b>T</b> <b>F</b> <b>P</b> <b>P</b> <b>T</b> <b>E</b> <b>P</b> <b>K</b> <b>K</b> <b>D</b> <b>K</b> <b>K</b> <b>K</b> <b>T</b> <b>D</b> <b>E</b> <b>A</b> <b>Q</b> <b>P</b> <b>L</b> <b>P</b> <b>Q</b> <b>R</b> <b>Q</b> <b>K</b> <b>K</b> <b>Q</b> <b>P</b> <b>T</b> <b>V</b> <b>T</b> <b>L</b> <b>L</b> <b>P</b> <b>A</b> <b>A</b> <b>D</b> <b>M</b> <b>D</b> <b>D</b> <b>F</b> <b>S</b> <b>R</b> <b>Q</b> <b>L</b> <b>Q</b> <b>N</b> <b>S</b> <b>M</b> <b>S</b>	
MERS	..... <b>T</b> <b>F</b> <b>P</b> <b>K</b> <b>K</b> <b>E</b> <b>K</b> <b>K</b> <b>Q</b> <b>K</b> <b>A</b> <b>P</b> <b>K</b> <b>E</b> ... <b>E</b> <b>S</b> <b>T</b> <b>D</b> <b>Q</b> <b>M</b> <b>S</b> <b>E</b> <b>P</b> <b>P</b> <b>K</b> <b>E</b> <b>Q</b> <b>R</b> <b>V</b> <b>Q</b> <b>G</b> ... <b>S</b> <b>I</b> <b>T</b> <b>Q</b> <b>R</b> <b>T</b> <b>R</b> <b>T</b> <b>R</b> <b>P</b> <b>S</b> <b>V</b> <b>Q</b> <b>P</b>	
NL63	..... <b>A</b> <b>F</b> <b>T</b> <b>K</b> <b>P</b> <b>S</b> <b>S</b> <b>I</b> <b>K</b> <b>E</b> <b>M</b> <b>Q</b> <b>S</b> <b>Q</b> <b>S</b> <b>S</b> <b>H</b> <b>V</b> <b>V</b> <b>Q</b> <b>N</b> <b>T</b> <b>V</b> <b>L</b> <b>N</b> <b>A</b> <b>S</b> <b>I</b> <b>P</b> <b>E</b> <b>S</b> <b>K</b> <b>P</b> ... <b>L</b> <b>A</b> <b>D</b> <b>D</b> <b>D</b> <b>S</b> <b>A</b> <b>I</b> <b>I</b> <b>E</b> <b>I</b> <b>V</b> <b>N</b>	
229E	..... <b>A</b> <b>F</b> <b>T</b> <b>R</b> <b>E</b> <b>M</b> <b>Q</b> <b>Q</b> <b>Q</b> <b>P</b> <b>L</b> <b>L</b> <b>N</b> <b>P</b> <b>S</b> <b>A</b> <b>L</b> <b>E</b> <b>F</b> <b>N</b> <b>P</b> <b>S</b> <b>Q</b> <b>T</b> <b>S</b> <b>P</b> <b>A</b> <b>T</b> <b>V</b> <b>E</b> ... <b>P</b> ... <b>V</b> <b>R</b> <b>D</b> <b>E</b> <b>V</b> <b>S</b> <b>I</b> <b>E</b> <b>T</b> <b>D</b> <b>I</b> <b>I</b> <b>D</b>	
OC43	<b>Q</b> <b>Q</b> <b>D</b> <b>G</b> <b>M</b> <b>M</b> ... <b>N</b> <b>M</b> <b>S</b> <b>P</b> <b>K</b> <b>P</b> <b>Q</b> <b>R</b> <b>Q</b> <b>R</b> <b>G</b> <b>H</b> <b>K</b> <b>N</b> <b>G</b> <b>Q</b> <b>G</b> <b>E</b> <b>N</b> <b>D</b> <b>N</b> <b>I</b> <b>S</b> <b>V</b> <b>A</b> <b>V</b> <b>P</b> <b>K</b> <b>S</b> <b>R</b> <b>V</b> <b>Q</b> <b>Q</b> <b>N</b> <b>K</b> <b>S</b> <b>I</b> <b>E</b> <b>L</b> <b>T</b> <b>A</b> <b>E</b> <b>D</b> <b>I</b> <b>S</b> <b>L</b> <b>L</b> <b>K</b> <b>K</b> <b>M</b> <b>D</b>	
HKU1	<b>N</b> <b>S</b> <b>N</b> <b>Q</b> <b>N</b> <b>T</b> <b>V</b> <b>S</b> <b>G</b> <b>S</b> <b>L</b> <b>S</b> <b>P</b> <b>K</b> <b>P</b> <b>Q</b> <b>R</b> <b>K</b> <b>R</b> <b>G</b> <b>V</b> <b>K</b> <b>Q</b> <b>S</b> <b>P</b> <b>E</b> <b>S</b> <b>F</b> <b>D</b> <b>S</b> <b>L</b> <b>N</b> <b>L</b> <b>S</b> <b>A</b> <b>D</b> <b>T</b> <b>Q</b> <b>H</b> <b>I</b> <b>S</b> <b>N</b> ... <b>D</b> <b>E</b> <b>T</b> <b>P</b> <b>E</b> <b>D</b> <b>H</b> <b>S</b> <b>L</b> <b>L</b> <b>A</b> <b>T</b> <b>L</b> <b>D</b>	

COV2	.....
SARS	GASADSTQA.
MERS	GPMIDVNTD.
NL63	EVLH.....
229E	EVN.....
OC43	EPYTEDTSEI
HKU1	DPYVEDSVA.

## D. Conservation of Spike Protein Sequence in HCoV

	1	10	20	
COV2	MFVFLVLL	PLV.....	SSQCVN.....	LTTRTQLPP.....A
SARS	MFIFLLFL	TLT.....	SGSDLD.....	RCTTFDDVQAPN.....YT
MERS	MFVFLVLL	TPTESYVDVGPDSVK	SACIEVDIQQTFFDKT	WFRP.....
NL63	MFVFLVLL	PLASCFFTCN.....	SNANLSMLQLGVPDNSS	ITVTGLLPHTHWICANQS
229E	MFVLLVA	.....	.....	.....
OC43	MFVLLVLL	LPT.AFAVIGD...LNCPL	DPRLKGSFNNRD	TGPFSI.....STDT
HKU1	MLLIIFIL	PTT..LAVIGDFNCTNFAL	IN.....DKN	TVTVPRI.....SEYV

	30	40	50	60	70	80
COV2	YTNSTFRGVYYP	PDVKFRS	SVLHST	QDLF..L	FFSNVTWFHATHV	GTNGTKRFDN....
SARS	QHTSSMRGVYYP	PDEIFRS	DTLYLT	QDLF..L	FFSNVTGFHT	INH.....FGN....
MERS	IDVSKADGIIY	POGRYS	NITITY	QGLF..L	PYQG.DHGDY	VYSAGHATGTPQK....
NL63	TSVYSANGFY	IDVGNHR	SAPALH	IGYDVNQ	YYIYVTNEIG	LNASVTLKICKFGINTTF
229E	.....	.....	.....	.....	.....	.....
OC43	VDDVINGLGTYY	VLDRVYL	NTTLFL	NGY...L	FTSGSTYRNMA	LKGTDLSTLWFKP....
HKU1	VDDVSYGLGTYY	ILDRVYL	NTTLFL	NGY...L	EKSGANFRDLS	LKGTITLSTLWYQK....

	90	100	110
COV2	PVLP.....	FNDGVYFASTEKSNIIRG.....	WIFGTTLDSEKT..
SARS	PVIP.....	FKDGIYFAATEKSNVVRG.....	WVFGSTMNKS..
MERS	LFVANYSQDVQK	FANGFVVRIGAAANSTGVI	ISPSTSATIRKIYPAFMLGSSVGNFSDG
NL63	DFLSNSSSSFD	IVNLLFTEQLGAPLGITI..	SGETVRLHLYNVTRTFYVFAA.....
229E	.....	.....	.....
OC43	PFLSD.....	FINGIFAKVKNTKVKFDG.....	VMYSEFPAITIGSTFVNTS..
HKU1	PFLSD.....	FNNGIFSRVKNTKLYVNK.....	TLYSEFSTIVIGSVFIINS..

	120	130	140	150
COV2	.....QSLTIVNN.....	ATNVVTKVCE	QFCND	PFLGVYYHKNNKSWMES
SARS	.....QSVIIINN.....	STNVVIRACN	FELCDN	PFFAVSKPMGTQ...T
MERS	KMGRFNNH	TLVLLP.....	DGCGTLLRAFYCILE	PRSGNHCPAGNS..YTS
NL63	.....YKLTKLS.....	VKCYFNYS	CVFSVFNAT	VIVNVITHNGR....
229E	.....YALLHIA.....	.....	GCQTIN	GTIN.....
OC43	.....YSVVVQ	PRNTINSTQDGVNKLQGLLE	VSVCQYNMCEY	PHTICHPNLGNH..FKE
HKU1	.....YTIVVQPH.....	.....NGVLEITACQYTM	CEYFHTICK	SKGSS..RNE

	160	170	180	190	200
COV2	EFRRVYSANN	CT.FEYVS	QPFLLMDLEGKQGNFKNLRE	FVFKNIDG...	Y..FKTYSKHT
SARS	HTMIFDNAFN	CT.FEYIS	DAFSLDVSEKSGNFKHLRE	FVFKNKDG...	F..LYVYKGYQ
MERS	FATYHTPAT	CDGNGYNNR	ASLNSFKEYFNLRNCTFM	YTYNITEDEILE	W..FGITQTAQ
NL63	..VVNYTV	CDDCN..GYTDN	IFSVQDGRIPNG	..FPFNN	..W..FLLTNGST
229E	..TSHSVCN	GV..GHSEN	VFAVESGGYIPSN	..FAFNN	..W..FLLTNTSS
OC43	LWHLDTGVVS	CL...YKRN	.....	FTYDVNAT	..YLYFHFYQEGG
HKU1	SWHFDKSEPI	GL...EKK	.....	ETYNVSTD	..WLYFHEYQERG

	210	220	230	240	250	260
COV2	PINLV..RDL	PQGFSAL	LEP	VDP	LP	TGINITR
SARS	PIDVV..RDL	PSGFNT	LKP	IFKLP	TGINITR	FRAILT....AFSP...AQDIWGTSA
MERS	GVHLFSSRY	VDLYGGN	MFQ	FATLP	VYDT	IKYYSIIFH..SIRSIQS...DRKAW...AA
NL63	LVDGVSRRL	YQPLRLTCL	WPV	PGLKS	STGFVYFNAT	GSVNCNGYQH...NSVADV...MR
229E	VVDGVVRS	FQPLLLNCL	LWS	SGS	RFTTG	FVYFNGTGRG.DCKGFYS...NASSDV...IR
OC43	TFYAY...F	DTGTFVKF	LFNVY	LGMA	LSHYVVMFL	..TC..ISR...RDIGFT...LE
HKU1	TFYAY...Y	ADSGMPT	TF	LFSLY	LGTL	LSHYVVMFL..TCNAIS...NTDNET...LQ

	270	280	290	300	310
COV2	YVVG...Y	.....	QPR	TFLLY	NENGT
SARS	YVVG...Y	.....	QPR	TFLLY	NENGT
MERS	YVVG...Y	.....	QPR	TFLLY	NENGT
NL63	YVVG...Y	.....	QPR	TFLLY	NENGT
229E	YVVG...Y	.....	QPR	TFLLY	NENGT
OC43	YVVG...Y	.....	QPR	TFLLY	NENGT
HKU1	YVVG...Y	.....	QPR	TFLLY	NENGT



320 330 340 350 360 370  
 COV2 FRV..QPTESIV.RFPNITNLCPFGE...VFNATRFASVYANNRKRISNCVADYSVLYN  
 SARS FRV..VPSGDVV.RFPNITNLCPFGE...VFNATKFPVYAWERKKISNCVADYSVLYN  
 MERS FEA..KPSGSGVVEQAEGV..ECDFSP...L.LSGTP.PQVYNFKRLVFTNCNLYNLTLLS  
 NL63 STINTTHVSTFVGVLPTTVREIVVARTGQFYINGFKYFDLGFIEAVNFNVTASATDFWT  
 229E TTIIGNETTSAFVGLPKITVREFVISRTGHFYINGYRYFSLGNVEAVNFNVTNAA.TTVCT  
 OC43 YTV..QPIADVYRRKPD..PNCNIEA...WLNKSVSPSLNWERKTFSNCNFMNSSLMS  
 HKU1 FTV..KEVATVHRRIPD..PDIDDK...WLNFNVPSPSLNWERKIFSNCFNLSTLLR

380 390 400 410 420  
 COV2 SASFSTFKCY.GVSPTKLNDLCFNVYADSFVIRGDEVVRQIAPGQTGKTADYNYKLPDDF  
 SARS STFFSTFKCY.GVSATKLNDLCFNVYADSFVVGKDDVRQIAPGQTGKTADYNYKLPDDF  
 MERS LFSVNDFTCS.QISPAATASNCYSLLIDYFVSYP.LSMKSDLSVSSAGPIISQFNKYQSFN  
 NL63 VAFATFVDVLVNVSATKIQNLCDSPFEKLQC.EHLOFGLQDGFYSANFLDDNVLPEYI  
 229E VALASVADVLVNVSATKIQNLCDSPFEKLQC.EHLOFGLQDGFYSANFLDDNVLPEYI  
 OC43 FLQADSTTCN.NIDAAKIYGMCFSSITIDKFAIPNRRKVDLQLGNLGYLQSSNYRIDTTA  
 HKU1 LVHTDSESCN.NFDESKIYGSCEKSIIVLDKFAIPNSRRSDLQLGSSGFLQSSNYRIDTTA

430 440 450 460 470  
 COV2 TGCVTANNSNNDSKVGG...NYNYLYRLFRKSNLKF...FERDITSTEIYQAGS  
 SARS MGCVLANNTRNIDATSTG...NYNYKYRYLRHGKLRP...FERDISNVFPSPDG  
 MERS PTCILILATVPHNLTITIK...PLKYSYINKCSRLLSD...DRTEVPO..LVNAN  
 NL63 VALPIYQHTDINFNTA...TASFG...GSCYVCKP...HGVNLSL...NG  
 229E VSLPVIYHKHTFVLYV...NFELRRGPGRCYNCRP...AVVNITLANFNETK  
 OC43 TSQQLYVLPAAANVSVSRFNPSTWNNKRFIEDSVFVPOPTGVFTNHSVYVYQHCFAKPK  
 HKU1 SSSQQLYVSLPALNVITINYNPSSWNNRRYGFNNFN...LSS...HSVVYYSRYCESVNN

480 490  
 COV2 T..PCN.....GVEGFNCYFPLQ  
 SARS K..PCN.....TPPALNCYWPLN  
 MERS QYSP...VSIPTVWEDG  
 NL63 NTSVVRVTSFHSIRYIYNRVKSGSPGDSWHI...YLKSGTCFSPFS  
 229E G.PLCLVDTSHTTQFVGKFD...RWSA...SINTGNCFSPFG  
 OC43 NCFPC...SSCPGKNNGIGTCPAGINYLTCNLDL...CTLDPIIFKAPD  
 HKU1 IFCFPCAIPSPASSCKSHKPPSASCPIGTNYRSCESTTVLDHTDWCRCSCLPDPIITAYDPR

500 510  
 COV2 SY...GFQPTNFGYQPYRVVLS.....  
 SARS DY...GFYTITGIGYQPYRVVLS.....  
 MERS DYYRKQLSPLEGGGWLVASGSTVA.....MTEQLQMG  
 NL63 .....KLNNFQKFKTICFSTVAVP.....GSCNFPLEATWHYTSYITVIGAL  
 229E .....KVNNEVFKFCVCFSLKIDIP.....GGCAMPIMANLVNHHKSHNIGSL  
 OC43 TYKCPQTKSLVGIGEHCSGLAVKSDYCG.....NNSCTCQPOAFLGWSADSCLGQDKCN  
 HKU1 S...CSQKKSLLVGGEHCAGFGVDEEKCGVLDGSYNVVSLCSTDALGWSYDTCVSNRCN

520 530 540 550 560  
 COV2 .FELL.....HAPATVCGPKKSTNLV..KNKCVNFFNGLTCTGVLTESNKKFLPFQ  
 SARS .FELL.....NAPATVCGPKKSTDLI..KNKCVNFFNGLTCTGVLTESNKKFLPFQ  
 MERS .FELL.....HAPATVCGPKKSTNLV..KNKCVNFFNGLTCTGVLTESNKKFLPFQ  
 NL63 .YVTVSEGNISICVPPVSGIREFSNLV..LNNCTKYNIDYVCTGIRSSNQSLAGGIT  
 229E .YVSWSDGDVITGVPKPEVGVSSFMNVT..LNKCTKYNIDYVSGVGVIRISNDTFLNGIT  
 OC43 IFANFILHDVNNGLTCTSDTLQKANTETE..LGVCVNVDLYGISGQGFVEVNATYYNSWQ  
 HKU1 IFSNFIINGINSGTCTCSNDLLQPNTEVF..TDVGVVDYDLYGITGQGFVEVSAVYYNSWQ

570 580 590 600 610 620  
 COV2 QFGRDIADTTDAVRDPQITLEILDITFCGSGF.GVSVITPGTNTSNQVAVLYQGVNCTEVPV.  
 SARS QFGRDVSDFDTSVRDPKITEILDISPCGSGF.GVSVITPGTINASSVAVLYQGVNCTDVT.  
 MERS RFVYDAYQNLVGYYS.DDGNYYCLRACVSVFVSVIY..DKETKTHATLFGSVACEHISST  
 NL63 YV..SNSSGNLGFKKNVSTGNIFIVTFPCNFPDQVAVY...QQSIIGAMTAVNESRYGLQ.  
 229E YT..STSGNLGFKKDVNTGTYISITPCNFPDQLVVY...QQAVVGAMLSNFSTSYGFS.  
 OC43 NLLYDSNGNLGFRDYITNRTFMHSCYSGRVSAAY..HANSEFALLFRNLIKCNVFN.  
 HKU1 NLLYDSNGNIGFKDFVITNKTYNIFCYAGRVSAAF..HQNASSLALLYRNLIKCSYVFN.

630 640 650 660 670  
COV2 .AIHADQLTPTWRVYSTGSNVFQTRAGCLIGAEHVNNNS..YECDFPIGAIGICASYQTQT.  
SARS .AIHADQLTPAWRIYSTGNVVFQTOAGCLIGAEHVDTS..YECDFPIGAIGICASYHTVS.  
MERS MSQYSRSTRSMLKRRDSTYGPLQTPVGCVLGLVNSSLF.VEDCKLP LQGQSLCALPDTPTST  
NL63 ...NLLQLPNFYVVSNGG.....NNCTTAVMTYSNF.....GICADGSLIP.  
229E ...NVVEMPKFYFASNGT.....YNCTDAVLTYSNF.....GVCAADGSIIA.  
OC43 .NSLTRQLQPINYSFDSY.....LGCVVNAYNSTAISVQTCDLTVGSGYVDY.....  
HKU1 ..NISLATQPYFDSY.....LGCVFNADNLTDISVSSCALRMGSGFCVDYNSPSS

680 690 700 710 720  
COV2 NSPFRARRSVASQSIIAYTMSLGAENSVAYSNNNS.....TAIPTNFTISVTTEILFV  
SARS ...LLRSTSQKSIVAYTMSLGADSSIAYSNNNT.....TAIPTNFSISITTEVMPV  
MERS LTPRSVR.SVPGEMRLASIAFNHPIQVDQLNSS.....YFKLSIPTNFSFGVTEYIQT  
NL63 VRPRNSSSDNGISAIITAN.....LSIPSNWTTTSVQVEYLQI  
229E VQPRNVSYDSVSAIVTAN.....LSIPSNWTTTSVQVEYLQI  
OC43 SKNRRSRRAITTGTYRFTNF...EPFTVNSVNDSELPVGGLYEIQIPSEFTTIGNMEFIQT  
HKU1 SSSRRKRRSISASVRFVTF...EPFNVSVFVNDSES VGGLYEIKIPTNFTIVGQEFIT

730 740 750 760 770 780  
COV2 SMTKTSVDCIMYICGDSTECGNLLOVGSFCTQLNRATGTAVEQDKN...TOEVFAOV  
SARS SMAKTSVDCIMYICGDSTECANLLOVGSFCTQLNRALSGIAAEQDRN...TREVFAOV  
MERS TIQKVTVDCQYVVCNGFQKCEQLLREYQFC SKINQALHGANLRQDD...VRNLFAV  
NL63 TSTPIVVDCAITYVCCNGNPRCKNLLKQYTSACKTIEDALRLSAHLETN...VSSMLTFD  
229E TSTPIVVDCAITYVCCNGNPRCKNLLKQYTSACKTIEDALRNSAMLESAD...VSEMLTFD  
OC43 SSPKVTIDCAAFVCGDYAA CKLQLVEYGSFCNINAILTEVNELLDTTQLQVANS LMNGV  
HKU1 NSPKVTIDCSLFCVCSNYAACHDLLSEYSTFCNININILDEVNGLLDTTQLHVAADLMQGV

790 800 810 820 830  
COV2 KQIYKTPPTKDFG.G.FNFSQLP...DPSKPSKRSFTEDELFNKVITADAGFTKQ.YG  
SARS KQMYKTPTLKDFG.G.FNFSQLP...DPLKPTKRSFTEDELFNKVITADAGFMKQ.YG  
MERS KSSQSSPIFGF.G.DFNLTLEPVSISTGSR.SARSAIEDLFNKVITADAGFMKG.YD  
NL63 SNAFSLANVTSFG.D.YNLSVLPQRNIHSRIAAGRSALDELFNKVITSGLGTVDVDYK  
229E KKAFTLANVSSFG.D.YNLSVLPVSLPRSGSRVAGRSALDELFNKVITSGLGTVDADYK  
OC43 TLSTKLKDGYNFNVDNINFSPVLGCLGSEC SKASSRSALDELFNKVITSGLGTVDVGFVEA.YN  
HKU1 TLSSNLNTNLHFDVDNINFKSLVGCLGPHCGS.SRSRFEDELFNKVITSGLGTVDVGFVEA.YN

840 850 860 870 880 890  
COV2 DCL..GDIAARDLICQKFNCHITVLPPLLTDemiaQYTSALLAGTITSGWTFGAGAA LOI  
SARS ECL..GDINARDLICQKFNCHITVLPPLLTDDMIAAYTAAALVSGTATAGWTFGAGAA LOI  
MERS DCMQGPASARDLICQYVAGYKVLPLMDVNMEAYTSSLLGSIAGVGTAGLSSFAAI  
NL63 SCT..KGLSIADLACQYNGIMVLPGVADAERMAMYTGS LIGGMVLGGLTSA...AAI  
229E KCT..KGLSIADLACQYNGIMVLPGVADAERMAMYTGS LIGGIALGGLTSA...ASI  
OC43 NCT..GGAERDILICQYSYKCHITVLPPLLSENQISGTLAATASLFPWTAA...AGV  
HKU1 NCT..GGSEIRDILICQYSFNGIKVLPPLISESQISGYTATATVAA MFPWSSAA...AGI

900 910 920 930 940  
COV2 PFAQMMAAYRENGIGVTONVLYENQKLIANQFNKAIQDSLSSTAS.....  
SARS PFAQMMAAYRENGIGVTONVLYENQKLIANQFNKAIQESLTITST.....  
MERS PFAQSIQFYRENGVGTQVLSGENQKLIANQFNKALGAMQTGFTTNE.....  
NL63 PFSALQARQNYVALQTDVLEQENQKLIASFNKAINNIVASFSSVNDAITQTAEAIHTVT  
229E PFSALQARQNYVALQTDVLEQENQKLIASFNKAMTNIVDAFTGVNDAITQTSQALQTV  
OC43 PFYLVNQYRENGLGVTMDVLSQENQKLIANAFNNALHAIQGGFDATNS.....  
HKU1 PFSLVNQYRENGLGVTMDVLSQENQKLIATAFNNALLS LQNGFSATNS.....

950 960 970 980 990 1000  
COV2 .ALGKIQDYVNVNQAQALNTLVKQLSNFSGAISSVLNDLSRLDKVEAEVQIDRLITGRLQ  
SARS .ALGKIQDYVNVNQAQALNTLVKQLSNFSGAISSVLNDLSRLDKVEAEVQIDRLITGRLQ  
MERS .AFREKQDYVNVNNAQALS KLASELSNTFGAISASIGDIQRLDVLEQDAQIDRLINGRLT  
NL63 IALNKIQDYVNVQOQSALNHLTSQLRHNFQAISNSIQATYDRLDSIQADQVDRITGRLA  
229E TALNKIQDYVNVQOQSALNHLTSQLRHNFQAISNSIQATYDRLDSIQADQVDRITGRLA  
OC43 .ALVRIQAVNVNANAQALNLLQQLSNRFGAISASLQELSRLLDALEAEVQIDRLINGRLT  
HKU1 .ALAKIQSVNVNNAQALNLSLQQLSNRFGAISSSLOELSRLLDALEAEVQIDRLINGRLT



## Conserved epitope 2

	1010	1020	1030	1040	1050	1060
COV2	S L Q T Y V T Q Q L I R A A E I R A S A N L A A T K M S E C V L G Q S K R V D F C G K G Y H L M S F F Q S A P H G V V F					
SARS	S L Q T Y V T Q Q L I R A A E I R A S A N L A A T K M S E C V L G Q S K R V D F C G K G Y H L M S F F Q A A P H G V V F					
MERS	T L N A F V A Q Q L V R S E S A A L S A Q L A K D K V N E C V K A Q S K R S G F C G Q G T H I V S F V V N A P N G L Y F					
NL63	A L N A F V S Q V L N K Y T E V R S S R R L A Q Q K I N E C V K S Q S N R Y G F C G N G T H I F S I V N S A P D G L L F					
229E	A L N V F V S H T L T K Y T E V R A S R Q L A Q Q K V N E C V K S Q S K R Y G F C G N G T H I F S L V N A P E G L V F					
OC43	A L N A Y V S Q Q L S D S T L V K F S A A Q A M E K V N E C V K S Q S S R I N F C G N G N H I I S L V Q N A P Y G L Y F					
HKU1	A L N A Y V S Q Q L S D I S L V K F G A A L A M E K V N E C V K S Q S P R I N F C G N G N H I L S L V Q N A P Y G L L F					

	1070	1080	1090	1100	1110
COV2	L H V T Y V P A Q E K N F T T A P A I C H D G . . . K A H F P R E G V F V S N G T H . . . W F V T Q R N F Y E P Q I				
SARS	L H V T Y V P S Q E R N F T T A P A I C H E G . . . K A Y F P R E G V F V F N G T S . . . W F I T Q R N F F S P Q I				
MERS	M H V G Y Y P S N H I E V S A Y G L C D A A N P T N C I A P V N G Y F I K T N N T R I V D E W S Y T G S S F Y A P E P				
NL63	L H T V L L P T D Y K N V K A W S G I C V D G I . . Y G Y V L R Q P N L V L Y S D N G V . . . F R V T S R V M F Q P R L				
229E	L H T V L L P T Q Y K D V E A W S G L C V D G I . . N G Y V L R Q P N L A L Y K E G N Y . . . Y R I T S R I M F E P R I				
OC43	I H F N Y V P T K Y V T A K V S P G L C I A G N . . R G I A P K S G Y F V N V N N T . . . W M Y T G S G Y Y P E P				
HKU1	M H F S Y K P I S F K T V L V S P G L C I S G D . . V G I A P K Q G Y F I K H N D H . . . W M F T G S S Y Y P E P				

	1120	1130	1140	1150	1160	1170
COV2	I T T D N T F V S G N C D V V I G I V N N T V Y D P L Q P E L . . D S F K E E L D K Y F K N H T S . P D V D L G . D I S					
SARS	I T T D N T F V S G N C D V V I G I I N N T V Y D P L Q P E L . . D S F K E E L D K Y F K N H T S . P D V D L G . D I S					
MERS	I T S L N T K Y V A P . Q V T Y Q N I S T N L P P P L L G N S T G I D F Q D E L D E F F K N V S T . S I P N F G . S L T					
NL63	P V L S D F V Q I Y N C N V T F V N I S R V E L H T V I P D Y . . V D V N K T L Q E F A Q N L P K Y V K P N F . . D L T					
229E	P T I A D F V Q I E N C N V T F V N I S R S E L Q T I V P E Y . . I D V N K T L Q E L S Y K L P N Y T V P D L . . V V E					
OC43	Y I N V T F L D L Q V E M N . . . T C A V N Y T K A P Y V M L N T S I P N L . . P D F K E E L D Q W F K N Q T S . V A P D L . . S L D					
HKU1	I S D K N V V F M N T C S V N F T K A P L V Y L N H S V P K L . . S D F E S E L S H W F K N Q T S . I A P N L T L N L H					

## Conserved epitope 1

	1180	1190	1200	1210
COV2	G I N A S V V N I Q K E I D . . . . . R L N E V A K N L N E S L I D L Q E L G K Y E C Y I K W P W Y I			
SARS	G I N A S V V N I Q K E I D . . . . . R L N E V A K N L N E S L I D L Q E L G K Y E C Y I K W P W Y V			
MERS	Q I N T L L D L T Y E M L . . . . . S L Q Q V V K A L N E S Y I D L K E L G N Y T Y Y N K W P W Y I			
NL63	P E N L T Y L N L S S E L K Q L E A K T A S L F Q T T V E L Q G L I D Q I N S T Y V D L K L L N R F E N Y I K W P W V			
229E	Q Y N Q T I L N L T S E I S T L E N K S A E L N Y T V Q K L Q T L I D N I N S T L V D L K W L N R V E T Y I K W P W V			
OC43	Y I N V T F L D L Q V E M N . . . . . R L Q E A I K V L N H S Y I N L K D I G T Y E Y Y V K W P W Y V			
HKU1	T I N A T F L D L Y Y E M N . . . . . L I Q E S I K S L N S Y I N L K D I G T Y E M Y V K W P W Y V			

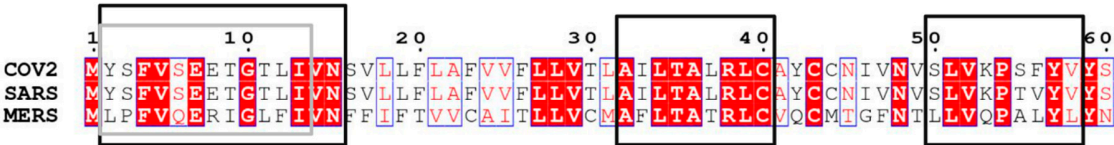
	1220	1230	1240	1250	1260	1270
COV2	W L G F I A G L I A I V M V T I M L C C M T S C S C L K G . C C S C G S C C K F D E D D S E P V L K G V K I H Y T					
SARS	W L G F I A G L I A I V M V T I L L C C M T S C S C L K G . A C S C G S C C K F D E D D S E P V L K G V K I H Y T					
MERS	W L G F I A G L V A L A L C V F F I L C T G C G T N C M G . K L K C N R C C D . R Y E E Y . . D L E P H K V H V H					
NL63	W L I I S V V F V V L L S L L V F C C L S T G C C G C C N C L T S S M R G C C D C G S T K L . P Y Y E F E K V H V Q					
229E	W L C I S V V L I F V V S M L L L C C S T G C G G F F S C F A S S I R G C C E . . S T K L . P Y Y D V E K I H I Q					
OC43	W L I C L A G V A M L V L L F F I C C T G C G T S C F K . . K C G G C C D . D Y T G Y . Q E L V I K T S H D D					
HKU1	W L I S E S F I I F L V L L F F I C C T G C G S A C F S . . . K C H N C C D . E Y G G H . H D F V I K T S H D D					

**Supplementary figure S3. Conservation of envelope, membrane, nucleocapsid and spike protein sequences in seven HCoV.** Consensus sequences of each protein were aligned and conserved regions were identified using ESPript 3.x. Conserved amino acids between HCoVs are colored in red. A. For envelope protein, no conserved epitopes were found for envelope protein among seven HCoVs. B. For membrane protein, 1 CD8<sup>+</sup> T cell epitope (Conserved epitope 1) was found to be conserved among seven HCoVs with ~55% conservancy, which is shown in the black square. C. For nucleocapsid protein, 1 CD8<sup>+</sup> T cell epitope (Conserved epitope 1) was found to be conserved among seven HCoVs with ~44% conservancy, which is shown in the black square. D. For spike protein, 2 CD8<sup>+</sup> T cell epitopes (Conserved epitope 1 and Conserved epitope 2) were found to be conserved among seven HCoVs, which are shown in the black squares. Epitope 1 is ~66% conserved, while epitope 2 is ~44% conserved. Epitopes are numbered with order of the highest conservancy.



A. Conservation of Envelope Protein Sequence in SARS-CoV-2, SARS-CoV, MERS-CoV

Conserved epitope 1  
(B cell epitope)



Conserved epitope 2  
(B cell epitope)

Conserved epitope 1      Conserved epitope 2



## B. Conservation of Membrane Protein Sequence in SARS-COV-2, SARS-COV, MERS

	1	10	20	30	40	50	60																																																					
COV2	M	A	D	S	N	G	T	I	T	V	E	E	L	K	K	L	L	E	Q	W	N	L	V	I	G	F	I	F	L	T	W	I	C	L	L	Q	F	A	Y	A	N	R	N	R	F	I	Y	I	I	K	L	I	F	L	W	L	L	W	P	V
SARS	M	A	D	.	N	G	T	I	T	V	E	E	L	K	Q	L	L	E	Q	W	N	L	V	I	G	F	I	F	L	A	W	I	M	L	L	Q	F	A	Y	S	N	R	N	R	F	I	Y	I	I	K	L	V	F	L	W	L	L	W	P	V
MERS	M	S	N	.	M	T	Q	I	T	E	A	Q	I	I	A	I	I	K	D	W	N	F	A	W	S	L	I	F	L	I	T	I	V	L	Q	Y	G	Y	P	S	R	S	M	T	V	Y	V	F	K	M	F	V	L	W	L	L	W	P	S	

	70	80	90	100	110	120																																																				
COV2	T	L	A	C	F	V	L	A	A	V	R	I	N	W	I	T	G	G	I	A	I	A	M	A	C	I	V	G	L	M	W	S	Y	F	I	A	S	F	R	L	F	A	R	T	E	S	M	W	S	F	N	P	E	T	N	I	L	L
SARS	T	L	A	C	F	V	L	A	A	V	R	I	N	W	V	T	G	G	I	A	I	A	M	A	C	I	V	G	L	M	W	S	Y	F	V	A	S	F	R	L	F	A	R	T	E	S	M	W	S	F	N	P	E	T	N	I	L	L
MERS	S	M	A	L	S	I	F	S	A	V	P	I	D	L	A	S	Q	I	I	S	G	I	V	A	A	V	S	A	M	M	W	S	Y	F	V	Q	S	I	R	L	F	M	R	T	E	S	W	S	F	N	P	E	T	N	C	L	L	

Conserved epitope 2      Conserved epitope 1

Conserved epitope 1 (B cell epitope)

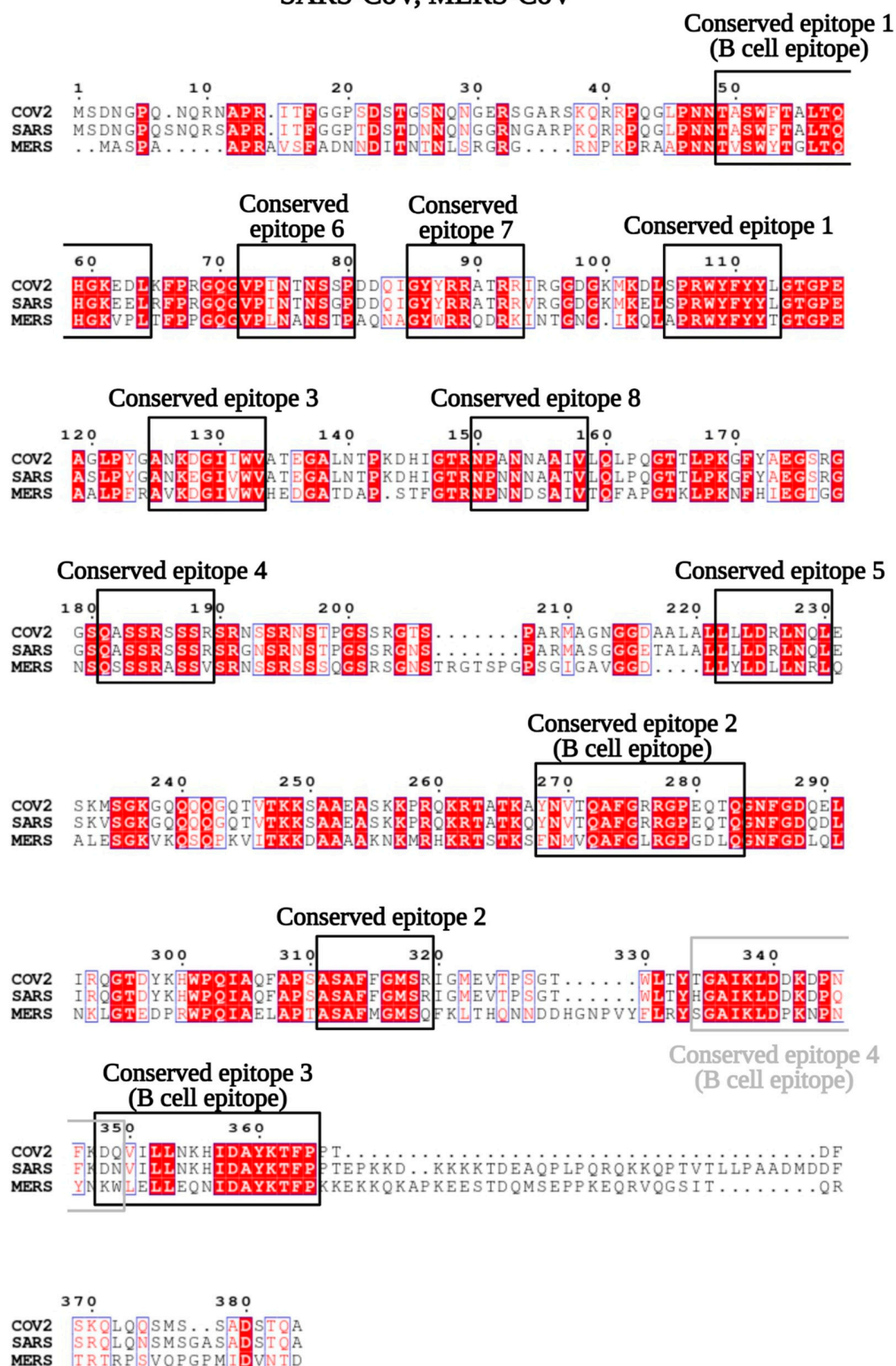
	130	140	150	160	170	180																																																						
COV2	N	V	P	L	H	G	T	I	L	T	R	P	L	L	E	S	E	L	V	I	G	A	V	I	L	R	G	H	L	R	I	A	G	H	H	L	G	R	C	D	I	K	D	L	P	K	E	I	T	V	A	T	S	R	T	L	S	Y	Y	K
SARS	N	V	P	L	R	G	T	I	V	T	R	P	L	M	E	S	E	L	V	I	G	A	V	I	I	R	G	H	L	R	M	A	G	H	S	L	G	R	C	D	I	K	D	L	P	K	E	I	T	V	A	T	S	R	T	L	S	Y	Y	K
MERS	N	V	P	F	G	T	V	V	R	P	L	V	E	D	S	T	S	V	T	A	V	V	T	N	G	H	L	K	M	A	G	M	H	F	G	A	C	D	I	D	R	L	E	N	E	V	T	V	A	K	P	N	V	L	I	A	L	K		

Conserved epitope 3

	190	200	210	220																																						
COV2	L	G	A	S	Q	R	V	A	G	D	S	G	F	A	A	S	R	Y	R	I	G	N	Y	K	L	N	T	D	H	S	S	S	S	D	N	I	A	L	L	V	Q	
SARS	L	G	A	S	Q	R	V	G	T	D	S	G	F	A	A	S	N	R	Y	R	I	G	N	Y	K	L	N	T	D	H	A	G	S	N	D	N	I	A	L	L	V	Q
MERS	M	V	K	R	Q	S	G	T	N	S	G	V	A	I	Y	H	R	Y	K	A	G	N	Y	R	.	S	P	P	I	T	A	D	I	E	L	A	L	L	R	A		

Conserved epitope 4

### C. Conservation of Nucleocapsid Protein Sequence in SARS-CoV-2, SARS-CoV, MERS-CoV



# D. Conservation of Spike Protein Sequence in SARS-COV-2, SARS-COV, MERS

	1	10	20	30	
COV2	..MFVFLVLLP	VSSQ.....	CVNLTTRTQL...	PAYTN..	STRGVYYP
SARS	..MFIFLLFLT	LTSGS.....	DLDRCTTFDDVQAP	PNYTQHT	SSMRGVYYP
MERS	MIHSVFLIMFL	LTPTESYVDVG	FDSVKSACIEVD	IQQTFFDKTW	PRPID.VSKADGIIYP

	40	50	60	70	80	90
COV2	DKVFRSVLHST	QDLFLPF...	FSNVTFHAIHV	SGINGTKR	....	DNFLVLPFNDGV..
SARS	DEIFRSDTLYLT	QDLFLPF...	YSNVTFHTINH	.....	....	GNFVIFPKDGI..
MERS	QGRITYSNITITY	QGLFLP...	HYQGDHGD	MYVYSAGHA	TGTTPOKL	EVANYSDVKQFANGFVV

	100	110	120			
COV2	.....YFASTEK	SNIRG...	WIFGTTLD	SKT.....	QLLVNNATNV	
SARS	.....YFAATEK	SNVVRG...	WVFGSTMNN	KS.....	QSVIIINNSTNV	
MERS	RIGAAANSTGT	VLISFST	SATLRKIYPA	FMLGSSVGN	FDGKMGRFFNH	TLVLLPDGCGT

	130	140	150	160	
COV2	VIVV.....	CEQFCND	PFLGVYHKN	KSWMSEFRV	TSANCT
SARS	VIRA.....	CNELCDN	PFFAVSKPMG	TQTHM...	IFDNAFNCT
MERS	LLRAYFYCILEPR	SGNHCFAGNSYTS	ATYHTPATDCSD	GNYNRNASLNSFKE	YFNINCT

	170	180	190	200	210	220
COV2	FEY...VSQFFL	MDLEGK	CNFKNLR	FEVFKNIDG	VFKHYSKHTPIN	VVRDLPQGSAL
SARS	FEY...ISDAFS	LDVSEK	SNFKHLR	FEVFKNKDGF	LYYKGYQPID	VVRDLPSGFNTK
MERS	FMYTYNIT	TEDEILEWFGI	HTAQGVHL	FSSR...	YVDLYGG...	NMFQ....SATL

	230	240	250	260	270	280
COV2	PVLDLPIGINIT	RQTLLALHRSYL	TPGDSSSC	WTAGAAVY	GYVOPRFF	LLKTYNENCT
SARS	PVFKLPLGINIT	FRAIL...TAF	SFAQDI..	WGTSAAVY	GYVOPRFF	LLKYDENCT
MERS	PVYD.....	TIKYYSILPHSIRS	ISDRKA...	W....AAEYVYK	OPRFFLLDE	SVDCY

	290	300	310	320	330	340
COV2	ITDAVDCALDP	LSETCTLM	SFTVERKGY	YVTSMFRVQPT	ESIVRFPNITNL	CFGGEVFNA
SARS	ITDAVDCSQNP	LAELKCSV	KSFEDIKGY	YVTSMFRVVP	SGDVVRFPNITNL	CFGGEVFNA
MERS	ITRAIDCGFND	LSQLKCSYB	SFDVESGUY	SVSEFAKPS	SGSVVEQAEGVE	.CQFSPLLSG

	350	360	370	380	390	400
COV2	TRFAVSVVAM	NRRKISNC	VADYSVLYN	SASFSTKCYG	VSPTKNDICFTNV	YADSIVIRG
SARS	TKFPVSVVAM	NRKKISNC	VADYSVLYN	STFFSTKCYG	VSAIKNDICFSNV	YADSIVVKG
MERS	TP.PGVVNF	KRLVFTNC	NYNLRLSLFSV	NDTCSQISPAAL	ASNAYSLLILDY	ESYPL

	410	420	430	440	450	460
COV2	DEVROAPGT	CTKLTADV	NYKLPDDFTG	CVIAWNN	SNLDSKVG	GNYNVLYRLFKSNLKPF
SARS	DDVROAPGT	CTKLTADV	NYKLPDDFTG	CVIAWNN	TRNIDAT	STGNYNVLYRLFKSNLKPF
MERS	SMKSDLVSV	SAGPISQ	FNYKQSF	SNFTCLILA	TVPHNLT	TITKPLKYSYINICSRLLSD

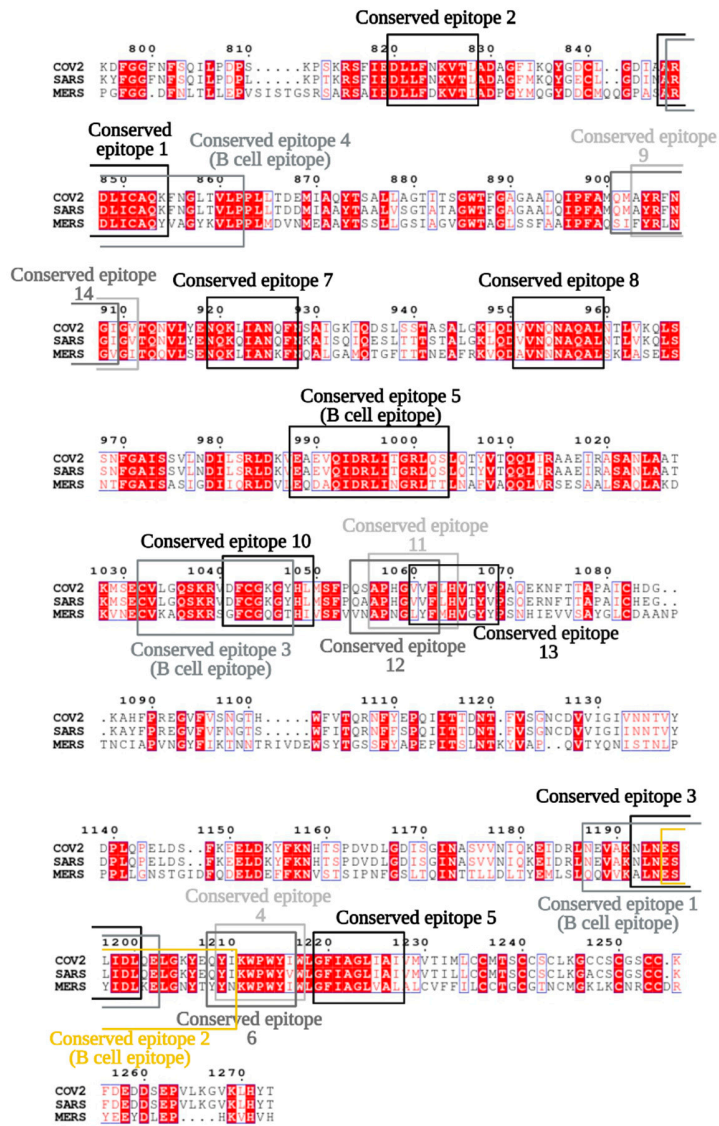
  

	470	480	490	500	510	
COV2	ERDISTE	EIQAGST	PCNGVEGF	NCV.....	FLQSYG	CFQPTNGVGYQPYRVVVL
SARS	ERDISTE	VFPSPDGK	PCP.PALN	CV.....	WPLNDY	CFYTTTCIGYQPYRVVVL
MERS	DRTEVP	GLVNANQYS	PCVSIVPS	TVMEDGDY	YRKQLSPLEGG	MLVASGSTVAMTEQLQM

	520	530	540	550	560	
COV2	SPELL...HAPA	TVC.....	GPKKSTN	LVKNKCVN	FNFNGLT	GTGVLTESNKKFLPF
SARS	SPELL...HAPA	TVC.....	GPKLSTN	LVKNKCVN	FNFNGLT	GTGVLTESNKKFLPF
MERS	GGGITVQYGTDTN	SVCPKLEFANDTKIASQL	..GNCV	EYSLYGVSGRVFQNCITAVGVRRQ		

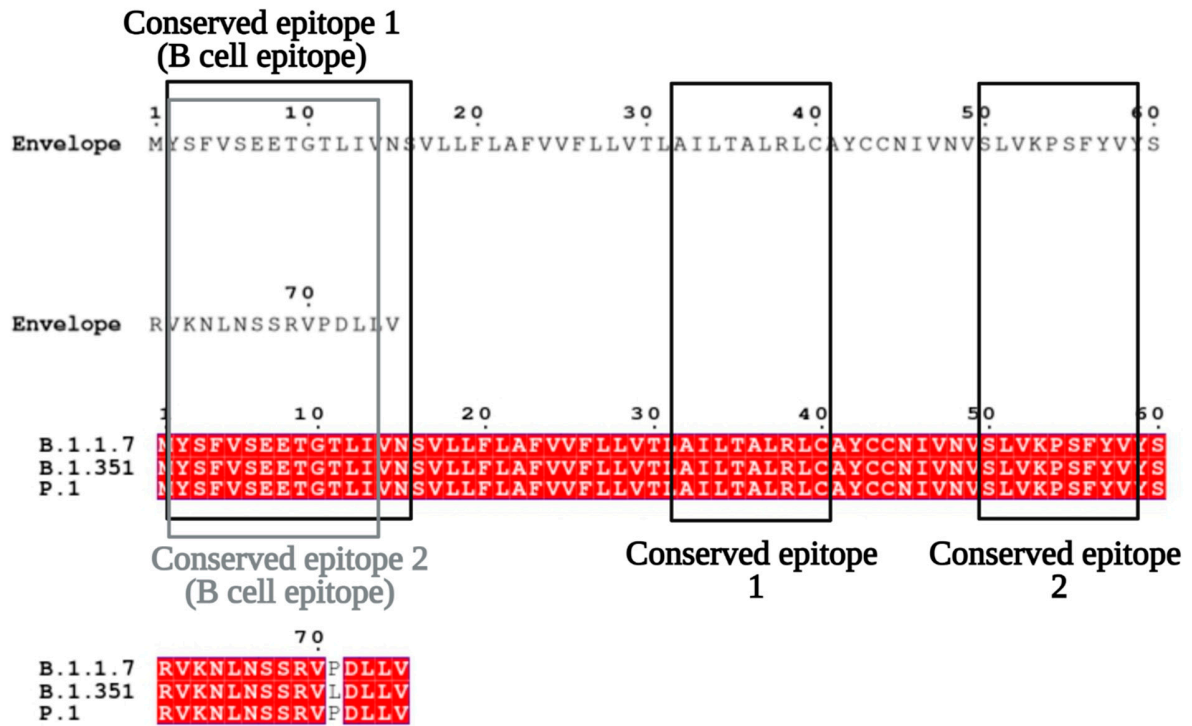




**Supplementary figure S4. Conservation of envelope, membrane, nucleocapsid, and spike protein sequences in SARS-CoV-2, SARS-CoV and MERS-CoV.**

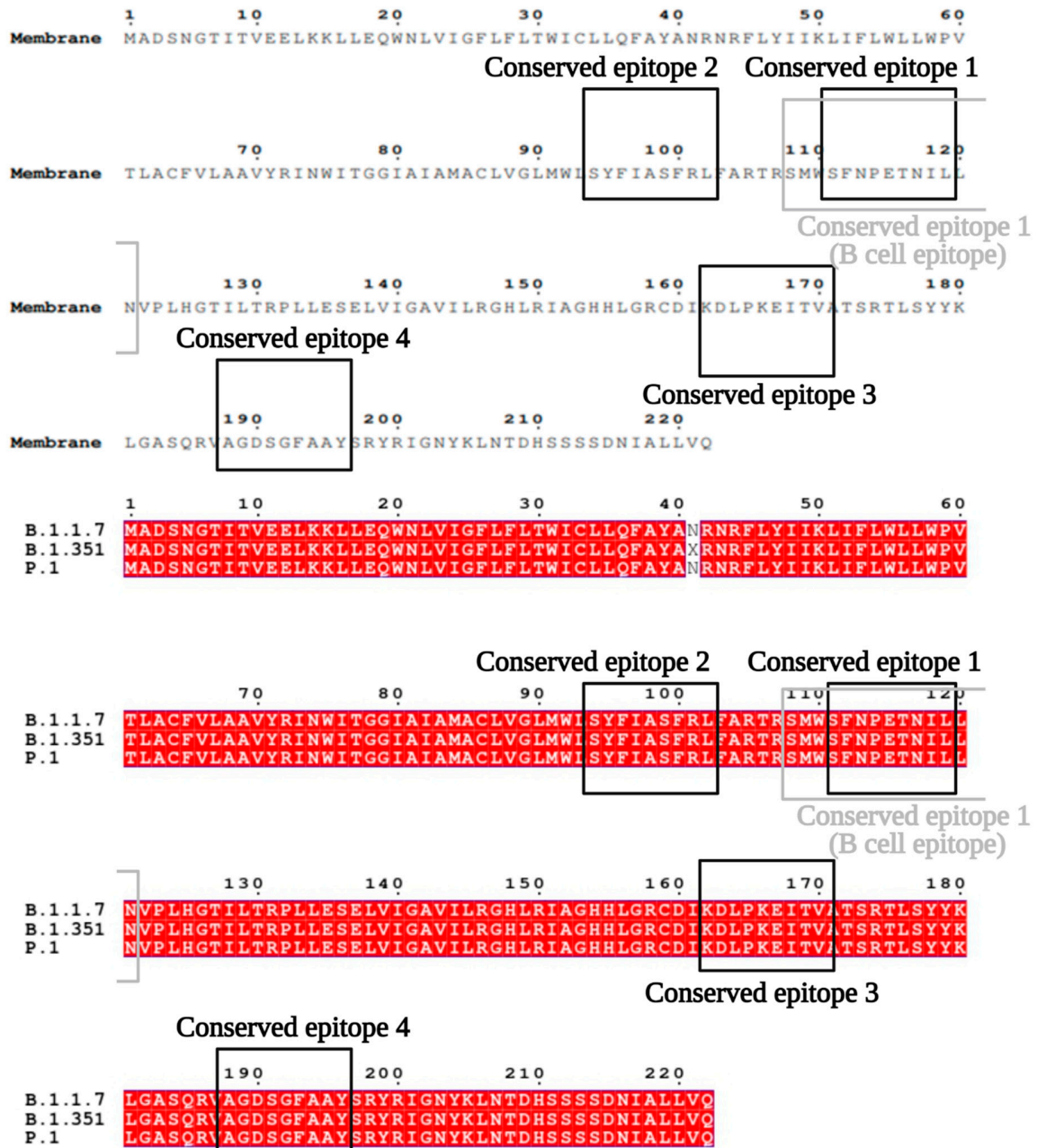
Consensus sequences of each protein were aligned and conserved regions were identified using ESPript 3.x. Conserved amino acids are colored in red. A. For envelope protein, two CD8<sup>+</sup> T cell epitopes 1 and 2, and two B cell epitopes 1 and 2, were found to be conserved among SARS-CoV-2, SARS-CoV and MERS-CoV, which are shown in black and grey squares. Two CD8<sup>+</sup> T cell epitopes, 1 and 2, were ~77% and ~44% conserved, respectively, while two B cell epitopes 1 and 2 were, respectively, ~50% and ~42% conserved. B. For membrane protein, four CD8<sup>+</sup> T cell epitopes 1, 2, 3 and 4, and one B cell epitope, were found to be conserved among SARS-CoV-2, SARS-CoV and MERS-CoV, which are shown in black and grey squares. Four CD8<sup>+</sup> T cell epitopes, 1, 2, 3, and 4 were, respectively, ~88%, ~66%, ~66% and ~44% conserved, while one B cell epitope was ~86% conserved. C. For nucleocapsid protein, eight CD8<sup>+</sup> T cell epitopes 1, 2, 3, 4, 5, 6, 7, and 8, and four B cell epitopes 1, 2, 3, and 4, were found to be conserved among SARS-CoV-2, SARS-CoV and MERS-CoV, which are shown in black and grey squares. Eight CD8<sup>+</sup> T cell epitopes 1, 2, 3, 4, 5, 6, 7, and 8 – were, respectively, ~77%, ~77%, ~66%, ~66%, ~66%, ~66%, ~55% and ~55% conserved, while four B cell epitope 1, 2, 3, and 4, were, respectively, ~69%, ~56%, ~56% and ~50% conserved. D. For spike protein, 14 CD8<sup>+</sup> T cell epitopes, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, and 14, and five B cell epitopes, 1, 2, 3, 4, and 5, were found to be conserved among SARS-CoV-2, SARS-CoV and MERS-CoV, which are shown in, respectively, black, grey, and orange squares. Fourteen CD8<sup>+</sup> T cell epitopes - 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, and 14 – were ~88%, ~77%, ~77%, ~77%, ~77%, ~66%, ~66%, ~66%, ~55%, ~55%, ~55%, ~44%, ~44%, and ~44% conserved, respectively, while five B cell epitope - Conserved epitopes 1, 2, 3, 4, and 5 - are ~63%, ~63%, ~56%, ~50%, and ~50% conserved. Epitopes are numbered in order of the highest conservancy.

**A.** Conservation of Envelope Protein Sequence  
in 5000 SARS-CoV-2 sequences and in B.1.1.7, B.1.351, P.1 variants

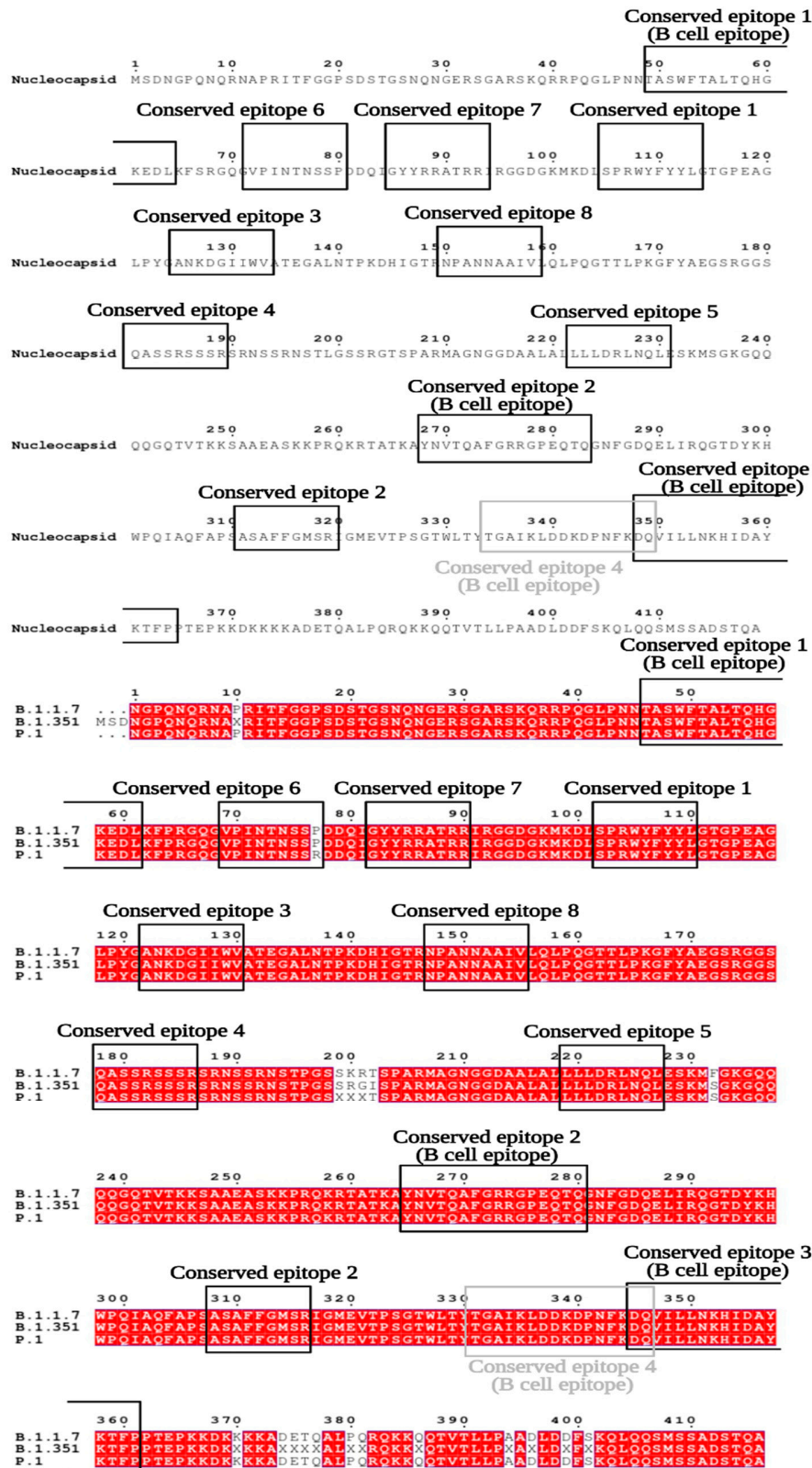




**B. Conservation of Membrane Protein Sequence  
in 5000 SARS-CoV-2 sequences and in B.1.1.7, B.1.351, P.1 variants**

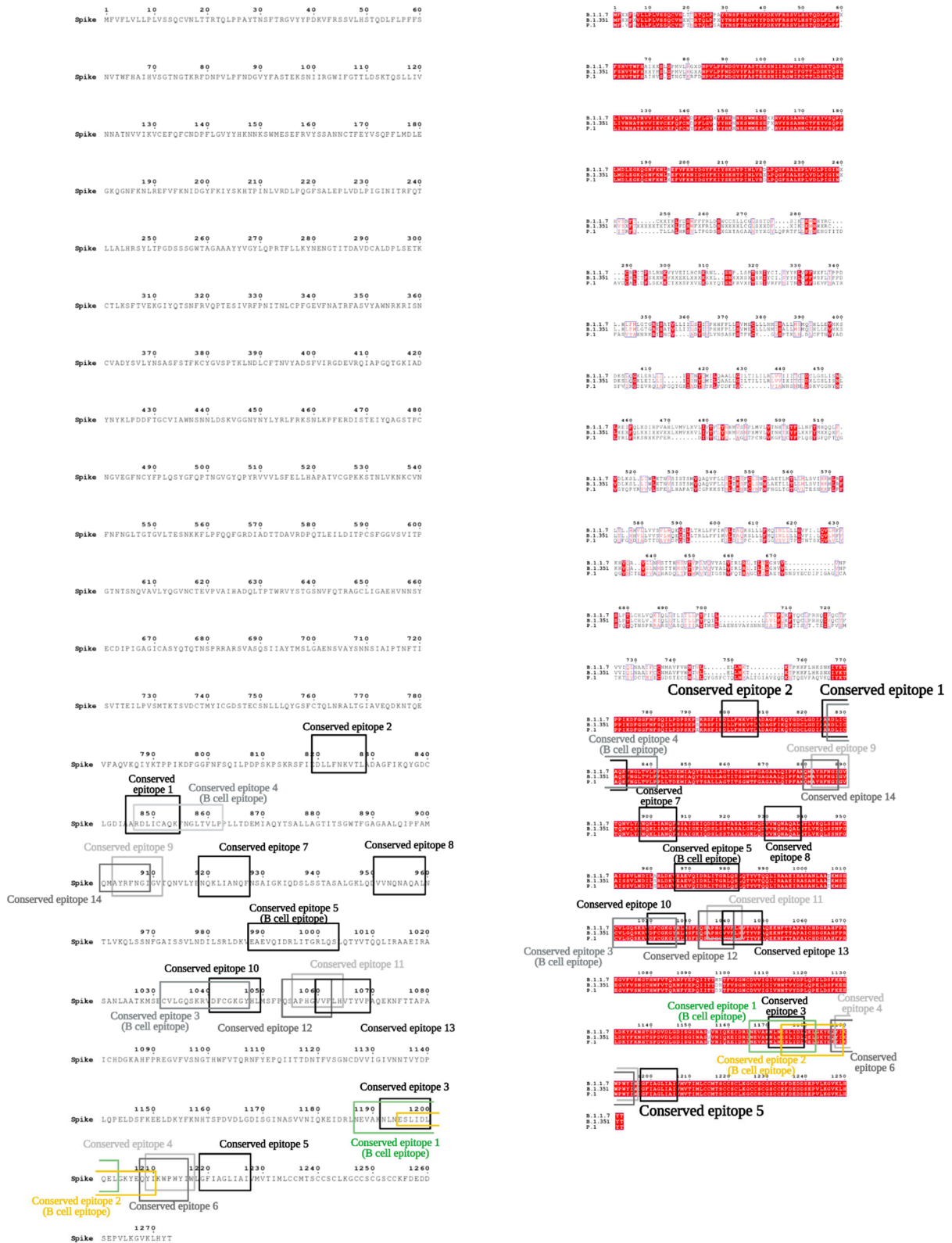


**C.**



D.

# Conservation of Spike Protein Sequence in 5000 SARS-CoV-2 sequences and in B.1.1.7, B.1.351, P.1 variants



**Supplementary figure S5. Conservation of envelope, membrane, nucleocapsid, and spike protein sequence in 5000 SARS-CoV-2 sequences and in B.1.1.7, B.1.351, and P.1 variants of SARS-CoV-2.** Consensus sequences of envelope protein were aligned and conserved regions were identified using ESPript 3.x. CD8+ T and B cell epitopes of SARS-CoV-2 predicted to show conserved amino acids in HCoV-229E were also found to be 100% conserved in different variants of SARS-CoV-2.

**Supplementary Table S1. Consensus sequences used to predict the epitopes for vaccine design.** Viral protein sequences for Envelop, Membrane, Nucleocapsid and Spike proteins were retrieved using National Center for Biotechnology Information (NCBI) reference sequences for all seven HCoV-229E: SARS-CoV-2 (NC\_045512), SARS-CoV (NC\_004718), MERS-CoV (NC\_019843), HCoV-NL63 (NC\_005831), HCoV-229E (NC\_002645), HCoV-HKU1 (NC\_006577), and HCoV-OC43 (NC\_006213). The protein sequences were then aligned using online MAFFT version 7 (<https://mafft.cbrc.jp/alignment/server/>) and consensus sequences were generated with Consensus Maker tool (<https://www.hiv.lanl.gov/content/sequence/CONSENSUS/SimpCon.html>)

Name of the virus	Consensus sequence
<b>Envelope</b>	
SARS-CoV-2	MYSFVSEETGTLIVNSVLLFLAFVVFLVTLAILTALRLCAYCCNIVNVSLVK PSFYVYSRVKNLNSSRVPDLLV
SARS-CoV	MYSFVSEETGTLIVNSVLLFLAFVVFLVTLAILTALRLCAYCCNIVNVSLVK PTVYVYSRVKNLNSSEGVPDLLV
MERS-CoV	MLPFVQERIGLFIVNFFITVVCATLLVCMAFLTATRLCVQCMTGFNTLLVQ PALYLYNTGRSVYVKFQDSKPPLPPDEWV
HCoV-NL63	MFLRLIDDNGIVLNSILWLLVMIFFFVLAMTFIKLIQLCFTCHYFFSRTLQPVY KIFLAYQDYMQUIAPVPAEVLNV
HCoV-229E	MFLKLVDHDLVNVNLLWCVVLIIVILLVCITIIKLIKLCFTCHMFCNRTVYGP IKNVYHIYQSYMHIDPFPPKRVIDF

HCoV-OC43	MFMA DAYLADTVWYVGQIIFIVAICLLVTIVVVAFLATFKLCIQICGMCNTL VLSPSIYVFNRGRQFYEFYNDIKPPVLDVDDV
HCoV-HKU1	MVDVFFTDTAWYIGQIFFLVLSCVIFLIFVVALLATIKLCIQICGFCNIFIISPA YVYNRGRQLYKSYSEHVIPSTLDDLI
<b>Membrane</b>	
SARS-CoV-2	MADSNGTITVEELKKLLEQWNLVIGFLFTWICLLQFAYANRNRFLYIIKLIF LWLLWPVTLACFVLA AVYRINWITGGIAIAMACLVGLMWLSYFIASFRLFAR TRSMWSFNPETNILLNVPLHGTILTRPLLESELVIGAVILRGHLRIAGHHLGR- CDIKDLPKEITVATSRTL SYYKLGASQRVAGDSGFAAYSRYRIGNYKLNTDH SSSDNIALLVQ
SARS-CoV	MADNGTITVEELKQLLEQWNLVIGFLFLAWIMLLQFAYSNNRNRFLYIIKLVF LWLLWPVTLACFVLA AVYRINWVTGGIAIAMACIVGLMWLSYFVASFRLFAR RTRSMWSFNPETNILLNVPLRG TIVTRPLMESELVIGAVIIRGHLRMAGHSLG RCDIKDLPKEITVATSRTL SYYKLGASQRVGTDSGFAAYNRYRIGNYKLNTD HAGSNDNIALLVQ
MERS-CoV	MSNMTQLTEAQIIAIKDWNFAWSLIFLLITIVLQYGYP SRSM TVYVFKMFVL WLLWPSSMALSI FSAVYPIDLASQIISGIVAAVSAMMWISYFVQSIRLFMRTG SWWSFNPETNCLLNVPFGGTTVVRPLVEDSTSVTAVVTNGHLKMAGMHFG ACDYDRLPNEVTVA KPNVLIALKMVKRQSYGTNSGVAIYHRYKAGNYRSPP ITADIELALLRA
HCoV-NL63	MSNSSVPLSEVYVHLRNWNFSWNLILTVFIVVLQYGHYKYSRLLYGLKMSV LWCLWPLVLALSIFDCFVNFNVDWVFFGSILMSIITLCLWVMYFVNSFRLW RRVKTFWAFNPETNAIISLQVYGHNYL PVM AAPTGVTLTLLSGVLLVDGH KIATRVQVGQLPKYVIVATPSTTIVCDRVGRSVNETSQTGWAFYVRAKHGD FSGVASQEGVLSEREKLLHLI

HCoV-229E	MSNDNCTGDIVTHLKNWNFGWNVILTIFIVILQFGHYKYSRLFYGLKMLVL WLLWPLVLALSIFDTWANWDSNWAFAFSFLMAVSTLVMWVMYFANSFR LFRRARTFWAWNPEVNAITVTTVLGQTYQPIQQAPTGITVTLISGVLVVDG HRLASGVQVHNLPEYMTVAVPSTTIYSRVGRSVNSQNSTGWVFYVRVKHG DFSAVSSPMSNMTENERLLHFF
HCoV-OC43	MSSKTTTPAPVYIWTADEAIKFLKEWNFSLGIILLFITIILQFGYTSRSMFVYVIK MIILWLMWPLTIILTIFNCVYALNNVYLGLSIVFTIVAIIMWIVYFVNSIRLFIR TGSFWSFNPETNNLMCIDMKGTMYVRPIIEDYHTLTVTIIRGHLIYQGIKLT GYSLADLPAYMTVAKVTHLCTYKRGFLDRISDTSGFAVYVKSKVGNYRLPS TQKGSGMDTALLRNNI
HCoV-HKU1	MNKSFFPQFTSDQATFLKEWNFSLGVILLFITIILQFGYTSRSMFVYLIKMIIL WLMWPLTITLTIFNCFYALNNAFLAFSIVFTIISIVIWILYFVNSIRLFIRTGSW WSFNPETNNLMCIDMKGKMFVRPVIEDYHTLTATVIRGHLIYQGVKLTGTGY TSLDLPVYVTVAKVQVLCTYKRAFLDKLDVNSGFAVFVKSKVGNYRLPSSK PSGMDTALLRA
<b>Nucleocapsid</b>	
SARS-CoV-2	MSDNGPQNQRNAPRITFGGPSDSTGSNQNGERSGARSKQRRPQGLPNNTAS WFTALTQHGKE- DLKFPRGQGVPIINTNSSPDDQIGYYRRATRRIRGGDGKMKDLSRWYFYYL GTGPEAGLPYGANKDGIIWVATEGALNTPKDHIGTRNPANNAIIVLQLPQGT TLPKGFYAEGSRGGSQASSRSSSRNSSRNSTPGSSRGTSARMAGNGGDA ALALLLDRLNQLESKMSGKGQQQQGQTVTKKSAAEASKKPRQKRTATKA- YNVTQAFGRRGPEQTQGNFGDQELIRQGTDYKHWPQIAQFAPSASAFFGMS RIGMEVTPSGTWLTYTGAIKLDDKDPNFKDQVILLNKHIDAYKTFPPTDFSK QLQQSMSSADSTQA
SARS-CoV	MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTA SWFTALTQHGKEELRFPRGQGVPIINTNSGPDDQIGYYRRATRRVRGGDGKM KELSPRWYFYYLGTGPEASLPYGANKEGIVWVATEGALNTPKDHIGTRNPN NNAATVLQLPQGTTLPKGFYAEGSRGGSQASSRSSSRGNSRNSTPGSSRG NSPARMASGGGETALALLLDRLNQLESKVSQKGGQQQQGQTVTKKSAAEA



	SKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYKHWPQI AQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKH IDAYKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQL QNSMSGASADSTQA
MERS-CoV	MASPAAPRAVSFADNNDITNTNLSRGRGRNPKPRAAPNNTVSWYTGLTQHG KVPLTFPPGQGVPLNANSTPAQNAGYWRRQDRKINTGNGIKQLAPRWYFYF TGTGPEAALPFRAVKDGVVWHEDGATDAPSTFGTRNPNNSAIVTQFAPGT KLKPNFHIEGTGGNSQSSSRASSVSRNSSRSSSQGSRSGNSTRGTSPGPSGIGA VGGDLLYDLLNRLQALESGKVQSQPKVITKDDAAAKNKMHRHKRTSTK SFNMVQAFGLRGPDLQGNFGDLQLNKLGTEDPRWPQIAELAPTASAFMG MSQFKLTHQNNDDHGNPVYFLRYSGAIKLDPKNPYNKWLELLEQNIDAY KTFPKKEKKQKAPKEESTDQMSEPPKEQRVQGSITQRTRTRPSVQPGPMIDV NTD
HCoV-NL63	MASVNWADDRAARKKFPPPSFYMPLLVSDDKAPYRVIPRNLVPIGKGNKDE QIGYWNVQERWRMRRGQRVDLPPKVHFYYLGTGPHKDLKFRQRSDGVVW VAKEGAKTVNTSLGNRKRNPQPLEPKFSIALPPELSVVEFEDRSNNSSRASSR SSTRNNSRDSSRSTSRQQRTRSDSNQSSSDLVAAVTLALKNLGFDNQSKSPS SSGTSTPKKPNKPLSQPRADKPSQLKKPRWKRVPTREENVIQCFGPRDFNHN MGDSDLVQNGVDAKGFPQLAELIPNQAALFFDSEVSTDEVGDNVQITYTYK MLVAKDNKNLPKFIEQISAF TKPSSIKEMQSQSSHVVQNTVLNASIPESKPLA DDDSAIIEIVNEVLH
HCoV-229E	MATVKWADASEPQRGRQGRIPYSLYSPLLVDSEQ-PWKVIPRNLVPINKK- DKNKLIGYWNVQKRFRTRKGRVDLSPKLHFYYLGTGPHKDAKFRERVEG VVWVAVDGAKTEPTGYGVRRKNSEPEIPHFNQKL PNGVTVVEEPDSRAPSR SQRSQSRSRGESKSQSRNPSSDRNHNSQDDIMKAVAAALKSLGFDKPQEKD KKSAGTGTPKPSRNQSPASSQSAKILARSQSSETKEQKHEMQKPRWKRQP NDDVTSNVTQCFGPRDL DHNFGSAGVVANGVKAKGYPQFAELVPSTAAML FD SHIVSKESGNTVVLTFTRVTVPKDHPLGKFLEELNAFTREMQQQPLL N PSALEFNPSQTSPATVEPVRDEVSIETDIIDEVN
HCoV-OC43	MSFTPGKQSSSRASSGNRSGNGILKWADQSDQFRNVQTRGRRAQPKQTATS QQPSGGNVVPYYSWFSGITQFQKGKEFEFAEGQGVPIAPGVPATEAKGYWY RHNRRSFKTADGNQRQLLPRWYFYLLGTGPHAKDQYGTIDGVYVWASN QADVNTPADIVDRDPSSDEAIPTRFPPGTVLPQGYIIEGSGRSAPNSRSTSRTS SRASSAGSRSRANSNRTPTSGVTPDMADQIASLV LAKLGKDATKPPQVTK HTAKEVRQKILNKPRQKRSPNKQCTVQQCFGKRGPNQNFGGGEMLKLGTS

	DPQFPILAEAPTAGAFFFGSRLELAKVQNLSGNPDEPQKDVYELRYNGAIRF DSTLSGFETIMKVLSENLNAYQQQDGMMNMSPKPQRQRGHKNGQGENDNI SVAVPKSRVQQNKSIeltaedisllkkmdepYTEDTSEI
HCoV-HKU1	MSYTPGHHAGSRSSSGNRSGILKKTswvdQsershQTYNrgRkpQpKFTVST QPQGNTIPHYswfSGITQFQKGRDFKFPDGQGVPIAYGIPPSEAKGYWYKHN RRSFKTADGQQKQLLPRWYFYyLGTGPYANASYGESHEGIFWVASHQADTS IPSDVSARDPTIQEAIPTRFSPGTILPQGYyVEGSGRSASNSRPGSRsQSRGPNN RSLSRSNSNFRHSDSIVKPDMADEIASLVLAKLGKDSKPQQVTKQNAKEIRH KILMKPRQKRTPNKFCNVQQCFGKRGPLQNFGNEMLLKLTNDPQFPILAE APTPGAFFFGSKLELFKRDSADSPSKDTFELRYSGSIRFDSTLPGFETIMKVL KENLDAYVNSNQNTVSGSLSPKPQRKRGVKQSPESFDSLNLsADTQHISNDF TPEDHSLlatLDDPYVEDSVA
<b>Spike</b>	
SARS-CoV-2	MFVFLVLLPLVSSQCVNLtTTRTQLPPAYtNSfTRGVYyPDKVFRSSVLHSTQ DLFLPFFSNVTWFHAIHVSGTNGTKRFDNPVLPFNDGVYFASTeKSNIIRGWI FGTTLDskTQsLLIVNNATNVVIKvCEfQFCNDPFLGVYyHKNNKSWMESEF RVYSSANNCTFEYVSQPFLMDLEGKQGNfKNLREFVfKNIDGYfKIYSKHTP INLVRDLPQGfSALEPLVDLPiGINITRFQTLlALHRSYLTPGDSSSGWtAGAA AYYVGyLQPRtFLLKYnENGtITDAVDCALDPLSETKCTLKSfTVEKGIYQT SNFRVQPTESIVRFPNITNLCPfGEVFNATRFASVYAWNRKRISNCVADYSVL YNSASfSTfKCYGVSPTKLNDLCfTNVYADSFVIRGDEVrQIAPGQTGKIAD YNYKLpDDfTGCVIAWNSNNLDSKVGGNYNYLYRLFRKSNLKPfERDISTEI YQAGSTPCNGVEGFNCYfPLQSYGFQPTNGVGYQPyrVVVLSfELLHAPAT VCGPKKSTNLVKNKCVNFNFGLTGTGVLTESNKKfLPfQQFGRDIADTTD AVRDPQTLEILDITPCSFGGVSVITPGTNTSNQVAVLYQGVNCTEVPVAIHAD QLTPTWRVYSTGSNVfQTRAGCLIGAeHVNNsYECdIPiGAGICAsYQTQTN SPRRARSVASQSIIAYTMSLGAENSVAYSNNsIAIPTNfTISVTTEILPVsMTKT SVDCTMYICGDSTECsNLLLQYGSfCTQLNRALTGiAVEQDKNTQEVfAQV KQIYKTPPIKDFGGFNfSQILPDPSKPSKRSfIEDLLfNKVTLADAGfIKYQGD CLGDIAARDLICAQKfNGLTVLPPLLTDEMIaQYTSALLAGTITSGWTFGAG AALQIPFAMQMAyRFNGIGVTQNVLYENQKLIANQFNsAIGKIQDSLSSTAS ALGKLQDVVNQNAQALNTLVKQLSSNfGAISSVLNDILSRLDKVEAEVQIDR LITGRLQSLQTYVTQQLIRAAEIRASANLAATKMSECVLGQSKRVDFCGKGY

	<p>HLMSFPQSAPHGVVFLHVTVPAQEKNFTTAPAICHDGKAHFPREGVFVSN  GTHWFTQRNFYEPQIITDNTFVSGNCDVVIGIVNNTVYDPLQPELDSFKEE  LDKYFKNHTSPDVDLGDISGINASVVNIQKEIDRLNEVAKNLNESLIDLQELG  KYEQYIKWPWYIWLGFIAGLIAIVMVTIMLCCMTSCCSCLKGCCSCGSCCKF  DEDDSEPVLKGVKLHYT</p>
SARS-CoV	<p>MFIFLLFLTLTSGSDLDRCTTFDDVQAPNYTQHTSSMRGVYYPDEIFRSDTLY  LTQDLFLPFYSNVTGFHTINHTFGNPVIPFKDGIYFAATEKSNVVRGWVFGST  MNNKSQSVIIINNSTNVIRACNFELCDNPFFAVSKPMGTQHTMIFDNAFNC  TFEYISDAFSLDVSEKSGNFKHLREFVFKNKDGFLYVYKGYQPIDVVRDLPS  GFNTLKPIFKLPLGINITNFRAILTAFSPAQDIWGTSAAYFVGYLKPTTFMLK  YDENGITIDAVDCSQNPLAELKCSVKSFEIDKGIYQTSNFRVVPSPGDVVRFPN  ITNLCPFGEVFNATKFPSVYAWERKKISNCVADYSVLYNSTFFSTFKCYGVS  ATKLNLDLCFSNVYADSFVVKGDDVRQIAPGQTGVIADYNYKLPPDFMGCV  LAWNTRNIDATSTGNYNKYRYLRHGKLRPFERDISNVPFSPDGKPCTPPAL  NCYWPLNDYGFYTTTGIGYQPYRVVLSFELLNAPATVCGPKLSTDLIKNCQ  VNFNFNGLTGTGVLTPSSKRFQPFQFGRDVSDFTDSVRDPKTSEILDSPCSF  GGVSVITPGTNASSEVAVLYQDVNCTDVSTAIHADQLTPAWRIYSTGNNVFQ  TQAGCLIGAEHVDTSYECDIPIGAGICASYHTVSLLRSTSQKSIVAYTMSLGA  DSSIAYSNNTIAIPTNFSISITTEVMPVSMAKTSVDCNMYICGDSTECANLLQ  YGSFCTQLNRALSGIAAEQDRNTREVFAQVKQMYKTPTLKYFGGFNFSQILP  DPLKPTKRSFIEDLLFNKVTLADAGFMKQYGECLGDINARDLICAQKFNGLT  VLPPLLTDDMIAAYTAALVSGTATAGWTFGAGAALQIPFAMQMAYRFNGIG  VTQNVLYENQKQIANQFNKAISQIQESLTTTSTALGKLQDVVNQNAQALNTL  VKQLSSNFGAISSVLNDILSRDLKVEAEVQIDRLITGRLQSLQTYVTQQLIRA  AEIRASANLAATKMSECVLGQSKRVDFCGKGYHLMSFPQAAPHGVVFLHVT  YVPSQERNFTTAPAICHEGKAYFPREGVFVFNGTSWFITQRNFFSPQIITDNT  FVSGNCDVVIGIINNTVYDPLQPELDSFKEELDKYFKNHTSPDVDLGDISGIN  ASVVNIQKEIDRLNEVAKNLNESLIDLQELGKYEQYIKWPWYVWLGFIAGLI  AIVMVTILLCCMTSCCSCLKGACSCGSCCKFDEDDSEPVLKGVKLHYT</p>
MERS-CoV	<p>MIHSVFLLMFLLTPTESYVDVGPDSVKSACIEVDIQQTFFDKTWPRPIDVSKA  DGIIYPQGRITYSNITITYQGLFPYQGDHGDYVYSAGHATGTPQKLFVANY  SQDVKQFANGFVVRIGAAANSTGTVIISPSTSATIRKIYPAFMLGSSVGNFSD  GKMGRFFNHTLVLLPDGCGTLLRAFYCILEPRSGNHCPAGNSYTSFATYHTP  ATDCSDGNYNRNASLNSFKEYFNLRNCTFMYTYNITEDEILEWFGITQTAQG  VHLFSSRYVDLYGGNMFQFATLPVYDTIKYYSIIPHSIRSIQSDRKAAAFYV  YKLQPLTFLDVSVDGYIRRAIDCGFNDLSQLHCSYESFDVESGVYSVSSFEA  KPSGSVVEQAEGVECDFSPLLSGTPPQVYNFKRLVFTNCNYNLTKLLSLFSV  NDFTCSQISPAAIASNCYSSLILDYFSYPLSMKSDLSVSSAGPISQFNYKQSFSN  PTCLILATVPHNLTTITKPLKYSYINKCSRLLSDDRTEVPQLVNANQYSPCVSI  VPSTVWEDGDYRQKQLSPLEGGGWLVASGSTVAMTEQLQMFGGITVQYGT</p>

	<p>DTNSVCPKLEFANDTKIASQLGNCVEYSLYGVSGRGVFQNCTAVGVRQQR  VYDAYQNLVGYYSDDGNYYCLRACVSVPSVIYDKETKTHATLFGSVACE  HISSTMSQYSRSTRSMLKRRDSTYGPLQTPVGCVLGLVNSSLFVEDCKLPLG  QSLCALPDTPSTLTPRSVRSVPGEMRLASIAFNHPIQVDQLNSSYFKLSIPTNF  SFGVTQEYIQTTIQKVTVDCKQYVCNGFQKCEQLLREYGQFCSKINQALHGA  NLRQDDSVRNLFASVKSSQSSPIIPGFGGDFNLTLLEPVSISTGSRARSASIEDL  LFDKVTIADPGYMQGYDDCMQQGPASARDLICAQYVAGYKVLPLMDVN  MEAAYTSSLLGSIAGVGWTAGLSSFAAIPFAQSIFYRLNGVGITQQVLSNQK  LIANKFNQALGAMQTGFTTTNEAFRKVQDAVNNNAQALSKLASELSNTFGA  ISASIGDIIQRLDVLEQDAQIDRLINGRLTTLNAFVAQQLVRSESAALSAQLAK  DKVNECVKAQSKRSGFCGQGTHIVSFVVNAPNGLYFMHVGYYPSNHIEVVS  AYGLCDAANPTNCIAPVNGYFIKTNNTRIVDEWSYTGSSFYAPEPITSLNTKY  VAPQVTYQNISTNLPPLLGNSTGIDFQDELDEFFKNVSTSIPNFGSLTQINTTL  LDLTYEMLSLQQVVKALNESYIDLKELGNYTYYNKWPWYIWLGFIAGLVAL  ALCVFFILCCTGCGTNCMGKLCNRCCDRYEEYDLEPHKVHVH</p>
HCov-NL63	<p>MKLFLILLVLPLASCFFTCNSNANLSMLQLGVDPDNSSTIVTGLLPTHWICANQ  STSVYSANGFFYIDVGNHRSAFALHTGYVDVNQYYIYVTNEIGLNASVTLKI  CKFGINTTFDFLSNSSSSSFDICVNLLFTEQLGAPLGITISGETVRLHLYNVTRTF  YVPAAYKLTKLSVKCYFNYSVFSVVNATVTVNVTTHNGRVVNYTVCDDC  NGYTDNIFSQQDGRIPNGFPFNNWFLLTNGSTLVDGVSRLYQPLRLTCLWP  VPGLKSSTGFVYFNATGSDVNCNGYQHNSVADVMRYNLNFSANSVDNLKS  GVIVFKTLQYDVLFYCSNSSSGVLDTTIPFGPSSQPYYCFINSTINTTHVSTFV  GVLPPTVREIVVARTGQFYINGFKYFDLGFIEAVNFNVTTASATDFWTVAF  TFVDVLVNVSATKIQNLLYCDSPFEKLQCEHLQFGLQDGFYSANFLDDNVLP  ETYVALPIYYQHTDINFATASFGGSCYVCKPHQVNISLNGNTSVCVRTSHFS  IRYIYNRVKSGSPGDSSWHIYKSGTCPFSSKLNNFQKFKTICFSTVAVPGSC  NFPLEATWHYTSYTIVGALYVTWSEGNISITGVPYPVSGIREFSNLVLNNCTK  YNIYDYVGTGIIRSSNQSLAGGITYVSNSGNLLGFKNVSTGNIFIVTPCNQPDQ  VAVYQQSIIGAMTAVNESRYGLQNLLQLPNFYVVSNGGNNCTTAVMTYSNF  GICADGSLIPVRPRNSSDNGISAHTANLSIPSNWTTSVQVEYLQITSTPIVDC  ATYVCNGNPRCKNLLKQYTSACKTIEDALRLSAHLETNDVSSMLTFDSNAFS  LANVTSFGDYNLSSVLPQRNIHSSRIAGRSALEDLLFSKVVTSGLGTVDDY  KSCTKGLSIADLACAQYYNGIMVLPGVADAERMAMYTGSLIGGMVLGGLT  SAAAIPFSLALQARLNYVALQTDVLQENQKILAASFNKAINNIVASFSSVND  ITQTAEAIHTVTIALNKIQDVVNQQGSALNHLTSQLRHNFAISNSIQAIYDRL  DSIQADQQVDRLITGRLAALNAFVSQVLNKYTEVRSSRRLAQKINECVKSQ  SNRYGFCGNGTHIFSIVNSAPDGLLFLHTVLLPTDYKNVKAWSGICVDGIYG  YVLRQPNVLVLYSDNGVFRVTSRVMFQPRLPVLSDFVQIYNCNVTFVNISRVE  LHTVIPDYVDVNKTLQEFAQNLPKYVKPNFDLTPFNLTYNLNSSELKQLEAK  TASLFQTTVELQGLIDQINSTYVDLKLLNRFENYIKWPWWVWLIIISVVFVVL  LSLLVFCCLSTGCCGCCNCLTSSMRGCCDCGSTKLPHYEFKVVHVQ</p>

HCoV-229E	<p>MFVLLVAYALLHIAGCQTTNGTNTSHSVCNGCVGHSENVFAVESGGYIPSNF  AFNNWFLLTNTSSVVDGVVRSFQPLLLNCLWSVSGSRFTTGFVYFNGTGRG  DCKGFYSNASSDVIRYNINFEENLRRGTILFKTSYGAVVFYCTNNTLVSGDA  HIPSGTVLGNFYCFVNTTIGNETTSAFVGALPKTVREFVISRTGHFYINGYRYF  SLGNVEAVNFNVTNAATTVCTVALASYADVLVNVSQTAIANIYCNSVINRL  RCDQLSFDVPDGFYSTSPIQSVELPVSIIVSLPVYHKHTFIVLYVNFELRRGPGR  CYNCRPAVVNITLANFNETKGPLCVDTSHTTTQFVGVKFDRWSASINTGNCP  FSFGKVNNFVKFGSVCFLKDIPGGCAMPIMANLVNHKSHNIGSLYVSWSDG  DVITGVPKPVEGVSSFMNVTNLKCTKYNIYDVSGVGVIRISNDTFLNGITYTS  TSGNLLGFKDVTNGTIYSITPCNPPDQLVVYQQA VVGAMLSENFSTSYGFSNV  VEMPKFFYASNGTYNCTDAVLTYSSFGVCADGSIIAVQPRNVSYDSVSAIVT  ANLSIPSNWTTSVQVEYLQITSTPIVDCSTYVCNGNVRCVELLKQYTSACK  TIEDALRNSAMLESADVSEMLTFDKKAFTLANVSSFGDYNLSSVIPSLPRSGS  RVAGRSAIEDILFSKLVTSGLGTVDADYKKCTKGLSIADLACAQYYNGIMVL  PGVADAERMAMYTGSLIGGIALGGLTSAASIPFSLAIQSRLNYVALQTDVLQ  ENQKILAASFNKAMTNIVDAFTGVNDAITQTSQALQTVATALNKIQDVVNQ  QGNSLNHLTSQLRQNFQAISSSIQAIYDRLDIIQADQQVDRLITGRLAALNVF  VSHTLTKYTEVRASRLAQQKVNECVKSQSKRYGFCGNGTHIFSLVNAAPE  GLVFLHTVLLPTQYKDVEAWSGLCVDGINGYVLRQPNLALYKEGNYYRITS  RIMFEPRIPTIADFVQIENCNVTFVNISRSELQTIVPEYIDVNKTLQELSYKLPN  YTVPDLVVEQYNQ TILNLTSEISTLENKSAELNYTVQKLQTLIDNINSTLVDL  KWLNRVETYIKWPWWVWLCISVVLI FVVSMLLLCCCSTGCCGFFSCFASSIR  GCCESTKL PYYDVEKIHQ</p>
HCoV-OC43	<p>MFLILLISLPTAFAVIGDLNCPLDPRLKGSFNNRDTGPPSISTDTVDVTNGLGT  YYVLDRVYLNNTLFLNGYYPTSGSTYRNMALKGTDLLSTLWFKPPFLSDFIN  GIFAKVKNTKVFKDGV MYSEFPAITIGSTFVNTSYSVVVQPTINSTQDGVN  KLQGLLEVSV CQYNMCEYPHTICHPNLGNHFKELWHLDTGVVSCLYKRNF  YDVNATYLYFHFYQEGGTFYAYFTDTGFVTKFLFNVYLGMA LSHYYVMPL  TCISRRDIGFTLEYWVTP LTPRQYLLAFNQDGIIFNAVDCMSDFMSEIKCKTQ  SIAPPTGVYELNGYTVQPIADVYRRKPDLPNCNIEAWLNDKSVPSPLNWERK  TFSNCNFMSSLM SFIQADSFTCNNIDA AKIYGMCFSSITIDKFAIPNRRKVDL  QLGNLGYLQSSNYRIDTTATSCQLYYNLPAANVSVS RFPSTWNKRFGFIED  SVFVPQPTGVFTNHSVVYAQHCFKAPKNFCPCSSCPGKNNGIGTCPAGTNYL  TCDNLCTLDPITFKAPDTYKCPQTKSLVGIGEHCSGLAVKSDYCGNNSCTCQ  PQAFLGWSAD SCLQGDKCNIFANFILHDVNNGLT CSTDLQKANTEIELGVCV  NYDLYGISGQGIFVEVNATYYNSWQNLLYDSNGNLYGFRDYITNRTFMIHSC  YSGRVSAAYHANSSEPALLFRNIKCNYVFNNSLTRQLQPINYSFDSYLGCVV  NAYNSTAISVQTCDLTVGSGYCVDY SKNRRSRRAITTYRFTNFEPFTVNSV  NDSLEPVGGLYEIQIPSEFTIGNMEEFIQTSSPKVTIDCAAFVCGDYAACKLQL  VEYGSFCDNINAILTEVNELDDTTQLQVANSLMNGVTLSTKLKDG VNFNVD</p>

	DINFSPVLGCLGSECSKASSRSAIEDLLFDKVKLSDVGFVEAYNNCTGGAEIR DLICVQSYKGIVLPPLLESENQISGYTLAATSASLFPPWTAAGVPFYLNVQY RINGLGVTMDVLSQNQKLIANAFNNALHAIQQGF DATNSALVKIQAVVNAN AEALNNLLQQLSNRFGAISASLQEILSRLDALEAEAQIDRLINGRLTALNAYV SQQLSDSL VKFSAAQAMEKVNCEVKSQSSRINFCGNGNHIISLVQNAPYGL YFIHFNYVPTKYVTAKVSPGLCIAGNRGIAPKSGYFVN VNNTWMYTGSGYY YPEPITENNVVVMSTCAVNYTKAPYVMLNTSIPNLPDFKEELDQWFKNQTS VAPDLSLDYINVTFLDLQVEMNRLQEAIKVLNHSYINLKDIGTYEYYVKWP WYVWLLICLAGVAMLVLLFFICCCTGCGTSCFKKCGGCCDDYTGYQELVIK TSHDD
HCoV-HKU1	MLLIIFILPTTLAVIGDFNCTNFAINDKNTTVPRISEYVVDVSYGLGTYIYILDR VYLNNTTILFTGYFPKSGANFRDLSLKGTTYLSTLWYQKPFLSDFNNGIFSRVK NTKLYVNKTLYSEFSTIVIGSVFINNSYITIVVQPHNGVLEITACQYTMCEYPH TICKSKGSSRNESWHFDKSEPLCLFKKNFTYNVSTDWLYFHFYQERGTFYAY YADSGMPTTFLFSLYLGTLSSHYYVLPLTCNAISSNTDNETLQYWVTPLSKR QYLLKFDNRGVITNAVDCSSSFFSEIQCKTKSLLPNTGVYDLSGFTVKPVATV HRRIPDLPCDIDKWLNNFNVP SPLNWERKIFSNCN FNLS TLLRLVHTDSFSC NNFDESKIYGSCFKSIVLDKFAIPNSRRSDLQLGSSGFLQSSNYKIDTTSSSCQ LYYSLPAINVTINNYNPSSWNRRYGFNNFNLSHSHSVVYSRYCFSVNNTFCPC AKPSFASSCKSHKPPSASCPIGTNYRSCESTTVLDHTDWCRCSCLPDPITAYD PRSCSQKKSLVGVGEGHCAGFGVDEEKCGLVDGSGYNV SCLCSTDAFLGWSYD TCVSNNR CNIFSNFILNGINSGTTCSNDLLQPNT EVFTDVCVDYDLYGITGQG IFKEVSAVYYNSWQNLLYDSNGNIIGFKDFVTNKTYNIFPCYAGRVSAAFHQ NASSLALLYRNLC KSYVLNNISLATQPYFDSYLGCVFNADNLTDYSVSSCAL RMGSGFCVDYNSPSSSSSRKRRSISASYRFVTFEPFNVSFVNDSIESVGGLYE IKIPTNFTIVGQEEFIQTNSPKVTIDCSLFVCSNYAACHDLLSEYGTFCDNINSI LDEVNGLLDTTQLHVADTLMQGVTLSSNLNTNLHFDVDNINFKSLVGCLGP HCGSSSRFFEDLLFDKVKLSDVGFVEAYNNCTGGSEIRDLLCVQSFNGIKVL PPILSESQISGYTTAATVAAMFPPWSAAAGIPFSLNVQYRINGLGVTMDVLN KNQKLIATAFNNALLSIQNGFSATNSALAKIQSVVNSNAQALNSLLQQLFNK FGAISSSLQEILSRLDALEAQVQIDRLINGRLTALNAYVSQQLSDISLVKFGAA LAMEKVNCEVKSQSPRINFCGNGNHILSLVQNAPYGLLFMHFSYKPI SFKTV LVSPGLCISGDVGIAPKQGYFIKHNDHWMFTGSSYYYPEPISDKNVVFMNTC SVNFTKAPLVYLNHSPKLSDFESELSHWFKNQTSIAPNLTLNLHTINATFLD LYYEMNLIQESIKSLNNSYINLKDIGTYEMYVKWPWYVWLLISFSFIIFLVLLF FICCCTGCGSACFSKCHNCCDEYGGHHDFVIKTSHDD

**Supplementary Table S2: List of all epitopes predicted with CTLpred.** Consensus sequences of Envelope, Membrane, Nucleocapsid, and Spike proteins from SARS-CoV-2, SARS-CoV, MERS-CoV, HCoV-NL63, HCoV-229E, HCoV-OC43, and HCoV-HKU1 were fed into the CTLpred (<https://webs.iiitd.edu.in/raghava/ctlpred/>) with ANN/SVM combined approach with default settings. Shortlisted epitopes used for the vaccine construct are highlighted in bold letters.

<b>SARS-CoV-2</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	13	IVNSVLLFL	0.89/1.3323721
	2	41	AYCCNIVNV	0.50/1.7081903
	3	51	LVKPSFYVY	0.70/0.7574663
	4	16	SVLLFLAFV	0.03/1.2319935
	5	57	YVYSRVKNL	0.43/0.65777634
	6	35	TALRLCAYC	0.80/0.27596158
	7	25	VLLVTLAI	0.99/0.065623615
	<b>8</b>	<b>50</b>	<b>SLVKPSFYV</b>	<b>0.00/0.99144443</b>
	9	38	RLCAYCCNI	0.27/0.70361798
	10	20	FLAFVVFL	0.02/0.94021965
	11	17	VLLFLAFVV	0.00/0.86079609
	12	23	FVVFLVTL	0.08/0.76653538
	13	26	FLLVTLAIL	0.10/0.72134398
	14	11	TLIVNSVLL	0.08/0.73778864
	15	24	VVFLVTLA	0.45/0.36571657
	16	34	LTALRLCAY	0.74/0.0034344671
	17	19	LFLAFVVFL	0.31/0.42809059
	18	53	KPSFYVYSR	0.02/0.66563238
	19	66	NSSRVPDLL	/0.67715744
	20	43	CCNIVNVSL	0.02/0.64476164
	21	44	CNIVNVSLV	0.10/0.55377124
	22	5	VSEETGTLI	0.77/-0.12141075
	23	12	LIVNSVLLF	0.64/-0.029654824
	24	18	LLFLAFVVF	0.01/0.57949258
	25	52	VKPSFYVYS	0.99/-0.45153825



	26	56	FYVYSRVKN	0.94/-0.42336705
	27	42	YCCNIVNVS	0.90/-0.39556418
	<b>28</b>	<b>32</b>	<b>AILTALRLC</b>	<b>0.61/-0.15241169</b>
	29	36	ALRLCAYCC	0.01/0.40105212
	30	37	LRLCAYCCN	0.97/-0.57229533
	31	30	TLAILTALR	0.00/0.36066322
	32	60	SRVKNLNSS	0.76/-0.51255186
	33	27	LLVTLAILT	0.95/-1.0998769
	34	2	YSFVSEETG	0.63/-0.89302618
<b>SARS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	13	IVNSVLLFL	0.89/1.3323721
	2	41	AYCCNIVNV	0.50/1.7081903
	<b>3</b>	<b>50</b>	<b>SLVKPTVYV</b>	<b>0.87/1.1235574</b>
	4	16	SVLLFLAFV	0.03/1.2319935
	5	51	LVKPTVYVY	0.45/0.78461679
	6	57	YVYSRVKNL	0.43/0.65777634
	7	35	TALRLCAYC	0.80/0.27596158
	8	25	VFLLVTLAI	0.99/0.065623615
	9	38	RLCAYCCNI	0.27/0.70361798
	10	20	FLAFVVFL	0.02/0.94021965
	11	17	VLLFLAFVV	0.00/0.86079609
	12	23	FVVFLLVTL	0.08/0.76653538
	13	26	FLLVTLAIL	0.10/0.72134398
	14	11	TLIVNSVLL	0.08/0.73778864
	15	24	VVFLLVTLA	0.45/0.36571657
	16	34	LTALRLCAY	0.74/0.0034344671
	17	19	LFLAFVVFL	0.31/0.42809059
	18	43	CCNIVNVSL	0.02/0.64476164
	19	44	CNIVNVSLV	0.10/0.55377124
	20	5	VSEETGTLI	0.77/-0.12141075
	21	12	LIVNSVLLF	0.64/-0.029654824
	22	18	LLFLAFVVF	0.01/0.57949258
	23	66	NSSEGVPDL	/0.58120937
	24	67	SSEGVPDLL	/0.57470504
	25	42	YCCNIVNVS	0.90/-0.39556418
	26	56	VYVYSRVKN	0.94/-0.46663835
	<b>27</b>	<b>32</b>	<b>AILTALRLC</b>	<b>0.61/-0.15241169</b>

	28	36	ALRLCAYCC	0.01/0.40105212
	29	48	NVSLVKPTV	0.00/0.40380231
	30	37	LRLCAYCCN	0.97/-0.57229533
	31	30	TLAILTALR	0.00/0.36066322
	32	52	VKPTVYVYS	0.97/-0.9738047
	33	27	LLVTLAILT	0.95/-1.0998769
	34	2	YSFVSEETG	0.63/-0.89302618
	35	47	VNVSLVKPT	0.69/-1.5062236
	36	65	LNSSEGVPD	0.90/-1.8194142
<b>MERS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	50	LLVQPALYL	0.79/0.84185633
	2	1	MLPFVQERI	0.94/0.49325186
	3	25	ITLLVCMAF	0.96/0.38790733
	4	24	AITLLVCMA	0.64/0.66448331
	5	9	IGLFIVNFF	0.44/0.78198539
	6	44	MTGFNTLLV	0.31/0.71660277
	7	8	RIGLFIVNF	0.05/0.97063083
	8	13	IVNFFIFTV	0.00/1.0043246
	9	60	NTGRSVYVK	0.31/0.65772969
	10	26	TLLVCMAFL	0.08/0.85346175
	11	57	YLYNTGRSV	0.00/0.89588326
	12	12	FIVNFFIFT	0.82/0.061943241
	13	42	QCMTGFNTL	0.91/-0.050979385
	14	36	ATRLCVQCM	0.00/0.84974322
	15	19	FTVVCAITL	0.04/0.80762511
	16	17	FIFTVVCAI	0.17/0.66048223
	17	16	FFIFTVVCA	0.25/0.56456289
	18	53	QPALYLYNT	0.52/0.26424758
	19	10	GLFIVNFFI	0.11/0.54082069
	20	3	PFVQERIGL	0.75/-0.14228684
	21	11	LFIVNFFIF	0.03/0.55199648
	22	47	FNTLLVQPA	0.85/-0.28167078
	23	58	LYNTGRSVY	0.05/0.51129716
	24	31	MAFLTATRL	0.13/0.40104314
	25	59	YNTGRSVYV	0.16/0.36364365
	26	33	FLTATRLCV	0.12/0.40129342
	27	55	ALYLYNTGR	0.99/-0.47537099

	28	20	TVVCAITLL	0.02/0.49219949
	29	6	QERIGLFIV	0.00/0.50545109
	<b>30</b>	<b>32</b>	<b>AFLTATRLC</b>	<b>0.59/-0.093565614</b>
	31	52	VQPALYLYN	0.96/-0.54955225
	32	30	CMAFLTATR	0.00/0.37374539
	33	27	LLVCMAFLT	0.67/-0.8292502
	34	41	VQCMTGFNT	0.66/-0.85129807
	35	45	TGFNTLLVQ	0.90/-1.1351984
	36	34	LTATRLCVQ	0.68/-1.0467289
	37	70	QDSKPPLPP	0.99/-1.5643758
	38	67	VKFQDSKPP	0.79/-1.6396241
<b>HCoV-NL63</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	30	MTFIKLIQL	0.85/1.1147816
	2	21	VMIFFFVLA	0.97/0.84537219
	3	12	VLNSILWLL	0.52/1.0318757
	4	5	LIDDNGIVL	0.89/0.42154725
	5	55	KIFLAYQDY	0.91/0.35324098
	6	19	LLVMIFFFV	0.48/0.70758604
	7	16	ILWLLVMIF	0.69/0.416904
	8	65	QIAPVPAEV	0.15/0.93282364
	9	37	QLCFTCHYF	0.57/0.49749676
	10	8	DNGIVLNSI	0.33/0.71998752
	11	24	FFFVLAMTF	0.70/0.34339497
	12	22	MIFFFVLAM	0.01/1.0168252
	13	48	RTLQPVYKI	0.09/0.9221873
	14	25	FFVLAMTFI	0.92/0.076178019
	15	4	RLIDDNGIV	0.06/0.88782061
	16	59	AYQDYMQIA	0.27/0.52539488
	17	43	HYFFSRTLQ	0.84/-0.04528681
	18	50	LQPVYKIFL	0.67/0.12251192
	19	28	LAMTFIKLI	0.97/-0.22110032
	20	27	VLAMTFIKL	0.01/0.7382466
	21	17	LWLLVMIFF	0.53/0.20745591
	22	66	IAPVPAEVL	/0.71392143
	23	49	TLQPVYKIF	0.01/0.60842696
	24	20	LVMIFFFVL	0.00/0.60981957
	25	13	LNSILWLLV	0.98/-0.45138175

	26	68	PVPAEVLNV	/0.52237808
	27	47	SRTLQPVYK	0.05/0.41973046
	28	34	KLIQLCFTC	0.02/0.37888815
	29	44	YFFSRTLQP	0.77/-0.47763152
	30	62	DYMQIAPVP	0.83/-0.76907218
<b>HCoV-229E</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	16	LLWCVVLI	0.97/1.4253784
	2	30	ITIILIKL	0.94/1.2826698
	3	7	DDHALVVNV	0.79/0.97984782
	4	66	HIDPFKRV	0.37/1.176099
	5	12	VNVLLWCV	0.95/0.50794883
	6	37	KLCFTCHMF	0.89/0.43601752
	7	14	NVLLWCVVL	0.94/0.24113564
	8	19	CVVLIVILL	0.26/0.76056346
	9	45	FCNRTVYGP	0.93/0.041517814
	10	15	VLLWCVVLI	0.18/0.79078298
	11	27	LVCITIKL	0.04/0.88417935
	12	20	VVLIVILLV	0.08/0.8400338
	13	4	KLVDHALV	0.46/0.45693719
	14	57	VYHIQSYM	0.01/0.89704074
	15	41	TCHMFCNRT	0.78/0.038413593
	16	24	VILLVCITI	0.31/0.49818172
	17	59	HIYQSYMHI	0.13/0.64915513
	18	13	VNVLLWCVV	0.67/0.00080872534
	19	49	TVYGPIKNV	0.20/0.46726731
	20	56	NVYHIYQSY	0.13/0.45685054
	21	8	DHALVVNVL	0.01/0.53900496
	22	40	FTCHMFCNR	0.03/0.48649853
	23	52	GPIKNVYHI	0.00/0.51594748
	24	34	KLIKLCFTC	0.04/0.46911404
	25	18	WCVVLIVIL	0.03/0.44295609
	26	60	IYQSYMHI	0.95/-0.53151165
	27	55	KNVYHIYQS	0.93/-0.51524004
	28	25	ILLVCITII	0.00/0.3944171
	29	58	YHIYQSYM	0.95/-0.59901297
	30	48	RTVYGPIKN	0.92/-0.58362683

	31	28	VCITHIKLI	0.51/-0.30207982
	32	61	YQSYMHDIP	0.83/-0.93658136
	33	33	IKLIKLCFT	0.56/-0.99293995
<b>HCoV-OC43</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	7	YLADTVWYV	0.31/2.3652348
	2	55	SPSIYVFNR	0.84/1.0516232
	3	27	LVTIVVVAF	0.80/1.0279233
	4	23	AICLLVTIV	0.74/0.93439759
	5	52	LVLSPSIYV	0.99/0.66790173
	6	66	QFYEFYNDI	0.71/0.79282771
	7	24	ICLLVTIVV	0.86/0.62629457
	8	13	WYVGQIIFI	0.54/0.92308938
	9	32	VVAFLATFK	0.95/0.49439065
	10	73	DIKPPVLDV	0.75/0.67326274
	11	19	IFIVAICLL	0.82/0.53378928
	12	31	VVVAFLATF	0.84/0.36277806
	13	6	AYLADTVWY	0.86/0.32145213
	14	22	VAICLLVTI	0.92/0.21636364
	15	18	IIFIVAICL	0.01/1.1021567
	16	63	RGRQFYEFY	0.93/0.089834301
	17	59	YVFNRGRQF	0.61/0.37921737
	18	29	TIVVVAFLA	0.95/0.0087496811
	19	40	KLCIQLCGM	0.50/0.45029766
	20	46	CGMCNTLVL	0.93/0.0047116177
	21	37	ATFKLCIQL	0.04/0.8126547
	22	25	CLLVTIVVV	0.02/0.79108189
	23	14	YVGQIIFIV	0.00/0.75333868
	24	70	FYNDIKPPV	0.59/0.16051728
	25	58	IYVFNRGRQ	0.89/-0.15046971
	26	17	QIIFIVAIC	0.28/0.44484173
	27	38	TFKLCIQLC	0.66/0.063805722
	28	28	VTIVVVAFL	0.02/0.63313612
	29	67	FYEFYNDIK	0.64/-0.024399741
	30	16	GQIIFIVAI	0.00/0.61229884
	31	12	VWYVGQIIF	0.63/-0.022659664
	32	15	VGQIIFIVA	0.62/-0.017997626
	33	62	NRGRQFYEF	0.01/0.55313485

	34	60	VFNRGRQFY	0.00/0.53977627
	35	53	VLSPSIYVF	0.01/0.52058673
	36	11	TVWYVGQII	0.00/0.51642822
	37	43	IQLCGMCNT	0.72/-0.2621705
	38	35	FLATFKLCI	0.00/0.41596129
	39	33	VAFLATFKL	0.61/-0.22762835
	40	45	LCGMCNTLV	0.58/-0.52434988
	41	54	LSPSIYVFN	0.85/-1.000583
	42	72	NDIKPPVLD	0.65/-0.93329455
<b>HCoV-HKU1</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Envelope</b>	1	5	FFTDTAWYI	0.97/0.81010327
	2	25	IFLIFVVAL	0.68/1.0860147
	3	19	LVLSCVIFL	0.81/0.66262758
	4	12	YIGQIFFLV	0.89/0.40362005
	5	21	LSCVIFLIF	0.81/0.46884947
	6	11	WYIGQIFFL	0.18/1.0211905
	7	50	FIISPSAYV	0.69/0.48887938
	8	24	VIFLIFVVA	0.74/0.4330104
	9	22	SCVIFLIFV	0.01/1.1354362
	10	43	ICGFCNIFI	0.87/0.22423108
	11	41	IQICGFCNI	0.37/0.63052644
	12	26	FLIFVVALL	0.03/0.89957281
	13	59	YNRGRQLYK	0.00/0.90229452
	14	10	AWYIGQIFF	0.47/0.42991415
	15	53	SPSAYVYNR	0.03/0.86586641
	16	14	GQIFFLVLS	0.97/-0.086289852
	17	52	ISPSAYVYN	0.96/-0.1175764
	18	20	VLSCVIFLI	0.05/0.78040404
	19	44	CGFCNIFII	0.19/0.62693374
	20	27	LIFVVALLA	0.98/-0.16718043
	21	58	VYNRGRQLY	0.04/0.75198527
	22	70	SEHVIPSTL	0.29/0.38945586
	23	8	DTAWYIGQI	0.02/0.65013492
	24	35	ATIKLCIQI	0.00/0.66442936
	25	55	SAYVYNRGR	0.98/-0.34168303
	26	1	MVDVFFTDT	0.55/0.064258598
	27	60	NRGRQLYKS	0.58/0.015687379

	28	48	NIFIISPSA	0.74/-0.15448485
	29	18	FLVLSCVIF	0.01/0.50903
	30	13	IGQIFFLVL	0.13/0.38814528
	31	3	DVFFTDTAW	0.01/0.46213887
	32	57	YVYNRGRQL	0.01/0.42747013
	33	73	VIPSTLDDL	/0.38726384
	34	7	TDTAWYIGQ	0.80/-0.75896706
	35	67	KSYSEHVIP	0.71/-0.69228345
<b>SARS-CoV-2</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	38	AYANRNRFL	0.94/1.534005
	2	41	NRNRFLYII	0.96/1.2950944
	3	43	NRFLYIIKL	0.79/1.3086849
	4	15	KLLEQWNLV	0.54/1.1477859
	5	104	ARTRSMWSF	0.87/0.79456645
	6	21	NLVIGFLFL	0.30/1.3302649
	7	83	AMACLVGLM	0.96/0.62580832
	8	130	TRPLLESEL	0.01/1.5277131
	9	46	LYIIKLIFL	0.00/1.4599298
	10	39	YANRNRFLY	0.70/0.75749834
	11	37	FAYANRNR	0.71/0.72206684
	12	99	SFRLFARTR	0.98/0.43485337
	13	136	SELVIGAVI	0.92/0.48139255
	14	50	KLIFLWLLW	0.97/0.41640893
	15	85	ACLVGLMWL	0.85/0.52376312
	16	172	TSRTLSTYYK	0.68/0.67723726
	17	193	FAAYSRYRI	0.84/0.50777994
	18	112	FNPETNILL	0.41/0.92818671
	19	45	FLYIIKLIF	0.83/0.5058283
	20	61	TLACFVLAA	0.43/0.89317424
	21	196	YSRYRIGNY	0.91/0.39062916
	22	40	ANRNRFLYI	0.05/1.2284371
	23	34	LLQFAYANR	0.94/0.28716981
	24	198	RYRIGNYKL	0.05/1.1734506
	25	173	SRTLSTYYKL	0.40/0.79618648
	26	92	WLSYFIASF	0.34/0.84325684
	27	96	FIASFRLFA	0.13/1.0490459
	28	49	IKLIFLWLL	0.53/0.6187025

29	212	SSSDNIALL	/1.1372182
30	5	NGTITVEEL	0.99/0.1463184
31	192	GFAAYSRYR	0.80/0.32961293
32	65	FVLAAYRI	0.00/1.1177805
33	168	ITVATSRTL	0.99/0.09512733
34	55	WLLWPVTLA	0.65/0.40702879
<b>35</b>	<b>94</b>	<b>SYFIASFRL</b>	<b>0.33/0.72395431</b>
36	73	INWITGGIA	0.90/0.14883025
37	52	IFLWLLWPV	0.91/0.13060966
<b>38</b>	<b>111</b>	<b>SFNPETNIL</b>	<b>0.18/0.83545969</b>
39	153	GHHLGRCDI	0.80/0.20706352
40	42	RNRFLYIIK	0.00/1.0055272
41	197	SRYRIGNYK	0.00/1.0036465
42	142	AVILRGHLR	0.31/0.6842731
43	48	IIKLIFLWL	0.25/0.73918596
44	26	FLFLTWICL	0.28/0.70891806
45	195	AYSRYRIGN	0.79/0.18502933
46	74	NWITGGIAI	0.93/0.039965501
47	171	ATSRTLSTYY	0.00/0.94722714
48	141	GAVILRGHL	0.84/0.095706564
49	72	RINWITGGI	0.91/0.022493064
50	68	AAVYRINWI	0.00/0.91891809
51	27	LFLTWICLL	0.89/-0.0071357757
52	98	ASFRLFART	0.41/0.46729682
53	75	WITGGIAIA	0.77/0.098162084
54	2	ADSNGTITV	0.68/0.1653799
55	137	ELVIGAVIL	0.01/0.83354512
56	138	LVIGAVILR	0.76/0.075370132
57	148	HLRIAGHHL	0.10/0.73433113
58	30	TWICLLQFA	0.94/-0.13203349
59	116	TNILLNVPL	0.59/0.2051763
60	56	LLWPVTLAC	0.62/0.17025753
61	58	WPVTLACFV	0.27/0.51367294
62	166	KEITVATSR	0.80/-0.040709
63	101	RLFARTRSM	0.04/0.71755901
64	211	SSSSDNIAL	/0.74666418
65	88	VGLMWLSYF	0.75/-0.032093333
66	24	IGFLFTWI	0.93/-0.21772675



67	8	ITVEELKKL	0.80/-0.095865837
68	19	QWNLVIGFL	0.89/-0.18860719
69	144	ILRGHLRIA	0.00/0.69453508
70	178	YYKLGASQR	0.02/0.66064316
71	79	GIAIAMACL	0.20/0.45983994
72	126	GTILTRPLL	0.00/0.65466291
73	194	AAYSRYRIG	0.08/0.55879723
74	103	FARTRSMWS	1.00/-0.39487449
75	36	QFAYANRNR	0.00/0.59525233
76	102	LFARTRSMW	0.09/0.50026909
77	186	RVAGDSGFA	0.97/-0.39718197
<b>78</b>	<b>162</b>	<b>KDLPKEITV</b>	<b>0.12/0.44772671</b>
79	76	ITGGIAIAM	0.03/0.52725066
80	97	IASFRLFAR	0.01/0.52982343
81	44	RFLYIIKLI	0.00/0.51353245
82	134	LESELVIGA	0.00/0.51224851
83	70	VYRINWITG	0.00/0.50995909
84	64	CFVLAAVYR	0.01/0.49918463
<b>85</b>	<b>188</b>	<b>AGDSGFAAY</b>	<b>0.06/0.43852094</b>
86	154	HHLGRCDIK	0.62/-0.14339661
87	119	LLNVPLHGT	0.99/-0.51433219
88	69	AVYRINWIT	0.04/0.43528962
89	67	LAAVYRINW	0.01/0.45284658
90	176	LSYYKLGAS	0.99/-0.53726447
91	201	IGNYKLNTD	0.70/-0.29728934
92	113	NPETNILLN	0.97/-0.62630241
93	179	YKLGASQRV	0.55/-0.22732784
94	87	LVGLMWLSY	0.62/-0.37756975
95	35	LQFAYANRN	0.98/-0.73842493
96	7	TITVEELKK	0.79/-0.58360765
97	81	AIAMACLVG	0.64/-0.44634314
98	203	NYKLNTDHS	0.54/-0.35367928
99	127	TILTRPLLE	0.60/-0.41466275
100	108	SMWSFNPET	0.60/-0.45458388
101	152	AGHHLGRCD	0.83/-0.69033051
102	164	LPKEITVAT	0.55/-0.57245695
103	118	ILLNVPLHG	0.57/-0.6114191
104	156	LGRCDIKDL	0.80/-0.87765572
105	165	PKEITVATS	0.74/-1.0843754

	106	133	LLESELVIG	0.71/-1.2327109
<b>SARS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	95	FVASFRLFA	0.68/1.2821694
	2	41	RNRFLYIIK	0.93/1.0055272
	3	25	FLFLAWIML	0.94/0.87487652
	4	196	NRYSRIGNYK	0.92/0.87277329
	5	125	GTIVTRPLM	0.88/0.9095859
	6	64	FVLAADVYRI	0.64/1.1177805
	7	40	NRNRFLYII	0.40/1.2950944
	8	14	QLLEQWNLV	0.82/0.68995958
	9	100	RLFARTRSM	0.79/0.71755901
	10	37	AYSNNRNRFL	0.09/1.3992408
	11	122	PLRGTIVTR	0.92/0.52673674
	12	1	MADNGTITV	0.69/0.72183355
	13	97	ASFRLFART	0.94/0.46729682
	14	129	TRPLMESEL	0.29/1.0952626
	15	20	NLVIGFLFL	0.01/1.3302649
	16	38	YSNNRNRFLY	0.33/0.99584681
	17	42	NRFLYIIKL	0.01/1.3086849
	18	197	RYRIGNYKL	0.13/1.1734506
	19	61	LACFVLAADV	0.95/0.3289888
	20	88	GLMWLSYFV	0.77/0.50853016
	21	172	SRTLSSYYKL	0.48/0.79618648
	22	211	GSNDNIAL	/1.2503836
	23	74	WVTGGIAIA	0.94/0.29104438
	24	177	YYKLGASQR	0.51/0.66064316
	25	193	AAYNRYRIG	0.92/0.22922506
	26	202	NYKLNTDHA	0.91/0.21813252
	27	143	IIRGHLRMA	0.87/0.19978509
	28	5	GTITVEELK	0.97/0.09780617
	29	36	FAYSNNRNR	0.03/1.0373734
	30	21	LVIGFLFLA	0.76/0.28726772
	31	8	TVEELKQLL	0.54/0.49652662
	32	39	SNRNRFLYI	0.10/0.92906309
	33	92	LSYFVASFR	0.80/0.2121516
	<b>34</b>	<b>187</b>	<b>GTDSGFAAY</b>	<b>0.24/0.75167131</b>
	35	138	VIGAVIIRG	0.96/0.027776638

36	170	ATSRTLSTYY	0.02/0.94722714
37	84	ACIVGLMWL	0.00/0.95329943
38	91	WLSYFVASF	0.23/0.72279954
39	165	KEITVATSR	0.99/-0.040709
40	111	FNPETNILL	0.02/0.92818671
41	136	ELVIGAVII	0.67/0.26083068
42	45	LYIIKLVFL	0.08/0.84998052
<b>43</b>	<b>161</b>	<b>KDLPKEITV</b>	<b>0.48/0.44772671</b>
44	159	DIKDLPKEI	0.99/-0.064095269
45	133	MESELVIGA	0.07/0.85535516
46	60	TLACFVLAA	0.01/0.89317424
47	54	WLLWPVTLA	0.48/0.40702879
<b>48</b>	<b>110</b>	<b>SFNPETNIL</b>	<b>0.05/0.83545969</b>
49	192	FAAYNRYRI	0.01/0.86589126
50	48	IKLVFLWLL	0.11/0.76135453
51	195	YNRYRIGNY	0.57/0.26868248
52	47	IILVFLWL	0.05/0.76608405
53	103	ARTRSMWSF	0.01/0.79456645
54	75	VTGGIAIAM	0.22/0.58328221
55	19	WNLVIGFLF	0.74/0.058311061
56	141	AVIIRGHLR	0.02/0.76535018
57	119	LVPLRGTI	0.92/-0.14422858
58	189	DSGFAAYNR	0.22/0.53397851
59	116	NILLNVPLR	0.11/0.61971381
60	194	AYNRYRIGN	0.53/0.19031694
61	101	LFARTRSMW	0.22/0.50026909
62	120	NVPLRGTV	0.05/0.66206936
63	171	TSRTLSTYYK	0.03/0.67723726
64	13	KQLLEQWNL	0.03/0.66756564
65	86	IVGLMWLSY	0.85/-0.16115119
66	67	AAVYRINWV	0.11/0.4817265
67	49	KLVLWLLW	0.08/0.49584252
68	178	YKLGASQRV	0.77/-0.22732784
69	28	LAWIMLLQF	0.55/-0.016850555
70	63	CFVLAAYVR	0.03/0.49918463
71	154	SLGRCDIKD	0.97/-0.44256032
72	57	WPVTLACFV	0.01/0.51367294
<b>73</b>	<b>93</b>	<b>SYFVASFRL</b>	<b>0.01/0.49716475</b>
74	135	SELVIGAVI	0.02/0.48139255

75	69	VYRINWVTG	0.00/0.48186639
76	145	RGHLRMAGH	0.12/0.36179993
77	76	TGGIAIAMA	0.84/-0.36138282
78	99	FRLFARTRS	0.66/-0.18296228
79	82	AMACIVGLM	0.00/0.4659084
80	98	SFRLFARTR	0.03/0.43485337
81	44	FLYIHKLVF	0.01/0.45288171
82	66	LAAVYRINW	0.00/0.45284658
83	53	LWLLWPVTL	0.98/-0.52996139
84	2	ADNGTITVE	0.99/-0.5458222
85	78	GIAIAMACI	0.01/0.41953955
86	96	VASFRLFAR	0.02/0.39749627
87	140	GAVIIRGHL	0.00/0.41030383
88	32	MLLQFAYSN	0.75/-0.34359439
89	149	RMAGHSLGR	0.52/-0.1219183
90	58	PVTLACFVL	0.71/-0.34015016
91	89	LMWLSYFVA	0.73/-0.36412405
92	200	IGNYKLNTD	0.63/-0.29728934
93	72	INWVTGGIA	0.67/-0.38085986
94	112	NPETNILLN	0.89/-0.62630241
95	16	LEQWNLVIG	0.72/-0.49429055
96	3	DNGTITVEE	0.97/-0.74767274
97	15	LLEQWNLVI	0.83/-0.62161766
98	139	IGAVIIRGH	0.84/-0.70317598
99	157	RCDIKDLPK	0.95/-0.82975361
100	155	LGRCDIKDL	0.99/-0.87765572
101	77	GGIAIAMAC	0.64/-0.5324313
102	121	VPLRGTIVT	0.53/-0.42807568
103	107	SMWSFNPET	0.55/-0.45458388
104	50	LVFLWLLWP	0.86/-0.88953572
105	203	YKLNTDHAG	0.59/-0.69082897
106	90	MWLSYFVAS	0.64/-0.81459416
107	201	GNYKLNTDH	0.99/-1.1881523
108	148	LRMAGHSLG	0.57/-0.78018021
109	144	IRGHLRMAG	0.52/-0.76672312
110	146	GHLRMAGHS	0.97/-1.2795936
111	176	SYYYKL GASQ	0.54/-0.90215334
<b>MERS-CoV</b>			

Viral protein name	Rank	Start position	Sequence	Score (ANN/SVM)
<b>Membrane</b>	<b>1</b>	<b>161</b>	<b>DRLPNEVTV</b>	<b>0.90/1.3325897</b>
	2	170	AKPNVLIAL	0.99/1.112663
	3	92	ISYFVQSIR	0.84/0.9756516
	4	138	SVTAVVTNG	0.96/0.50240003
	5	20	NFAWSLIFL	0.02/1.3750567
	6	15	IKDWNFAW	0.76/0.63209891
	<b>7</b>	<b>187</b>	<b>GTNSGVAIY</b>	<b>0.29/1.043693</b>
	8	22	AWSLIFLLI	0.80/0.4916732
	9	38	YPSRSMTVY	0.93/0.30431911
	10	44	TVYVFKMFV	0.29/0.9305296
	11	133	VEDSTSVTA	0.74/0.47422488
	12	61	MALSIFSAV	0.66/0.5519665
	13	91	WISYFVQSI	0.62/0.58502457
	14	171	KPNVLIALK	0.67/0.48465964
	15	163	LPNEVTVAK	0.63/0.51851449
	<b>16</b>	<b>93</b>	<b>SYFVQSIRL</b>	<b>0.01/1.1261978</b>
	17	124	GGTTVVRPL	0.67/0.39598193
	18	189	NSGVAIYHR	0.33/0.73296251
	19	175	LIALKMVKR	0.92/0.11586787
	20	191	GVAIYHRYK	0.02/1.0027127
	21	24	SLIFLLITI	0.06/0.9604213
	22	64	SIFSAVYPI	0.76/0.25040794
	23	76	SQIISGIVA	0.75/0.2054042
	24	95	FVQSIRLFM	0.29/0.64863165
	25	41	RSMTVYVFK	0.64/0.29572551
	26	19	WNFAWSLIF	0.99/-0.059008351
	27	188	TNSGVAIYH	0.15/0.7426577
	28	7	LTEAQIIAI	0.96/-0.075540835
	29	155	FGACDYDRL	0.99/-0.12423854
	30	159	DYDRLPNEV	0.14/0.69761321
	31	45	VYVFKMFVL	0.02/0.803342
	32	26	IFLLITIVL	0.02/0.7820036
	33	47	VFKMFVLWL	0.36/0.42789595
	34	200	AGNYRSPPI	0.84/-0.068816791
	35	209	TADIELALL	/0.76173539
	36	111	FPETNCLL	0.20/0.55943845
	37	86	VSAMMWISY	0.72/0.037550757
	38	21	FAWSLIFLL	0.11/0.61443046

39	37	GYPSRSMTV	0.05/0.65965946
40	5	TQLTEAQII	0.88/-0.17703249
41	8	TEAQIIAII	0.04/0.63874655
42	120	NVPFGGTTV	0.01/0.66645184
43	66	FSAVYPIDL	0.01/0.6514081
44	193	AIYHRYKAG	0.70/-0.038623293
45	154	HFGACDYDR	0.92/-0.26473748
46	110	SFNPETNCL	0.05/0.6027168
47	55	LLWPSSMAL	0.10/0.54332774
48	167	VTVAKPNVL	0.53/0.099630592
49	4	MTQLTEAQI	0.58/0.04746237
50	88	AMMWISYFV	0.25/0.37056163
51	196	HRYKAGNYR	0.15/0.46866787
52	202	NYRSPPIA	/0.60523431
53	184	QSYGTNSGV	0.82/-0.21985573
54	122	PFGGTTVVR	0.78/-0.19356706
55	152	GMHFGACDY	0.59/-0.010658696
56	116	NCLLNVPFG	0.53/0.024311399
57	57	WPSSMALSI	0.04/0.51276495
58	147	HLKMAGMHF	0.10/0.44982024
59	204	RSPPITADI	/0.53618799
60	121	VPFGGTTVV	0.06/0.46007594
61	168	TVAKPNVLI	0.01/0.49683601
62	62	ALSIFSAVY	0.00/0.50497952
63	208	ITADIELAL	/0.50382266
64	50	MFVLWLLWP	1.00/-0.49992406
65	13	IAIKDWNF	0.03/0.46314214
66	141	AVVTNGHLK	0.00/0.48952038
67	149	KMAGMHFGA	0.68/-0.20106563
68	14	AIKDWNFA	0.69/-0.21587273
69	169	VAKPNVLIA	0.06/0.4127837
70	42	SMTVYVFKM	0.00/0.47270089
71	17	KDWNFAWSL	0.00/0.41420002
72	143	VTNGHLKMA	0.68/-0.3130819
73	74	LASQIISGI	0.72/-0.39172178
74	99	IRLFMRTGS	0.85/-0.61346618
75	23	WSLIFLLIT	0.78/-0.55058213
76	54	WLLWPSSMA	0.77/-0.6485796
77	119	LNVPFGGTT	0.98/-0.88103088

	78	32	IVLQYGYP	0.95/-0.85753772
	79	127	TVVRPLVED	0.66/-0.58089465
	80	90	MWISYFVQS	0.81/-0.79029278
	81	112	NPETNCLLN	0.83/-0.81810302
	82	153	MHFGACDYD	0.53/-0.57221429
	83	157	ACDYDRLPN	0.93/-1.234207
	84	146	GHLKMAGMH	0.91/-1.2450366
	85	176	IALKMVKRQ	0.52/-0.98755935
	86	165	NEVTVAKPN	0.58/-1.1145372
	87	56	LWPSSMALS	0.75/-1.2934473
<b>HCoV-NL63</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	61	ALSIFDCFV	0.42/1.3269396
	2	27	TVFIVVLQY	0.96/0.75983923
	3	39	KYSRLLYGL	0.09/1.6269526
	4	194	GWAFYVRAK	0.61/1.1032724
	5	143	TLLSGVLLV	0.54/1.0866306
	6	72	NVDWVFFGF	0.93/0.6902797
	7	68	FVNFNVDWV	0.17/1.3164491
	8	103	WRRVKTFWA	0.99/0.41672656
	9	24	LILTVFIVV	0.74/0.62932871
	10	112	FNPETNAII	0.99/0.36572179
	11	23	NLILTVFIV	0.05/1.2803939
	12	163	GQLPKYVIV	0.28/1.0441602
	13	217	SEREKLLHL	/1.2478922
	14	134	MAAPTGVTL	0.01/1.2020559
	15	54	CLWPLVLAL	0.00/1.2025543
	16	15	LRNWNFSWN	0.91/0.29116747
	17	193	TGWAFYVRA	0.76/0.43350684
	18	43	LLYGLKMSV	0.99/0.19587974
	19	92	WVMYFVNSF	0.20/0.97214623
	20	50	SVLWCLWPL	0.66/0.50112474
	21	46	GLKMSVLWC	0.81/0.32713326
	22	41	SRLLYGLKM	0.58/0.55680081
	23	208	GVASQEGVL	0.88/0.23328951
	24	117	NAIISLQVY	0.79/0.32261677
	25	19	NFSWNLILT	0.95/0.13441361
	26	88	TLCLWVMYF	0.46/0.60965105

27	96	FVNSFRLWR	0.03/1.0157158
28	25	ILTVFIVVL	0.08/0.95244546
29	31	VVLQYGHYK	0.74/0.28403801
30	104	RRVKTFWAF	0.47/0.5453457
31	21	SWNLILTVF	0.76/0.21671489
32	191	SQTGWAFYV	0.00/0.95134123
33	93	VMYFVNSFR	0.79/0.12575145
34	76	VFFGFSILM	0.32/0.58884796
35	60	LALSIFDCF	0.70/0.20465324
36	101	RLWRRVKTF	0.01/0.89108233
37	47	LKMSVLWCL	1.00/-0.1038001
38	192	QTGWAFYVR	0.30/0.58530988
39	5	SVPLSEVYV	0.00/0.88332248
40	35	YGHYKYSRL	0.47/0.40854334
41	98	NSFRLWRRV	0.18/0.69216339
42	152	DGHKIATRV	0.93/-0.058796914
43	190	TSQTGWAFY	0.16/0.71048752
44	57	PLVLALSIF	0.96/-0.090172178
45	22	WNLILTVFI	0.60/0.26016217
46	97	VNSFRLWRR	0.19/0.66920512
47	51	VLWCLWPLV	0.00/0.85091173
48	81	SILMSIITL	0.06/0.78407937
49	7	PLSEVYVHL	0.33/0.50358441
50	10	EVYVHLRNW	0.76/0.06737552
<b>51</b>	<b>94</b>	<b>MYFVNSFRL</b>	<b>0.00/0.81397405</b>
52	16	RNWNFSWNL	0.00/0.78986837
53	12	YVHLRNWNF	0.02/0.75910697
54	95	YFVNSFRLW	0.93/-0.16742967
55	99	SFRLWRRVK	0.03/0.73100482
56	155	KIATRVQVG	0.79/-0.041900185
57	70	NFNVDWVFF	0.01/0.7324852
58	136	APTGVTLTL	0.13/0.61061023
59	4	SSVPLSEVY	0.96/-0.22283825
60	172	ATPSTTIVC	0.15/0.57706808
61	6	VPLSEVYVH	0.77/-0.04931276
62	128	NYYLPVMAA	0.30/0.41639443
63	89	LCLWVMYFV	0.01/0.69801811
64	175	STTIVCDRV	0.16/0.5326677
65	201	AKHGDFSGV	0.58/0.087165888



66	189	ETSQTGWAF	0.22/0.44474166
67	63	SIFDCFVNF	0.10/0.54780813
68	75	WVFFGFSIL	0.12/0.52189011
69	176	TTIVCDRVG	0.91/-0.27882271
70	67	CFVNFNVDW	0.66/-0.042369151
71	182	RVGRSVNET	0.64/-0.022524297
72	129	YYLPVMAAP	0.18/0.42472047
73	111	AFNPETNAI	0.01/0.57068859
74	73	VDWVFFGFS	0.95/-0.38416166
75	102	LWRRVKTFW	0.88/-0.33399253
76	79	GFSILMSII	0.14/0.40177837
77	44	LYGLKMSVL	0.05/0.48935835
78	36	GHYKYSRLL	0.60/-0.063998079
79	157	ATRVQVGQL	0.04/0.4959503
80	119	IISLQVYGH	0.78/-0.25341414
81	38	YKYSRLLYG	0.54/-0.038425177
82	20	FSWNLILTV	0.55/-0.055016499
83	87	ITLCLWVMY	0.00/0.49009803
84	71	FNVDWVFFG	0.00/0.47647061
85	123	QVYGHNYYL	0.02/0.43768443
86	32	VLQYGHYKY	0.76/-0.34582979
87	13	VHLRNWNFS	0.92/-0.5134316
88	211	SQEGVLSE	/0.40520158
89	82	ILMSIITLC	0.68/-0.33428279
90	108	TFWAFNPET	0.93/-0.58450234
91	174	PSTTIVCDR	0.64/-0.40569235
92	207	SGVASQEGV	0.87/-0.68502309
93	33	LQYGHYKYS	0.82/-0.70260336
94	114	PETNAIISL	0.57/-0.51338289
95	109	FWAFNPETN	0.52/-0.52429078
96	185	RSVNETSQT	0.87/-0.90286
97	11	VYVHLRNWN	0.51/-0.54535812
98	181	DRVGRSVNE	0.63/-0.71633178
99	184	GRSVNETSQ	0.83/-0.95171498
100	45	YGLKMSVLW	0.70/-0.84469286
101	133	VMAAPTGVT	0.69/-1.0697739
102	106	VKTFWAFNP	0.53/-1.0908489
103	138	TGVTLTLLS	0.78/-1.4291627
<b>HCoV-229E</b>			

<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	53	LLWPLVLAL	0.90/1.0172013
	2	207	AVSSPMSNM	0.65/1.1315979
	3	100	RLFRRARTF	0.69/1.0834146
	4	43	FYGLKMLVL	0.73/1.0416194
	5	62	SIFDTWANW	0.86/0.86647004
	6	69	NWDSNWAFV	0.77/0.91060155
	7	45	GLKMLVLWL	0.85/0.80022528
	8	38	KYSRLFYGL	0.07/1.5660843
	9	153	HRLASGVQV	0.98/0.59855888
	10	50	VLWLLWPLV	0.98/0.58891615
	11	24	ILTIFIVIL	0.06/1.4954772
	12	22	NVILTIFIV	0.74/0.8008092
	13	162	HNLPEYMTV	0.72/0.81596502
	14	178	IYSRVGRSV	0.57/0.93221588
	15	23	VILTIFIVI	0.98/0.49982791
	16	41	RLFYGLKML	0.93/0.5402551
	17	171	AVPSTTIY	0.48/0.93651537
	18	199	RVKHGDFSA	0.90/0.50488593
	19	40	SRLFYGLKM	0.69/0.7118252
	20	16	NWNFGWNVI	0.70/0.69004979
	21	142	TLLSGVLYV	0.03/1.3396451
	22	200	VKHGDFSAV	0.99/0.36784466
	23	103	RRARTFWAW	0.19/1.1631307
	24	71	DSNWAFVAF	0.02/1.1416897
	25	55	WPLVLALSI	0.98/0.15944072
	26	154	RLASGVQVH	0.67/0.4606884
	27	21	WNVILTIFI	0.82/0.30367738
	28	181	RVGRSVNSQ	0.98/0.14283932
	29	52	WLLWPLVLA	0.82/0.294418
	30	189	QNSTGWVIFY	0.76/0.32083315
	31	192	TGWVIFYVRV	0.31/0.76078895
	32	7	TGDIVTHLK	0.82/0.24707595
	33	92	VMYFANSFR	0.97/0.072860827
	34	133	QQAPTGITV	0.14/0.86735107
	35	98	SFRLFRRAR	0.12/0.87329753
	36	84	AVSTLVMWV	0.00/0.97860228
	37	177	IIYSRVGRS	0.98/-0.0067330453

38	216	TENERLLHF	/0.95511253
39	6	CTGDIVTHL	0.42/0.53178046
40	76	FVAFSFLMA	0.16/0.78893424
41	91	WVMYFANSF	0.19/0.73912126
42	93	<b>MYFANSFRL</b>	<b>0.23/0.69655934</b>
43	29	IVILQFGHY	0.95/-0.026529684
44	135	APTGITVTL	0.00/0.90960118
45	97	NSFRLFRA	0.06/0.8416029
46	96	ANSFRLFRR	0.01/0.88298762
47	47	KMLVLWLLW	0.87/0.021222742
48	121	TTVLGQTYT	0.80/0.079188078
49	156	ASGVQVHNL	0.25/0.62625185
50	26	TIFIVILQF	0.07/0.80272298
51	151	DGHRLASGV	0.90/-0.029760481
52	122	TVLGQTYTQ	0.87/-0.0033528359
53	112	NPEVNAITV	0.01/0.85169738
54	107	TFWAWNPEV	0.75/0.099752668
55	159	VQVHNLPEY	0.92/-0.091567921
56	95	FANSFRLFRR	0.07/0.75279398
57	115	VNAITVTTV	0.03/0.78849761
58	176	THYSRVGR	0.74/0.075212733
59	15	KNWNFGWNV	0.17/0.63687585
60	110	AWNPEVNAI	0.00/0.76448507
61	102	FRRARTFWA	0.00/0.73206915
62	214	NMTENERLL	/0.7228851
63	60	ALSIFDTWA	0.02/0.70106824
64	74	WAFVAFSFL	0.04/0.67644523
65	81	FLMAVSTLV	0.29/0.38369605
66	147	VLYVDGHR	0.56/0.087799519
67	99	FRLFRRART	0.25/0.394557
68	191	STGWVIFYVR	0.19/0.44185124
69	73	NWAFVAFSF	0.12/0.50684428
70	193	GWVIFYVRVK	0.01/0.61394467
71	30	VILQFGHYK	0.18/0.38876175
72	169	TVAVPSTTI	0.16/0.38178865
73	58	VLALSIFDT	0.60/-0.083623343
74	163	NLPEYMTVA	0.77/-0.27216171
75	188	SQNSTGWVF	0.04/0.45088961

	76	11	VTHLKNWNF	0.00/0.49021607
	77	68	ANWDSNWAF	0.00/0.48682911
	78	126	QTYYPPIQQ	0.77/-0.2954456
	79	87	TLVMWVMYF	0.01/0.46406701
	80	143	LLSGVLYVD	0.70/-0.25662713
	81	168	MTVAVPSTT	1.00/-0.56747993
	82	34	FGHYKYSRL	0.00/0.42878079
	83	139	ITVTLLSGV	0.71/-0.30838919
	84	184	RSVNSQNST	0.99/-0.60372976
	85	120	VTTVLGQTY	0.97/-0.6509589
	86	203	GDFSAVSSP	0.98/-0.70057856
	87	10	IVTHLKNWN	0.88/-0.60266118
	88	119	TVTTVLGQT	0.97/-0.75396264
	89	187	NSQNSTGWV	0.69/-0.52883165
	90	186	VNSQNSTGW	0.74/-0.5944038
	91	183	GRSVNSQNS	0.55/-0.43389174
	92	82	LMAVSTLVM	0.55/-0.44613325
	93	165	PEYMTVAVP	0.95/-0.91303044
	94	175	TTHYSRVG	0.92/-0.91126139
	95	123	VLGQTYYPQ	0.70/-0.69320026
	96	5	NCTGDIVTH	0.61/-0.68737295
	97	57	LVLALSIFD	0.80/-0.9154223
	98	118	ITVTTVLGQ	0.51/-0.81357846
	99	152	GHRLASGVQ	0.69/-1.0067898
	100	205	FSAVSSPMS	0.82/-1.1698711
	101	131	PIQQAPTGI	0.64/-1.2193975
	102	144	LSGVLYVDG	0.77/-1.4398635
<b>HCoV-OC43</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	204	KSKVGNYRL	0.87/1.0800295
	2	153	HLYIQGIKL	0.91/0.99884889
	3	101	FVNSIRLFI	0.72/1.0052845
	4	98	IVYFVNSIR	0.72/0.97294785
	5	88	VFTIVAIIIM	0.89/0.77527236
	6	202	YVKSKVGNY	0.01/1.5368575
	7	32	ILLFITHIL	0.82/0.67852814
	8	84	GLSIVFTIV	0.00/1.4977617
	9	195	DTSGFAVYV	0.01/1.4556836

10	196	TSGFAVYVK	0.82/0.64395629
11	117	FNPETNNLM	0.76/0.65010619
12	75	VYALNNVYL	0.08/1.3192874
<b>13</b>	<b>99</b>	<b>VYFVNSIRL</b>	<b>0.19/1.1841596</b>
14	90	TIVAIIMWI	0.67/0.687161
15	47	RSMFVYVIK	0.45/0.89755685
16	13	WTADEAIKF	0.43/0.91282428
17	168	ADLPAYMTV	0.65/0.68062847
18	26	NFSLGILL	0.52/0.75929494
19	76	YALNNVYLG	0.99/0.28536806
20	106	RLFIRTGSF	0.86/0.38885643
21	57	IILWLMWPL	0.78/0.45035621
22	51	VYVIKMIL	0.00/1.2206353
23	53	VIKMILWL	0.91/0.31035067
24	5	TTPAPVYIW	0.65/0.56357336
25	166	SLADLPAYM	0.25/0.93801696
26	28	SLGILLFI	0.32/0.86143284
27	185	YKRGFLDRI	0.87/0.30619036
28	139	IEDYHTLTV	0.83/0.3012014
29	109	IRTGSFWSF	0.58/0.54674803
30	121	TNNLMCIDM	0.54/0.5838765
31	91	IVAIIMWIV	0.00/1.1008378
32	87	IVFTIVAI	0.22/0.86440487
33	97	WIVYFVNSI	0.31/0.73599333
34	45	TSRSMFVYV	0.67/0.375493
35	41	QFGYTSRSM	0.85/0.16598633
36	61	LMWPLTHL	0.30/0.68843628
37	107	LFIRTGSFW	0.97/0.0054854584
38	4	KTTPAPVYI	0.49/0.45991837
39	46	SRSMFVYVI	0.33/0.61400396
40	116	SFNPETNNL	0.07/0.8711914
41	73	NCVYALNNV	0.19/0.73995041
42	48	SMFVYVIKM	0.18/0.7496604
43	147	VTIIRGHLY	0.63/0.29023927
44	141	DYHTLTVTI	0.14/0.76679025
45	44	YTSRSMFVY	0.52/0.38320157
46	50	FVYVIKMII	0.55/0.34820031
47	148	TIIRGHLYI	0.62/0.27756513
48	70	TIFNCVYAL	0.03/0.86500586

49	43	GYTSRSMFV	0.23/0.62312688
50	60	WLMWPLTII	0.26/0.5798484
51	86	SIVFTIVAI	0.33/0.50061561
52	128	DMKGTMYVR	0.52/0.31026034
53	81	VYLGLSIVF	0.70/0.12251951
54	63	WPLTIILTI	0.00/0.81833073
55	14	TADIAIKFL	0.23/0.58809773
56	77	ALNNVYLGL	0.01/0.80614231
57	80	NVYLGLSIV	0.02/0.78781589
58	222	DTALLRNNI	/0.74298675
59	131	GTMVVRPII	0.06/0.68201434
60	6	TPAPVYIWT	0.01/0.69695802
61	71	IFNCVYALN	0.91/-0.21001346
62	27	FSLGIILF	0.16/0.47991135
63	162	GTGYSLADL	0.00/0.62454384
64	171	PAYMTVAKV	0.59/0.029421572
65	89	FTIVAIIMW	0.01/0.59649437
66	102	VNSIRLFIR	0.00/0.60407438
67	16	DEAIKFLKE	0.60/-0.0035159999
68	83	LGLSIVFTI	0.08/0.51618789
69	156	IQIKLGTG	0.92/-0.32930791
70	67	IILTIFNCV	0.20/0.38686145
71	23	KEWNFSLGI	0.11/0.46300213
72	52	YVIKMILW	0.59/-0.02569042
73	33	LLFITIILQ	0.96/-0.39815528
74	198	GFAVYVKSK	0.01/0.54861816
75	30	GIILLFITI	0.00/0.55003067
76	170	LPAYMTVAK	0.90/-0.36265672
77	64	PLTIILTIF	0.74/-0.20273025
78	2	SSKTPAPV	0.10/0.43423301
79	79	NNVYLGLSI	0.90/-0.38064983
80	125	MCIDMKGTM	0.92/-0.40415557
81	165	YSLADLPAY	0.80/-0.28568776
82	113	SFWSFNPET	0.96/-0.44918176
83	93	AIIMWIVYF	0.01/0.49439466
84	39	ILQFGYTSR	0.51/-0.0063597186
85	94	IIMWIVYFV	0.00/0.49695658



	86	118	NPETNNLMC	0.99/-0.52216283
	87	82	YLGLSIVFT	0.99/-0.52451338
	88	25	WNFSLGIL	0.03/0.43323177
	89	21	FLKEWNFSL	0.05/0.38922126
	90	194	SDTSGFAVY	0.70/-0.26941699
	91	184	TYKRGFLDR	0.03/0.38589956
	92	186	KRGFLDRIS	0.97/-0.56233208
	93	209	NYRLPSTQK	0.00/0.39768487
	94	143	HTLTVTIIR	0.02/0.37679545
	95	167	LADLPAYMT	0.98/-0.58678549
	96	31	IILLFITII	0.01/0.38045256
	97	178	KVTHLCTYK	0.01/0.37415561
	98	103	NSIRLFIRT	0.00/0.37253474
	99	142	YHTLTVTII	0.60/-0.23463653
	100	172	AYMTVAKVT	0.76/-0.40286962
	101	12	IWTADEAIK	0.65/-0.41066048
	102	182	LCTYKRGFL	0.85/-0.6361775
	103	133	MYVRPIIED	0.61/-0.40004934
	104	37	TIILQFGYT	0.86/-0.67178448
	105	108	FIRTGSFWS	0.88/-0.72882034
	106	200	AVYVKSKVG	0.92/-0.82483856
	107	65	LTIILTIFN	0.98/-1.0027422
	108	138	IIEDYHTLT	0.95/-1.0002105
	109	150	IRGHLYIQG	0.65/-0.71544181
	110	145	LTVTIIRGH	0.84/-1.0417056
	111	7	PAPVYIWTA	0.53/-0.81669081
	112	197	SGFAVYVKS	0.62/-0.95789293
	113	40	LQFGYTSRS	0.54/-1.0393013
	114	124	LMCIDMKGT	0.63/-1.2106251
	115	29	LGILLFIT	0.60/-1.3935892
<b>HCoV-HKU1</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score (ANN/SVM)</b>
<b>Membrane</b>	1	161	TLSDLPVYV	0.82/1.6426512
	2	96	FVNSIRLFI	0.86/1.0052845
	3	75	NAFLAFSIV	0.96/0.88310432
	4	25	GVILLFITI	0.89/0.83050669
	5	111	SFNPETNNL	0.80/0.8711914
	6	163	SDLPVYVTV	0.59/1.0252376

7	112	FNPETNNLM	0.96/0.65010619
8	38	GYTSRSMFV	0.98/0.62312688
9	79	AFSIVFTII	0.73/0.86282911
10	136	DYHTLTATV	0.70/0.82891884
11	86	IISIVIWIL	0.06/1.4653727
12	1	MNKSFFPQF	0.91/0.54186681
13	72	ALNNAFLAF	0.88/0.56512118
14	41	SRSMFVYLI	0.81/0.61535157
15	14	ATFLKEWNF	0.16/1.2182794
16	21	NFSLGVILL	0.75/0.62512415
17	63	TLTIFNCFY	0.99/0.37497442
18	62	ITLTIFNCF	0.78/0.57908849
19	84	FTIISIVIW	0.60/0.74460738
20	104	IRTGSWWSF	0.87/0.41438112
21	70	FYALNNAFL	0.01/1.241901
22	43	SMFVYLIK M	0.68/0.57181132
23	197	FVKSKVGNY	0.00/1.21182
24	179	TYKRAFLDK	0.94/0.27057003
25	94	LYFVNSIRL	0.55/0.61934416
26	141	TATVIRGHL	0.85/0.31455757
27	99	SIRLFIRTG	0.99/0.16760057
28	74	NNAFLAFSI	0.36/0.78130362
29	189	DVNSGFAVF	0.03/1.1105628
30	148	HLYIQGVKL	0.24/0.89431328
31	85	TIISIVIW I	0.21/0.92006925
32	98	NSIRLFIRT	0.75/0.37253474
33	123	DMKGKMFVR	0.08/1.0423894
34	199	KSKVGNYRL	0.04/1.0800295
35	81	SIVFTIISI	0.18/0.86866348
36	65	TIFNCFYAL	0.00/0.99832674
37	190	VNSGFAVFV	0.03/0.91457699
38	46	VYLIKMIIL	0.00/0.92964441
39	69	CFYALNNAF	1.00/-0.071757212
40	83	VFTIISIVI	0.63/0.27647494
41	157	GTGYTLSDL	0.01/0.89257713
42	12	DQATFLKEW	0.84/0.055260038
43	23	SLGVILLFI	0.04/0.82424265
44	50	KMIILWLMW	0.93/-0.074516348
45	211	KPSGMDTAL	/0.8355206

46	102	LFIRTGSWW	0.69/0.13616864
47	82	IVFTIISIV	0.20/0.60767042
48	61	TITLTIFNC	0.55/0.2542018
49	8	QFTSDQATF	0.78/0.01411364
50	67	FNCFYALNN	0.92/-0.13202853
51	193	GFAVFKSK	0.00/0.78456363
52	45	FVYLIKMI	0.37/0.40929747
53	66	IFNCFYALN	0.82/-0.074265626
54	39	YTSRSMFVY	0.36/0.38320157
55	88	SIVIWILYF	0.09/0.65029146
56	154	VKLGTGYTL	0.92/-0.18144495
57	164	DLPVYVTVA	0.99/-0.25351136
58	92	WILYFVNSI	0.07/0.65480057
59	56	LMWPLTITL	0.07/0.65168812
60	89	IVIWILYFV	0.00/0.71849902
61	76	AFLAFSIVF	0.51/0.20705242
62	27	ILLFITIIL	0.01/0.67852814
63	9	FTSDQATFL	0.63/0.032709597
64	52	IILWLMWPL	0.20/0.45035621
<b>65</b>	<b>95</b>	<b>YFVNSIRLF</b>	<b>0.77/-0.13241026</b>
66	40	TSRSMFVYL	0.08/0.55538402
67	93	ILYFVNSIR	0.02/0.59878913
68	97	VNSIRLFIR	0.00/0.60407438
69	116	TNNLMCIDM	0.02/0.5838765
70	78	LAFSIVFTI	0.02/0.57733795
71	87	ISIVIWILY	0.21/0.38150171
72	22	FSLGVILLF	0.08/0.44594864
73	16	FLKEWNFSL	0.11/0.38922126
74	101	RLFIRTGSW	0.11/0.38242376
75	73	LNN AFLAFS	0.92/-0.42902267
76	149	LYIQGVKLG	0.67/-0.22461178
77	28	LLFITIILQ	0.80/-0.39815528
78	42	RSMFVYLIK	0.00/0.39421498
79	166	PVYVTVAKV	0.00/0.36871985
80	109	WWSFNPETN	0.77/-0.40375519
81	188	LDVNSGFAV	0.96/-0.61386228
82	127	KMFVRPVIE	0.64/-0.33198466
83	144	VIRGHLYIQ	0.81/-0.50687677
84	103	FIRTGSWWS	0.75/-0.45677222

	85	15	TFLKEWNFS	0.75/-0.47347113
	86	30	FITIILQFG	0.73/-0.46932787
	87	130	VRPVIEDYH	0.64/-0.38282906
	88	106	TGSWWSFNP	0.87/-0.64308731
	89	140	LTATVIRGH	0.94/-0.76560859
	90	182	RAFLDKLDV	0.89/-0.75765784
	91	131	RPVIEDYHT	0.68/-0.67948815
	92	108	SWWSFNPET	0.71/-0.72766196
	93	35	LQFGYTSRS	0.91/-1.0393013
	94	181	KRAFLDKLD	0.79/-0.93814167
	95	201	KVGNYRLPS	0.69/-0.89105159
	96	133	VIEDYHTLT	0.84/-1.2414406
	97	137	YHTLTATVI	0.56/-1.0434784
	98	158	TGYTLSDLP	0.55/-1.1778233
	99	156	LGTGYTLSD	0.58/-1.8489281
<b>SARS-CoV-2</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	262	RTATKAYNV	0.89/0.93090592
	2	222	LLDRLNQL	1.00/0.77917492
	3	85	GYRRTARR	0.96/0.64584436
	4	105	SPRWYFYLY	0.09/1.2901952
	5	165	TTLPKGFYA	0.84/0.47343878
	6	269	NVTQAFGR	0.99/0.29166166
	7	150	NPANNAIV	0.84/0.43927091
	8	266	KAYNVTQAF	0.83/0.43425876
	9	78	SSPDDQIGY	0.84/0.40764252
	10	215	GDAALALL	0.90/0.34336961
	11	211	AGNGGDAAL	0.87/0.36326844
	12	87	YRRATRRIR	0.36/0.86386896
	13	216	DAALALLL	0.19/1.026872
	14	153	NNAAIVLQL	0.70/0.43817133
	15	201	SSRGTSPAR	0.62/0.51098887
	16	131	IWVATEGAL	0.97/0.12227819
	17	226	RLNQLESKM	0.55/0.5287504
	18	138	ALNTPKDHI	0.49/0.54832454
	19	202	SRGTSPARM	0.19/0.84455909
	20	345	NFKDQVILL	0.04/0.93817097
	21	355	KHIDAYKTF	0.59/0.38618339

22	37	SKQRRPQGL	0.82/0.15310833
23	47	NNTASWFTA	0.99/-0.023318872
24	312	SAFFGMSRI	0.00/0.93379976
25	146	IGTRNPANN	0.95/-0.020760697
26	258	PRQKRTATK	0.93/-0.0085201677
27	81	DDQIGYYRR	0.60/0.31359257
28	86	YYRRATTRI	0.03/0.87920722
29	188	SRSRNSSRN	0.74/0.15874619
30	45	LPNNTASWF	0.87/0.026589893
31	323	EVTPSGTWL	0.50/0.38083972
32	186	SSSRNRNSS	0.95/-0.077380571
<b>33</b>	<b>125</b>	<b>ANKDGIWV</b>	<b>0.02/0.85111885</b>
34	166	TLPKGFYAE	0.89/-0.029340939
35	240	QQQGQTVTK	0.98/-0.12221078
36	124	GANKDGIW	0.55/0.29607236
37	9	QRNAPRITF	0.15/0.69591987
38	30	GERSGARSK	0.12/0.72256846
<b>39</b>	<b>311</b>	<b>ASAFFGMSR</b>	<b>0.02/0.78351091</b>
40	299	KHWPQIAQF	0.69/0.1045106
41	102	KDLSPRWYF	0.11/0.68097968
42	89	RATRRIRGG	0.85/-0.095326758
43	93	RIRGGDGKM	0.98/-0.24295114
<b>44</b>	<b>181</b>	<b>QASSRSSSR</b>	<b>0.32/0.40488402</b>
45	122	PYGANKDGI	0.96/-0.23589992
46	141	TPKDHIGTR	0.79/-0.070754546
47	96	GGDGKMKDL	0.05/0.65008429
48	68	RGQGVPI NT	0.83/-0.13269028
49	84	IGYYRRATR	0.01/0.66944734
50	338	KLDDKDPNF	0.01/0.66715268
51	2	SDNGPQNQR	0.64/0.036463719
52	53	FTALTQHGK	0.08/0.59448868
53	329	TWLTYTGAI	0.99/-0.31632651
54	213	NGGDAALAL	0.67/0.0028763949
55	48	NTASWFTAL	0.71/-0.042997835
56	173	AEGSRGGSQ	0.98/-0.34218131
57	7	QNQRNAPRI	0.14/0.48978888
58	306	QFAPSASAF	0.00/0.6216748
59	218	ALALLLLDR	0.19/0.42058446

60	210	MAGNGGDAA	0.82/-0.21169807
61	119	AGLPYGANK	0.21/0.38539776
62	148	TRNPANNAA	0.04/0.54723853
63	362	TFPPTDFSK	/0.55405184
64	175	GSRGGSQAS	0.96/-0.41600347
65	117	PEAGLPYGA	0.62/-0.093465736
66	88	RRATTRRIRG	0.01/0.51061059
67	314	FFGMSRIGM	0.04/0.44248594
68	368	FSKQLQQSM	/0.48014359
69	343	DPNFKDQVI	0.00/0.46907794
70	103	DLSRWYFY	0.00/0.45513017
71	289	QELIRQGT	0.96/-0.50787165
72	309	PSASAFFGM	0.53/-0.087196442
73	229	QLESKMSGK	0.79/-0.36665693
74	4	NGPQNQRNA	0.02/0.397257
75	270	VTQAFGRRG	0.82/-0.4093242
76	66	FPRGQGVPI	0.05/0.36004115
77	26	SNQNGERSG	0.91/-0.50282377
78	156	AIVLQLPQG	0.54/-0.14856364
79	50	ASWFTALTQ	0.71/-0.31887616
80	298	YKHWPQIAQ	0.94/-0.55888964
81	143	KDHIGTRNP	0.74/-0.35924171
82	205	TSPARMAGN	0.80/-0.46091766
83	349	QVILLNKHI	0.54/-0.23291994
84	253	EASKKPRQK	0.57/-0.2697544
85	246	VTKKSAAEA	0.88/-0.58975552
86	115	TGPEAGLPY	1.00/-0.72582022
87	256	KKPRQKRTA	0.55/-0.31949939
88	339	LDDKDPNFK	1.00/-0.79829376
89	217	AALALLLD	0.82/-0.66387806
90	12	APRITFGGP	0.64/-0.51055721
91	320	IGMEVTPSG	0.93/-0.80199944
92	268	YNVTQAFGR	0.66/-0.54408881
93	233	KMSGKGQQQ	0.66/-0.5681714
94	301	WPQIAQFAP	0.86/-0.7731242
95	260	QKRTATKAY	0.82/-0.73406697
96	305	AQFAPSASA	0.68/-0.63194405
97	244	QTVTKKSAA	0.71/-0.6637588
98	22	DSTGSNQNG	0.97/-0.92411209

	99	3	DNGPQNQRN	0.97/-0.92525482
	100	231	ESKMSGKGQ	0.85/-0.84680824
	101	107	RWYFYYLGT	0.57/-0.57892917
	102	60	GKEDLKFP	0.83/-0.85866559
	103	332	TYTGAIKLD	0.96/-1.0332853
	104	139	LNTPKDHIG	0.68/-0.76483333
	105	197	STPGSSRGT	0.96/-1.0609933
	106	278	GPEQTQGNF	0.56/-0.74934502
	107	17	FGGPSDSTG	0.85/-1.0477316
	108	137	GALNTPKDH	0.94/-1.1694863
	109	157	IVLQLPQGT	0.65/-0.94266092
	110	97	GDGKMKDLS	0.79/-1.096834
	111	279	PEQTQGNFG	0.91/-1.2228358
	112	198	TPGSSRGTS	0.65/-0.9769109
	113	16	TFGGPSDST	0.88/-1.2239998
	114	239	QQQQGQTVT	0.59/-0.95994625
	115	340	DDKDPNFKD	0.52/-0.90141856
	116	158	VLQLPQGTT	0.91/-1.2940397
	117	333	YTGAIKLDD	0.67/-1.0851174
	118	247	TKKSAAEAS	0.78/-1.2140661
	<b>119</b>	<b>72</b>	<b>VPINTNSSP</b>	<b>0.55/-1.0398301</b>
	120	294	QGTDYKHWP	0.73/-1.2211872
	121	335	GAIKLDDKD	0.94/-1.4612102
	122	302	PQIAQFAPS	0.61/-1.2992434
	123	168	PKGFYAEGS	0.80/-1.7322271
<b>SARS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	87	YYRRATRRV	0.96/1.0113912
	2	212	ASGGGETAL	0.91/0.90898954
	<b>3</b>	<b>106</b>	<b>SPRWYFYYL</b>	<b>0.44/1.2901952</b>
	4	103	KELSPRWYF	0.56/1.1574925
	5	203	SRGNSPARM	0.96/0.68383146
	6	85	IGYYRRATR	0.94/0.66944734
	<b>7</b>	<b>151</b>	<b>NPNNNAATV</b>	<b>0.90/0.64775325</b>
	8	346	QFKDNVILL	0.20/1.2729819
	9	166	TTLPKGFYA	0.90/0.47343878
	10	315	FFGMSRIGM	0.91/0.44248594
	<b>11</b>	<b>312</b>	<b>ASAFFGMSR</b>	<b>0.55/0.78351091</b>

12	255	ASKKPRQKR	0.98/0.35330623
13	104	ELSPRWYFY	0.70/0.63184645
<b>14</b>	<b>182</b>	<b>QASSRSSSR</b>	<b>0.79/0.40488402</b>
15	313	SAFFGMSRI	0.23/0.93379976
16	149	TRNPNNNAA	0.06/1.0992146
17	381	QPLPQRQKK	0.81/0.331546
18	348	KDNVILLNK	0.73/0.4042565
19	217	ETALALLLL	0.00/1.1212241
20	242	QQGQTVTKK	0.95/0.15472685
21	344	DPQFKDNVI	0.93/0.15510863
22	97	GGDGKMKEL	0.04/1.0139312
23	216	GETALALLL	0.79/0.25443743
24	332	LTYHGAIKL	0.63/0.40861273
25	308	FAPSASAFF	0.68/0.3213651
26	101	KMKELSPRW	1.00/-0.021813224
27	7	QSNQRSAPR	0.35/0.62090318
28	329	GTWLTYHGA	0.80/0.15791305
29	62	KEELRFPRG	0.96/-0.045242636
30	120	ASLPYGANK	0.08/0.82674523
31	183	ASSRSSRS	1.00/-0.10019775
32	65	LRFPRGQGV	0.66/0.23783152
33	353	LLNKHIDAY	0.85/0.021288078
34	80	GPDDQIGYY	0.84/0.025720532
35	227	RLNQLESKV	0.00/0.86225984
<b>36</b>	<b>223</b>	<b>LLDRLNQL</b>	<b>0.07/0.77917492</b>
37	362	KTFPTEPK	0.60/0.24733102
38	128	KEGIVWVAT	0.59/0.24617659
39	263	RTATKQYNV	0.00/0.82832214
40	10	QRSAPRITF	0.09/0.73651318
41	307	QFAPSASAF	0.19/0.6216748
42	380	AQPLPQRQK	0.23/0.5753709
43	350	NVILLNKHI	0.98/-0.19546084
44	267	KQYNVTQAF	0.18/0.58266537
45	142	TPKDHIGTR	0.83/-0.070754546
46	202	SSRGNSPAR	0.09/0.66714286
<b>47</b>	<b>126</b>	<b>ANKEGIVWV</b>	<b>0.20/0.55428711</b>
48	160	LQLPQGTTL	0.98/-0.23496764
49	205	GNSPARMAS	0.99/-0.24525216
50	152	PNNNAATVL	0.97/-0.25317817



51	26	DNNQNGGRN	0.97/-0.27184383
<b>52</b>	<b>86</b>	<b>GYRRATTR</b>	<b>0.05/0.64584436</b>
53	400	DMDDFSRQL	/0.6943173
54	33	RNGARPKQR	0.97/-0.28343228
55	88	YRRATTRVR	0.00/0.68621587
56	272	TQAFGRRGP	0.89/-0.20410199
57	260	RQKRTATKQ	0.98/-0.3004523
58	249	KKSAAEASK	0.93/-0.27044377
59	139	ALNTPKDHI	0.11/0.54832454
60	154	NNAATVLQL	0.06/0.5857138
61	8	SNQRSAPRI	0.16/0.4740182
62	360	AYKTFPTE	0.52/0.11141851
63	35	GARPKQRRP	0.83/-0.20998679
64	29	QNGGRNGAR	0.96/-0.36319524
65	54	FTALTQHGK	0.00/0.59448868
66	9	NQRSAPRIT	0.75/-0.16130801
67	292	LIRQGTDYK	0.51/0.072620614
68	42	RPQGLPNNT	0.81/-0.24416525
69	339	KLDDKDPQF	0.10/0.45884604
70	136	TEGALNTPK	0.72/-0.16914112
71	57	LTQHGKEEL	0.58/-0.042653612
72	186	RSSSRSGN	0.90/-0.37107981
73	90	RATRRVRGG	0.10/0.42004926
74	207	SPARMASGG	0.94/-0.42016665
75	286	NFGDQDLIR	0.60/-0.083027619
76	201	GSSRGNSPA	0.64/-0.1233026
77	220	LALLLLDRL	0.62/-0.11574675
78	56	ALTQHGKEE	0.94/-0.44868483
79	84	QIGYYRRAT	0.91/-0.42881926
80	363	TFPTEPKK	0.03/0.4507155
81	112	YYLGTGPEA	0.55/-0.077572032
82	294	RQGTDYKHW	0.95/-0.48354644
83	324	EVTSGTWL	0.08/0.38083972
84	296	GTDYKHWPQ	0.85/-0.38989884
85	146	HIGTRNPNN	0.99/-0.54574633
86	176	GSRGGSQAS	0.86/-0.41600347
87	404	FSRQLQNSM	/0.43846516
88	219	ALALLLLDR	0.01/0.42058446
89	214	GGGETALAL	0.02/0.40188538

90	11	RSAPRITFG	0.02/0.39113805
91	2	SDNGPQSNQ	0.99/-0.58342809
92	356	KHIDAYKTF	0.00/0.38618339
93	3	DNGPQSNQR	0.89/-0.50840327
94	67	FPRGQGVPI	0.02/0.36004115
95	63	EELRFPRGQ	1.00/-0.63332203
96	52	SWFTALTQH	0.99/-0.62969617
97	293	IRQGTDYKH	0.87/-0.5127469
98	175	EGSRGGSQA	0.55/-0.19442815
99	210	RMASGGGET	0.88/-0.53626227
100	316	FGMSRIGME	0.76/-0.43050044
101	121	SLPYGANKE	0.59/-0.28571486
102	92	TRRVRGGDG	0.57/-0.27342213
103	129	EGIVWVATE	0.63/-0.34245656
104	165	GTTLPKGFI	0.96/-0.68259645
105	269	YNVTQAFGR	0.82/-0.54408881
106	274	AFGRRGPEQ	0.96/-0.69079218
107	235	VSGKGQQQQ	0.93/-0.6682894
108	24	STDNNQNGG	0.66/-0.40356691
109	190	RSRGNSRNS	0.52/-0.28449689
110	291	DLIRQGTDY	0.66/-0.42705044
111	199	TPGSSRGNS	0.87/-0.69928417
112	351	VILLNKHID	0.90/-0.76166859
113	20	GPTDSTDNN	0.75/-0.62044357
114	91	ATRRVRGGD	0.55/-0.4368316
115	247	VTKKSAAEA	0.70/-0.58975552
116	40	QRRPQGLPN	0.96/-0.85617741
117	390	QPTVTLLPA	0.58/-0.50072892
118	114	LGTGPEASL	0.87/-0.80322609
119	43	PQGLPNNTA	0.97/-0.93230078
120	22	TDSTDNNQN	0.84/-0.81465668
121	252	AAEASKKPR	0.76/-0.74082171
122	229	NQLESKVSG	0.91/-0.90777161
123	158	TVLQLPQGT	0.91/-0.9130794
124	302	WPQIAQFAP	0.77/-0.7731242
125	240	QQQQGQTVT	0.93/-0.95994625
126	78	NSGPDDQIG	0.52/-0.57410796
127	47	PNNTASWFT	0.83/-0.8916636
128	66	RFPRGQGVF	0.85/-0.92044246

	129	232	ESKVSGKGQ	0.68/-0.80324094
	130	233	SKVSGKGQQ	0.98/-1.1173681
	131	283	TQGNFGDQD	0.64/-0.8435021
	132	342	DKDPQFKDN	0.69/-0.90895165
	133	277	RRGPEQTQG	0.99/-1.216101
	134	295	QGTDYKHWP	0.96/-1.2211872
	135	382	PLPQRQKKQ	0.94/-1.2017533
	136	200	PGSSRGNSP	0.93/-1.1938617
	137	282	QTQGNFGDQ	0.99/-1.2588899
	138	320	RIGMEVTPS	0.99/-1.2663348
	139	377	TDEAQPLPQ	0.97/-1.3052823
	140	141	NTPKDHIGT	0.56/-0.95695803
	141	244	GQTVTKKSA	0.66/-1.143769
	142	238	KGQQQQGQT	0.95/-1.4372311
	143	322	GMEVTPSGT	0.71/-1.2057264
	<b>144</b>	<b>73</b>	<b>VPINTNSGP</b>	<b>0.67/-1.1760564</b>
	145	180	GSQASSRSS	0.75/-1.2942226
	146	6	PQSNQRSAP	0.87/-1.4717376
	147	198	STPGSSRGN	0.77/-1.3867119
	148	60	HGKEELRFP	0.59/-1.2712352
	149	74	PINTNSGPD	0.69/-1.3928335
	150	143	PKDHIGTRN	0.98/-1.6828756
	151	395	LLPAADMDD	0.60/-1.3249575
	152	173	YAEGSRGGS	0.98/-1.804953
	153	163	PQGTTLPKG	0.75/-1.7411809
<b>MERS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	326	VYFLRYSGA	0.82/0.85173683
	2	288	TEDPRWPQI	0.60/1.020664
	3	329	LRYSGAIKL	0.36/1.2090184
	<b>4</b>	<b>214</b>	<b>LYLDLLNRL</b>	<b>0.61/0.93922265</b>
	5	342	PNYNKWLEL	1.00/0.47691486
	<b>6</b>	<b>170</b>	<b>QSSSRASSV</b>	<b>0.92/0.48670336</b>
	7	92	KQLAPRWYF	0.55/0.78160836
	8	15	NNDITNTNL	0.87/0.4309186
	<b>9</b>	<b>115</b>	<b>AVKDGIVWV</b>	<b>0.01/1.2876187</b>
	10	347	WLELLEQNI	0.97/0.30634976
	11	35	AAPNNTVSW	0.96/0.26512505

12	343	NYNKWLELL	0.57/0.63619978
13	28	GRNPKPRAA	0.00/1.1641891
14	211	GDLLYLDLL	0.90/0.25642104
15	353	QNIDAYKTF	0.98/0.16272627
16	155	KLPKNFHIE	0.82/0.30335762
17	233	SQPKVITKK	0.79/0.32852037
18	245	AAKNKMRHK	0.51/0.60005073
19	109	AALPFRAVK	0.29/0.79870845
20	82	DRKINTGNG	0.78/0.29116261
21	327	YFLRYSGAI	0.70/0.37093526
22	72	AQNAGYWRR	0.39/0.67261704
23	130	DAPSTFGTR	0.51/0.53477132
24	173	SRASSVSRN	0.99/0.023642934
25	210	GGDLLYLDL	0.94/0.065766746
26	90	GIKQLAPRW	0.96/0.017562599
27	208	AVGGDLLYL	0.05/0.92649247
<b>28</b>	<b>95</b>	<b>APRWYFYT</b>	<b>0.60/0.37406648</b>
29	246	AKNKMRRHK	0.99/-0.027664866
30	143	DSAIVTQFA	0.49/0.46452033
31	198	GTSPGPSGI	0.98/-0.034719191
32	380	SEPPKEQRV	0.18/0.7439029
33	356	DAYKTFPKK	0.84/0.082428678
34	305	AFMGMSQFK	0.69/0.22322831
35	250	MRHKRTSTK	0.00/0.90434598
36	149	QFAPGTKLP	0.98/-0.084491668
37	150	FAPGTKLPK	0.97/-0.084591696
<b>38</b>	<b>139</b>	<b>NPNNSAIV</b>	<b>0.69/0.19349234</b>
39	18	ITNTNLSRG	0.96/-0.10557402
40	221	RLQALESBK	0.97/-0.15275972
41	322	HGNPVYFLR	0.33/0.47551321
42	350	LLEQNIDAY	0.98/-0.17770976
43	217	DLLNRLQAL	0.00/0.7886719
44	154	TKLPKNFHI	0.33/0.43450264
<b>45</b>	<b>303</b>	<b>ASAFMGMSQ</b>	<b>0.95/-0.1961539</b>
46	183	SRSSSQGSR	0.74/-0.017063041
47	328	FLRYSGAIK	0.00/0.71581396
48	33	PRAAPNNTV	0.91/-0.19875788
49	200	SPGPSGIGA	0.30/0.39825848
50	176	SSVSRNSSR	0.31/0.38038282

51	148	TQFAPGTKL	0.00/0.68511766
52	330	RYSGAIKLD	1.00/-0.32297433
53	291	PRWPQIAEL	0.16/0.51109243
54	36	APNNTVSWY	0.00/0.66729812
55	318	NNDDHGPNV	0.53/0.13346147
56	80	RQDRKINTG	0.59/0.053859252
57	57	FPPGQGVPL	0.14/0.45757756
58	2	ASPAAPRAV	0.02/0.57689358
59	39	NTVSWYTGL	0.83/-0.2415175
60	24	SRGRGRNPK	0.06/0.5133712
61	1	MASPAAPRA	0.52/0.042181518
62	249	KMRHKRTST	0.97/-0.40798952
63	259	SFNMVQAFG	0.70/-0.1449044
64	384	KEQRVQGS	0.00/0.55401551
65	306	FMGMSQFKL	0.87/-0.3252675
66	295	QIAELAPTA	0.59/-0.053972092
67	394	QRTRTRPSV	/0.52141346
68	60	GQGVPLNAN	0.74/-0.23063099
<b>69</b>	<b>76</b>	<b>GYWRRQDRK</b>	<b>0.00/0.50733206</b>
70	52	KVPLTFPPG	0.88/-0.37641247
71	107	PEAALPFRA	0.93/-0.43063521
72	77	YWRRQDRKI	0.72/-0.22493933
73	43	WYTGLTQHG	0.62/-0.12625376
74	340	KNPNYNKWL	0.67/-0.17807239
75	338	DPKNPNYNK	0.00/0.48988479
76	137	TRNPNNDSA	0.09/0.38771361
77	213	LLYLDLLNR	0.02/0.43876131
78	349	ELLEQNIDA	0.92/-0.4646237
79	320	DDHGPNVYF	0.08/0.37157273
80	304	SAFMGMSQF	0.03/0.40607493
81	27	RGRNPKPRA	0.84/-0.41637797
82	84	KINTGNGIK	0.00/0.41259831
83	402	VQPGPMIDV	/0.41215303
84	242	DAAAAKNKM	0.73/-0.32337359
85	75	AGYWRRQDR	0.01/0.39004073
86	106	GPEAALPFR	0.93/-0.53419678
87	113	FRAVKDGIV	0.00/0.39224958
88	292	RWPQIAELA	0.97/-0.60231607
89	232	QSQPKVITK	0.00/0.36478831

90	23	LSRGRGRNP	0.75/-0.40775681
91	360	TFPKKEKKQ	0.92/-0.59649026
92	83	RKINTGNGI	0.95/-0.62975784
93	372	KEESTDQMS	0.82/-0.50634084
94	201	PGPSGIGAV	0.59/-0.28998395
95	252	HKRTSTKSF	0.97/-0.67059799
96	117	KDGIVWVHE	0.51/-0.2218636
97	332	SGAIKLDPK	0.95/-0.69689508
98	185	SSSQGSRSG	0.56/-0.31672703
99	247	KNKMRHKRT	0.52/-0.28027281
100	37	PNNTVSWYT	0.88/-0.65141131
101	42	SWYTGLTQH	0.82/-0.62385673
102	122	WVHEDGATD	0.72/-0.52644832
103	319	NDDHGNPVY	0.61/-0.42396956
104	263	VQAFGLRGP	0.74/-0.61483983
105	123	VHEDGATDA	0.80/-0.67587854
106	227	SGKVKQSQP	0.99/-0.91028653
107	194	NSTRGTSPG	0.96/-0.88518292
108	98	WYFYTGTTG	0.92/-0.88490721
109	224	ALESGKVKQ	1.00/-0.97315219
110	223	QALESGKVK	0.79/-0.78521188
111	310	SQFKLTHQN	0.81/-0.82783131
112	317	QNNDHGNP	0.77/-0.79513552
113	192	SGNSTRGTS	0.96/-1.0468348
114	203	PSGIGAVGG	0.74/-0.82836864
115	212	DLLYDLLN	0.55/-0.65856866
116	13	ADNNDITNT	0.54/-0.6704073
117	280	DLQLNKLGT	0.77/-0.94074599
118	231	KQSQPKVIT	0.51/-0.68078812
119	40	TVSWYTGLT	0.84/-1.0114911
120	168	NSQSSSRAS	0.67/-0.88895741
121	269	RGPGDLQGN	0.92/-1.1405124
122	335	IKLDPKNPN	0.85/-1.0728028
123	365	EKKQKAPKE	0.79/-1.014063
124	309	MSQFKLTHQ	0.62/-0.85540747
125	16	NDITNTNLS	0.97/-1.2251622
126	222	LQALESGKV	0.78/-1.082446
127	46	GLTQHKGVP	0.95/-1.2622989
128	331	YSGAIKLDP	0.94/-1.2708713

	129	119	GIVVWHEDG	0.60/-0.98102059
	130	363	KKEKKQKAP	0.93/-1.3267215
	131	7	PRAVSFADN	0.65/-1.0473359
	132	271	PGDLQGNFG	0.62/-1.0298644
	133	333	GAIKLDPKN	0.85/-1.2743699
	134	237	VITKKDAAA	0.89/-1.3561122
	135	196	TRGTSPGPS	0.60/-1.0713156
	136	65	LNANSTPAQ	0.66/-1.1731177
	137	375	STDQMSEPP	0.86/-1.5075913
	138	204	SGIGAVGGD	0.93/-1.5873422
	139	121	VVWHEDGAT	0.60/-1.2582851
	140	366	KKQKAPKEE	0.53/-1.2082515
	<b>141</b>	<b>63</b>	<b>VPLNANSTP</b>	<b>0.70/-1.4579462</b>
<b>HCoV-NL63</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	93	QRSDGVVWV	0.83/1.0503585
	2	27	VSSDKAPYR	0.89/0.85441578
	3	153	RSSTRNNSR	0.98/0.53378298
	4	297	EVGDNVQIT	0.81/0.65935549
	5	64	MRRGQRVDL	0.64/0.82563036
	6	38	PRNLVPIGK	0.99/0.3316211
	7	251	GPRDFNHNM	0.07/1.2149976
	8	7	ADDRAARKK	0.95/0.33291945
	9	167	TSRQQSRTR	0.59/0.6626187
	10	33	PYRVIPRNL	0.89/0.34361091
	11	32	APYRVIPRN	0.90/0.31070761
	12	181	SSSDLVAAV	0.22/0.98207037
	13	127	FSIALPPEL	0.96/0.22155007
	14	166	STSRQQSRT	0.91/0.26624856
	15	272	KGFPQLAEL	0.63/0.48881282
	16	61	RWRMRRGQR	0.88/0.237016
	17	53	IGYWNVQER	0.31/0.788897
	18	279	ELIPNQAAL	0.00/1.0575762
	19	296	DEVGDNVQI	0.88/0.17056271
	20	278	AELIPNQAA	0.79/0.25853901
	21	94	RSDGVVWVA	0.16/0.88619516
	22	6	WADDRAARK	0.00/1.0073102
	23	238	RVPTREENV	0.00/0.99866868

24	327	SAFTKPSSI	0.70/0.26711755
25	247	IQCFGPRDF	0.43/0.52897559
26	321	KFIEQISAF	0.25/0.70443494
27	47	GNKDEQIGY	0.57/0.37815546
28	230	QLKKPRWKR	0.21/0.73360993
29	62	WRMRRGQRV	0.00/0.91669726
30	3	SVNWADDRA	0.63/0.28523363
31	50	DEQIGYWNV	0.00/0.89996169
32	29	SDKAPYRVI	0.67/0.22325403
33	281	IPNQAALFF	0.02/0.87146481
34	188	AVTLALKNL	0.04/0.84887309
35	195	NLGFDNQSK	0.90/-0.022352029
36	95	SDGVVWVAK	0.52/0.33841478
37	318	NLPKFIEQI	0.04/0.79643602
38	243	EENVIQCFG	0.96/-0.12436364
39	70	VDLPPKVHF	0.89/-0.066727275
40	367	AIIEIVNEV	/0.82051619
41	245	NVIQCFGPR	0.59/0.2096198
42	68	QRVDLPPKV	0.72/0.078024545
43	264	LVQNGVDAK	0.76/0.028387027
44	294	STDEVGDNV	0.00/0.78810974
45	58	VQERWRMRR	0.60/0.17273593
46	28	SSDKAPYRV	0.08/0.6794857
47	57	NVQERWRMR	0.02/0.7372599
48	224	RADKPSQLK	0.20/0.55370903
49	330	TKPSSIKEM	0.86/-0.10663569
50	146	NSSRASSRS	0.74/0.0073732169
51	36	VIPRNLVPI	0.94/-0.19830235
52	129	IALPPELSV	0.64/0.094165838
53	145	NNSSRASSR	0.00/0.72792832
54	149	RASSRSSTR	0.01/0.71577376
55	314	KDNKNLPKF	0.75/-0.029560779
56	183	SDLVAAVTL	0.95/-0.23214497
57	269	VDAKGFPQL	0.67/0.047815037
58	123	LEPKFSIAL	0.22/0.49253334
59	302	VQITYTYKM	0.98/-0.27260668
60	121	KPLEPKFSI	0.29/0.40952758
61	165	RSTSRQQSR	0.01/0.65243185
62	56	WNVQERWRM	0.03/0.60837944



63	265	VQNGVDAKG	1.00/-0.3761421
64	105	GAKTVNTSL	0.03/0.58121056
65	180	QSSSDLVAA	0.77/-0.17987904
66	157	RNNSRDSSR	0.07/0.51667575
67	177	DSNQSSSDL	0.14/0.43242551
68	159	NSRDSSRST	0.56/-0.0050559884
69	108	TVNTSLGNR	0.01/0.54170818
70	133	PELSVVEFE	0.93/-0.39332569
71	334	SIKEMQSQS	0.98/-0.44833146
72	48	NKDEQIGYW	0.58/-0.059508967
73	79	YYLGTGPHK	0.02/0.49258046
74	15	KFPPPSFYM	0.06/0.44951581
75	342	SSHVVQNTV	0.04/0.45734456
76	114	GNRKRNQKP	0.96/-0.46925538
77	112	SLGNRKRNQ	0.93/-0.4486876
78	301	NVQITYTYK	0.00/0.47327988
79	241	TREENVIQC	0.00/0.46984244
80	63	RMRRGQRVD	0.94/-0.47648794
81	368	IIEIVNEVL	/0.45993555
82	347	QNTVLNASI	0.98/-0.52255007
83	252	PRDFNHNMG	0.72/-0.27256097
84	323	IEQISAF TK	0.07/0.37570906
85	115	NRKRNQKPL	0.00/0.40656426
86	242	REENVIQCF	0.01/0.3949619
<b>87</b>	<b>73</b>	<b>PPKVHFYYL</b>	<b>0.03/0.36963986</b>
88	59	QERWRMRRG	0.01/0.38920753
89	162	DSSRSTSRQ	0.52/-0.15910388
90	324	EQISAF TKP	0.94/-0.58828055
91	336	KEMQSQSSH	0.59/-0.2934749
92	109	VNTSLGNRK	0.75/-0.45733365
93	80	YLGTGPHKD	0.98/-0.70409094
94	174	TRSDSNQSS	0.69/-0.42839095
95	348	NTVLNASIP	0.67/-0.41687283
96	307	TYKMLVAKD	0.90/-0.65446786
97	26	LVSSDKAPY	0.89/-0.65067805
98	256	NHNMGDSDL	0.52/-0.29159951
99	136	SVVEFEDRS	0.59/-0.36166442
100	309	KMLVAKDNK	0.92/-0.69753357

101	92	RQRSDGVVW	0.55/-0.33061775
102	134	ELSVVEFED	0.70/-0.4834477
103	350	VLNASIPES	0.62/-0.41936453
104	190	TLALKNLGF	0.76/-0.57306937
105	122	PLEPKFSIA	0.83/-0.64468962
106	103	KEGAKTVNT	0.72/-0.54231507
107	72	LPPKVHFYY	0.69/-0.52054071
108	144	SNNSSRASS	0.61/-0.46571218
109	333	SSIKEMQSQ	0.94/-0.82165389
110	99	VWVAKEGAK	0.75/-0.64096712
111	311	LVAKDNKNL	0.65/-0.56632874
112	312	VAKDNKNLP	0.59/-0.50910295
113	106	AKTVNTSLG	0.65/-0.58682727
114	193	LKNLGFDNQ	0.62/-0.5680716
115	258	NMGDSDLVQ	0.84/-0.78932
116	204	SPSSSGTST	0.57/-0.55455441
117	178	SNQSSSDLV	0.72/-0.72102631
118	89	LKFRQRSDG	0.94/-0.97484454
119	274	FPQLAELIP	0.89/-0.93373417
120	280	LIPNQAALF	0.55/-0.60569392
121	332	PSSIKEMQS	0.92/-1.0260017
122	40	NLVPIGKGN	0.99/-1.1594635
123	19	PSFYMPLLV	0.53/-0.7327578
124	352	NASIPESKP	0.53/-0.77936507
125	196	LGFDNQSKS	0.93/-1.1827807
126	49	KDEQIGYWN	0.93/-1.1991049
127	207	SSGTSTPKK	0.56/-0.83177295
128	155	STRNNSRDS	0.64/-1.0013112
129	213	PKKPNKPLS	0.94/-1.3310662
130	10	RAARKKFPP	0.81/-1.2379601
131	197	GFDNQSKSP	0.62/-1.0696074
132	313	AKDNKNLPK	0.67/-1.1337899
133	52	QIGYWNVQE	0.81/-1.2762632
134	217	NKPLSQPRA	0.62/-1.1000601
135	77	HFYYLGTGP	0.92/-1.4174491
136	232	KKPRWKRPV	0.74/-1.3135443
137	222	QPRADKPSQ	0.62/-1.2730379
138	176	SDSNQSSSD	0.87/-1.5739187
139	308	YKMLVAKDN	0.81/-1.5219762

	140	275	PQLAELIPN	0.75/-1.5298008
	141	102	AKEGAKTVN	0.73/-1.5281956
	142	208	SGTSTPKKP	0.61/-1.4323565
	143	335	IKEMQSQSS	0.72/-1.6078258
	144	24	PLLVSSDKA	0.63/-1.8420451
	145	262	SDLVQNGVD	0.67/-1.9171155
<b>HCoV-229E</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	96	RVEGVVWVA	0.99/0.91622308
	2	333	HLGKFLEEL	0.77/1.0331021
	3	16	RQGRIPYSL	0.95/0.69755833
	4	131	QKLPNGVTV	0.93/0.65267196
	5	340	ELNAFTREM	0.90/0.61756135
	6	146	RAPSRSQSR	0.60/0.90146006
	7	142	EPDSRAPSR	0.65/0.82474115
	8	40	PRNLVPINK	0.88/0.54735297
	9	60	VQKRFRTRK	0.91/0.49418632
	<b>10</b>	<b>75</b>	<b>SPKLHFYYL</b>	<b>0.15/1.2036463</b>
	11	125	EIPHFNQKL	1.00/0.34615981
	12	55	IGYWNVQKR	0.95/0.3801915
	13	49	KDKNKLIGY	0.59/0.64992196
	14	97	VEGVVWVAV	0.14/1.0911548
	15	301	AMLFDSHIV	0.90/0.32027775
	16	178	SQDDIMKAV	0.18/1.0281547
	17	316	TVVLTFTTR	0.55/0.63669774
	18	262	TQCFGPRDL	0.04/1.1314081
	19	182	IMKAVAAAL	0.70/0.46065499
	20	295	LVPSTAAML	0.91/0.24603278
	21	66	TRKGKRVDL	0.00/1.0673474
	22	70	KRVDLSPKL	0.98/0.086905042
	23	12	PQRGRQGRI	0.93/0.11614714
	24	288	GYPQFAELV	0.38/0.66351077
	25	84	GTGPHKDAK	0.91/0.042910867
	26	311	KESGNTVVL	0.10/0.83520062
	27	73	DLSPKLHFY	0.36/0.57360838
	28	57	YWNVQKRFR	0.97/-0.040738699
	29	242	HEMQKPRWK	0.92/0.0075692109
	30	30	VDSEQPWKV	0.87/0.047847944

31	276	SAGVVANGV	0.88/0.033896012
32	62	KRFRTRKGK	0.01/0.87190794
33	266	GPRDL DHNF	0.85/0.018575467
34	85	TGPHKDAKF	0.95/-0.094427052
35	88	HKDAKFRER	0.66/0.17563285
36	269	DLDHNF GSA	0.95/-0.12192039
37	287	KGY PQFAEL	0.01/0.81610108
38	351	QPLL NPSAL	0.44/0.37436007
39	95	ERVEGVVWV	0.02/0.77795782
40	153	SRSQSR SRG	0.87/-0.072772925
41	160	RGESKSQSR	0.94/-0.14344557
42	258	TSNVTQCFG	0.95/-0.17865933
43	336	KFLEELNAF	0.17/0.58639788
44	8	DASEPQRGR	0.82/-0.065278576
45	147	APSRSQSRS	0.59/0.15949376
46	212	KPSRNQSPA	0.01/0.72358528
47	313	SGNTVVLTF	0.94/-0.21218693
48	312	ESGNTVVLT	0.68/- 0.00053717926
49	132	KLPNGVTVV	0.04/0.63929904
50	189	ALKSLGFDK	0.22/0.45660609
51	207	KTGTPKPSR	0.64/0.0092939722
52	94	RERVEGVVW	0.84/-0.19877636
53	156	QSR SRGESK	0.72/-0.094187755
54	81	YYLGTGPHK	0.13/0.49258046
55	92	KFRERVEGV	0.23/0.3870249
56	294	ELVPSTAAM	0.07/0.54195853
57	219	PASSQSAAK	1.00/-0.39375896
58	331	HPHLGKFLE	0.93/-0.32704756
59	166	QSRNPSSDR	0.82/-0.22002781
60	246	KPRWKRQPN	0.67/-0.073549592
61	204	KSAKTGTPK	0.85/-0.25520001
62	41	RNLVPINKK	0.11/0.47153736
63	174	RNHNSQDDI	0.71/-0.14818412
64	59	NVQKRFRTR	0.10/0.46031065
65	61	QKRFRTRKG	0.95/-0.41058221
66	366	TSPATVEPV	/0.53685795
67	253	PNDDVTSNV	0.01/0.52487438
68	52	NKLIGYWNV	0.04/0.4910114

69	317	VVLTFTTRV	0.00/0.52411543
70	303	LFDSHIVSK	0.91/-0.38718112
71	236	ETKEQKHEM	0.00/0.52163142
72	335	GKFLEELNA	0.88/-0.3764547
73	155	SQSRSRGES	0.98/-0.48969212
74	210	TPKPSRNQS	0.90/-0.41812113
75	118	RRKNSEPEI	0.06/0.41790866
76	370	TVEPVRDEV	/0.46443112
77	279	VVANGVKAK	0.02/0.44383583
78	305	DSHIVSKES	0.97/-0.51205745
79	309	VSKESGNTV	0.01/0.44037585
80	223	QSAAKILAR	0.00/0.43185349
81	128	HFNQKLPNG	0.60/-0.17257577
82	230	ARSQSSETK	0.02/0.40465747
83	176	HNSQDDIMK	0.57/-0.15094078
84	7	ADASEPQRG	0.92/-0.50517523
85	46	INKKDKNKL	0.01/0.39938551
86	151	SQSRSQSRS	0.65/-0.25747986
87	233	QSSETKEQK	0.79/-0.39868484
88	374	VRDEVSIET	/0.37206472
89	193	LGFDKPQEK	0.67/-0.29807002
90	76	PKLHFYYLG	0.97/-0.5998392
91	36	WKVIPRNLV	0.64/-0.271922
92	264	CFGPRDLDH	0.96/-0.59305311
93	222	SQSAAKILA	0.98/-0.61830664
94	169	NPSSDRNHN	0.62/-0.27202627
95	343	AFTREMQQQ	0.99/-0.64317551
96	310	SKESGNTVV	0.92/-0.58600038
97	183	MKAVAAALK	0.77/-0.43764113
98	162	ESKSQSRNP	0.71/-0.4052145
99	144	DSRAPSRSQ	0.97/-0.67182608
100	58	WNVQKRFRT	0.83/-0.53626171
101	120	KNSEPEIPH	0.75/-0.46313565
102	32	SEQPWKVIP	0.63/-0.37191504
103	77	KLHFYYLGT	0.51/-0.26014265
104	214	SRNQSPASS	0.61/-0.37235207
105	255	DDVTSNVTQ	0.94/-0.70386792
106	249	WKRQPNDDV	0.68/-0.53077535
107	80	FYYLGTGPH	0.54/-0.39331188

108	197	KPQEKDKKS	0.83/-0.71545221
109	299	TAAMLFDSDH	0.54/-0.42606284
110	126	IPHFNQKLP	0.78/-0.68352791
111	123	EPEIPHFNQ	0.75/-0.65410199
112	209	GTPKPSRNQ	0.90/-0.82823563
113	19	RIPYSLYSP	0.87/-0.81368293
114	195	FDKPQEKDK	0.58/-0.53498434
115	99	GVVWVAVDG	0.64/-0.60336552
116	320	TFTTRVTVP	0.68/-0.65033092
117	42	NLVPINKKD	0.86/-0.8308423
118	239	EQKHEMQKP	0.86/-0.84002514
119	328	PKDHPHLGK	0.97/-0.9627777
120	263	QCFGPRDL	0.93/-0.93474488
121	339	EELNAFTRE	0.56/-0.56800581
122	360	EFNPSQTSP	0.68/-0.69223836
123	307	HIVSKESGN	0.56/-0.58140183
124	281	ANGVKAKGY	0.52/-0.55856737
125	215	RNQSPASSQ	0.51/-0.5617737
126	304	FDSHIVSKE	0.75/-0.88250981
127	179	QDDIMKAVA	0.51/-0.65401886
128	2	ATVKWADAS	0.94/-1.1097636
129	184	KAVAAALKS	0.76/-0.95242894
130	136	GVTVVEEPD	0.55/-0.80294364
131	251	RQPNDDVTS	0.75/-1.0132369
132	332	PHLGKFL	0.56/-0.82481605
133	188	AALKSLGFD	0.93/-1.214132
134	363	PSQTSPATV	0.51/-0.8004713
135	148	PSRSQSR	0.97/-1.2882981
136	78	LHFYYLGTG	0.99/-1.3365084
137	213	PSRNQSPAS	0.99/-1.352974
138	5	KWADASEPQ	0.72/-1.1760276
139	190	LKSLGFDKP	0.93/-1.4108754
140	356	PSALEFNPS	0.62/-1.1466631
141	163	SKSQSRNPS	0.67/-1.2017927
142	237	TKEQKHEMQ	0.91/-1.5050854
143	354	LNPSALEFN	0.97/-1.6062851
144	334	LGKFLEELN	0.75/-1.5378383
145	4	VKWADASEP	0.60/-1.4799163
146	198	PQEKDKKSA	0.54/-1.7502832

	147	290	PQFAELVPS	0.56/-1.8487869
	148	27	PLLVDSEQP	0.77/-2.2873173
<b>HCoV-OC43</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	372	VLSENLNAY	0.98/1.1589641
	2	111	TADGNQRQL	0.86/1.0763098
	3	195	APNSRSTSR	0.99/0.72578835
	4	305	TSDPQFPIL	0.53/1.0784335
	5	355	GAIRFDSTL	1.00/0.59758371
	6	101	WYRHNRRSF	0.60/0.88504665
	7	320	AGAFFFGSR	0.98/0.49611093
	8	95	TEAKGYWYR	0.69/0.7854361
	9	348	VYELRYNGA	0.96/0.51262726
	10	214	RSRANSNGNR	0.91/0.5574033
	11	174	TRFPPGTVL	0.33/1.1317061
	12	70	TQFQKGKEF	0.95/0.49619917
	13	127	YLG TGPHAK	0.92/0.50505149
	14	410	SVAVPKSRV	0.82/0.55273831
	15	102	YRHNRRSFK	0.56/0.78670916
	<b>16</b>	<b>120</b>	<b>LPRWYFYYL</b>	<b>0.00/1.3445772</b>
	17	33	FRNVQTRGR	0.65/0.69112238
	18	140	TDIDGVYWV	0.60/0.66153271
	19	350	ELRYNGAIR	0.66/0.57692708
	20	389	MSPKPQRQR	0.82/0.40238414
	21	212	GSRSRANSNG	0.99/0.19801971
	22	180	TVLPQGYYI	0.11/1.071776
	23	292	NQNFGGGEM	0.86/0.24436914
	24	28	DQSDQFRNV	0.16/0.9288344
	25	362	TLSGFETIM	0.11/0.95739281
	26	248	ATKPQQVTK	0.02/1.0266931
	27	261	EVRQKILNK	0.85/0.18807761
	28	26	WADQSDQFR	0.94/0.067667846
	29	317	APTAGAFFF	0.00/0.99957346
	30	378	NAYQQQDGM	0.93/0.063430407
	31	38	TRGRRAPK	0.63/0.36004039
	32	235	QIASLV LAK	0.37/0.60747635
	33	104	HNRRSFKTA	0.58/0.38779782
	34	374	SEN LNAYQQ	0.54/0.415616

35	118	QLLPRWYFY	0.98/-0.029293394
36	35	NVQTRGRRA	0.95/-0.020171979
37	407	DNISVAVPK	0.99/-0.071861209
38	232	MADQIASLV	0.90/-0.0013435781
39	57	GNVVPYYSW	0.83/0.065642286
40	87	PIAPGVPAT	0.70/0.19052465
41	405	ENDNISVAV	0.95/-0.066271
42	236	IASLVLAKL	0.03/0.83832741
43	72	FQKGKEFEF	0.15/0.71231577
44	15	SGNRSGNGI	0.98/-0.12225342
45	98	KGYWYRHNH	0.37/0.46659975
46	281	TVQQCFGKR	0.76/0.075971959
47	52	QQPSGGNVV	0.43/0.40136015
48	117	RQLLPRWYF	0.12/0.69126931
49	119	LLPRWYFYY	0.66/0.14958306
50	144	GVYWVASNQ	0.74/0.064004786
51	61	PYYSWFSGI	0.33/0.42787822
52	203	RTSSRASSA	0.01/0.73618624
53	298	GEMLKLGTS	0.91/-0.16625057
54	112	ADGNQRQLL	0.65/0.092373213
55	358	RFDSTLSGF	0.64/0.10180045
56	170	EAIPTRFPP	0.96/-0.21946597
57	76	KEFEFAEGQ	0.85/-0.11156586
58	404	GENDNISVA	0.15/0.58622941
59	18	RSGNGILKW	0.36/0.37499581
60	200	STSRTSSRA	0.57/0.16279576
61	81	AEGQGVPIA	0.00/0.73152769
62	415	KSRVQQNKS	0.88/-0.16747608
63	414	PKSRVQQNK	0.88/-0.17052663
64	329	LELAKVQNL	0.15/0.54371414
65	110	KTADGNQRQ	0.88/-0.18966282
66	51	SQQPSGGNV	0.02/0.67014926
67	323	FFFGSRLEL	0.30/0.38427807
68	139	GTDIDGVYW	0.52/0.16174201
69	280	CTVQQCFGK	0.89/-0.21336165
70	258	TAKEVRQKI	0.03/0.64565508
71	351	LRYNGAIRF	0.01/0.66297897
72	204	TSSRASSAG	0.97/-0.29827064



73	306	SDPQFPILA	0.80/-0.12835707
74	326	GSRLELAKV	0.56/0.10387762
75	17	NRSGNGILK	0.00/0.66350694
76	154	DVNTPADIV	0.00/0.66288667
77	322	AFFFGSRLE	0.80/-0.14400224
78	308	PQFPILAEI	0.66/-0.015817526
79	99	GYWYRHNRR	0.16/0.47660875
80	418	VQQNKSIEL	0.19/0.43088901
81	369	IMKVLSENL	0.11/0.5046064
82	265	KILNKPRQK	0.67/-0.05586895
83	115	NQRQLLPRW	0.86/-0.24916488
84	65	WFSGITQFQ	0.98/-0.38070944
85	66	FSGITQFQK	0.75/-0.15600666
86	94	ATEAKGYWY	0.04/0.54584638
87	16	GNRSGNGIL	0.98/-0.40156265
88	365	GFETIMKVL	0.12/0.45554363
89	188	IEGSGRSAP	0.83/-0.26854815
90	349	YELRYNGAI	0.00/0.55373259
91	392	KPQRQRGHK	0.08/0.46484825
92	316	LAPTAGAFF	0.89/-0.3492627
93	371	KVLSENLNA	0.02/0.51461096
94	234	DQIASLVLA	0.87/-0.33606481
95	229	TPDMADQIA	0.93/-0.39632129
96	8	QSSSRASSG	0.68/-0.14745512
97	233	ADQIASLVL	0.85/-0.32030753
98	116	QRQLLPRWY	0.98/-0.45161339
99	321	GAFFFGSRL	0.00/0.52367268
100	31	DQFRNVQTR	0.01/0.47579119
101	147	WVASNQADV	0.00/0.47524332
102	156	NTPADIVDR	0.85/-0.37584138
103	381	QQQDGMMNM	0.00/0.47333015
104	210	SAGSRSRAN	0.92/-0.44776397
105	145	VYWVASNQA	0.91/-0.45971292
106	86	VPIAPGVPA	0.52/-0.081605456
107	208	ASSAGSRSR	0.00/0.42280948
108	344	PQKDVYELR	0.94/-0.51876437
109	266	ILNKPRQKR	0.01/0.40169297
110	48	TATSQQPSG	0.87/-0.46990769
111	88	IAPGVPATE	0.66/-0.266765

112	103	RHNRRSFKT	0.94/-0.55384533
113	202	SRTSSRASS	0.67/-0.28627804
114	263	RQKILNKPR	0.98/-0.59824813
115	409	ISVAVPKSR	0.67/-0.29111441
116	78	FEFAEGQGV	0.74/-0.41562362
117	227	GVTPDMADQ	0.99/-0.67107157
118	105	NRRSFKTAD	0.88/-0.56190498
119	39	RGRRAQPKQ	0.68/-0.37835187
120	217	ANSGNRTPT	0.99/-0.69366685
121	380	YQQQDGMMN	0.92/-0.64246371
122	260	KEVRQKILN	0.95/-0.68944282
123	198	SRSTSRTSS	1.00/-0.78806077
124	176	FPPGTVLPQ	0.92/-0.70942411
125	223	TPTSGVTPD	0.66/-0.47617744
126	417	RVQQNKSIE	0.92/-0.74387469
127	332	AKVQNLSGN	0.90/-0.74889762
128	272	QKRSPNKQC	0.84/-0.6929533
129	262	VRQKILNKP	0.85/-0.71258341
130	346	KDVYELRYN	0.84/-0.71158156
131	143	DGVYWVASN	0.96/-0.85322507
132	9	SSSRASSGN	0.83/-0.73755023
133	160	DIVDRDPSS	0.93/-0.84494347
134	269	KPRQKRSPN	0.77/-0.70191533
135	131	GPHAKDQYG	0.51/-0.4490066
136	382	QQDGMMNMS	0.61/-0.57713077
137	368	TIMKVLSEN	0.61/-0.5775436
138	275	SPNKQCTVQ	0.88/-0.86039888
139	186	YYIEGSGRS	0.89/-0.8913775
140	100	YWYRHNRRS	0.74/-0.80281384
141	218	NSGNRTPTS	0.73/-0.8219241
142	196	PNSRSTSRT	0.72/-0.82214697
143	113	DGNQRQLLP	0.77/-0.87846238
144	370	MKVLSENLN	0.99/-1.1110516
145	408	NISVAVPKS	0.60/-0.73476361
146	59	VVPYYSWFS	0.75/-0.88913655
147	393	PQRQRGHKN	0.62/-0.76562887
148	264	QKILNKPRQ	0.95/-1.1080764
149	20	GNGILKWAD	0.78/-0.94466833
150	395	RQRGHKNGQ	0.53/-0.72963819

	151	333	KVQNLSGNP	0.86/-1.0843094
	152	221	NRTPTSGVT	0.59/-0.81525772
	153	6	GKQSSSRAS	0.94/-1.1654096
	154	159	ADIVDRDPS	0.97/-1.2393525
	155	166	PSSDEAIPT	0.83/-1.1086528
	156	285	CFGKRGPNQ	0.67/-0.94935807
	157	135	KDQYGTDID	0.91/-1.1913038
	158	69	ITQFQKGKE	0.57/-0.87735713
	159	253	QVTKHTAKE	0.59/-0.90283739
	160	42	RAQPKQTAT	0.52/-0.83385458
	161	327	SRLELAKVQ	0.75/-1.0871909
	162	330	ELAKVQNLS	0.55/-0.91879418
	163	283	QQCFGKRG	0.52/-0.90087492
	164	162	VDRDPSSDE	0.96/-1.3437042
	165	375	ENLNAYQQQ	0.71/-1.0946865
	166	396	QRGHKNGQG	0.79/-1.1780586
	167	226	SGVTPDMAD	0.66/-1.1779906
	168	341	PDEPQKDVY	0.85/-1.4332236
	169	245	GKDATKPQQ	0.58/-1.1995382
	170	247	DATKPQQVT	0.51/-1.1432332
	171	290	GPNQNFEGG	0.56/-1.2629712
	172	331	LAKVQNLSG	0.75/-1.4736735
	173	385	GMMNMSPKP	0.85/-1.6132945
	174	178	PGTVLPQGY	0.65/-1.5461551
	175	383	QDGMMNMSP	0.61/-1.5386503
	176	337	LSGNPDEPQ	0.95/-1.9712696
	177	297	GGEMCLKGT	0.91/-1.9332697
	178	45	PKQTATSQQ	0.84/-2.0144397
<b>HCoV-HKU1</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Nucleocapsid</b>	1	314	APTPGAFFF	1.00/1.4961973
	2	355	STLPGFETI	1.00/0.97182876
	3	32	ERSHQTYNR	0.82/0.86769761
	4	100	WYKHNRSSF	0.70/0.97369498
	5	345	LRYSGSIRF	0.96/0.71316192
	6	194	ASNSRPGSR	0.88/0.72719808
	7	16	GNRSGILKK	0.90/0.66710451
	8	69	TQFQKGRDF	0.98/0.4921251

9	395	GVKQSPESF	0.90/0.55619454
10	51	STQPQGNTI	0.88/0.489655
<b>11</b>	<b>119</b>	<b>LPRWYFYYL</b>	<b>0.00/1.3445772</b>
12	166	TIQEAIPTTR	0.94/0.38718563
13	337	SPSKDTFEL	0.78/0.51268009
14	141	HEGIFWVAS	0.97/0.31072606
15	162	ARDPTIQEA	0.09/1.1728461
16	205	SRGPNNRSL	0.01/1.178805
17	13	SSSGNRSGI	0.71/0.47744339
18	275	NKFCNVQQC	0.88/0.29567564
19	366	VLKENLDAY	0.30/0.85327468
20	88	AYGIPPSEA	0.47/0.66066661
21	94	SEAKGYWYK	0.34/0.77827761
22	110	TADGQQKQL	0.88/0.23068034
23	116	KQLLPRWYF	0.38/0.72259239
24	356	TLPGFETIM	0.00/1.0953365
25	385	SLSPKPQRK	0.38/0.70228852
26	407	NLSADTQHI	0.86/0.21778056
27	246	DSKPQQVTK	0.16/0.8846158
28	125	YYLGTGPYA	0.44/0.60022141
29	264	ILMKPRQKR	0.96/0.059271792
30	276	KFCNVQQCF	0.70/0.30659755
31	269	RQKRTPNKF	0.41/0.59205225
32	235	IASLVLAKL	0.15/0.83832741
33	363	IMKVLKENL	0.16/0.76137225
34	139	ESHEGIFWV	0.08/0.83099216
35	230	DMADEIASL	0.87/0.03858306
36	231	MADEIASLV	0.64/0.26806972
37	138	GESHEGIFW	0.81/0.094269955
38	290	LQNFGNEML	0.81/0.091112397
39	349	GSIRFDSTL	0.02/0.87707935
40	272	RTPNKFCNV	0.04/0.85664392
41	173	TRFSPGTIL	0.00/0.89520939
42	163	RDPTIQEAI	0.98/-0.091534088
43	136	SYGESHEGI	0.98/-0.09514377
44	198	RPGSRSQSR	0.23/0.6542641
45	344	ELRYSGSIR	0.82/0.052614872
46	126	YLGTPGYAN	0.93/-0.062409977
47	179	TILPQGYV	0.01/0.85514416

48	170	AIPTRFSPG	0.87/-0.0074589008
49	39	NRGRKPQPK	0.79/0.059153307
50	15	SGNRSGILK	0.31/0.53667082
51	299	KLGTNDPQF	0.94/-0.093433378
52	374	YVNSNQNTV	0.03/0.80286352
53	373	AYVNSNQNT	0.72/0.10713683
54	292	NFGNEMCLK	0.82/0.0061638397
55	35	HQTYNRGRK	0.63/0.19607071
56	302	TNDPQFPIL	0.00/0.82315451
57	263	KILMKPRQK	0.68/0.112952
58	343	FELRYSGSI	0.36/0.40478423
59	42	RKPQPKFTV	0.56/0.1987804
60	289	PLQNFGNEM	0.97/-0.23728854
61	24	KTSWVDQSE	0.93/-0.21097773
62	285	GKRGPLQNF	0.54/0.17107091
63	2	SYTPGHHAG	0.28/0.41662649
64	48	FTVSTQPQG	0.91/-0.21785461
65	98	GYWYKHNRR	0.04/0.64327472
66	203	SQSRGPNNR	0.28/0.39298196
67	359	GFETIMKVL	0.21/0.45554363
68	227	VKPDMADEI	0.88/-0.240664
69	256	NAKEIRHKI	0.00/0.63518759
70	321	FFGSKLELF	0.70/-0.082833362
71	214	SRSNSNFRH	0.54/0.077132849
72	60	PHYSWFSGI	0.96/-0.36043775
73	252	VTKQNAKEI	0.53/0.067310078
74	306	QFPILAEAL	0.78/-0.1828691
75	164	DPTIQEAIP	0.98/-0.38787822
76	422	EDHSLLATL	/0.57708186
77	142	EGIFWVASH	0.83/-0.2631943
78	370	NLDAYVNSN	0.77/-0.20748117
79	174	RFSPGTILP	0.84/-0.28600685
80	124	FYYLGTGPY	0.68/-0.12632572
81	319	AFFFGSKLE	0.96/-0.40723022
82	117	QLLPRWYFY	0.57/-0.029293394
83	381	TVSGSLSPK	0.16/0.37354508
84	301	GTNDPQFPI	0.09/0.44294619
85	234	EIASLVLAK	0.14/0.37554337

86	57	NTIPHYSWF	0.00/0.48730721
87	192	RSASNSRPG	0.94/-0.45394261
88	190	SGRSASNSR	0.88/-0.39746696
89	391	QRKRGVKQS	0.72/-0.24787469
90	130	GPYANASYG	0.92/-0.45072802
91	259	EIRHKILMK	0.06/0.3990221
92	291	QNFGNEMLK	0.00/0.45211586
93	405	SLNLSADTQ	0.83/-0.38015142
94	122	WYFYYLGTG	1.00/-0.5715081
95	206	RGPNNRSLs	0.98/-0.56476422
96	212	SLSRNSNF	0.04/0.37027036
97	143	GIFWVASHQ	0.63/-0.23044717
98	103	HNRRSFKTA	0.00/0.38779782
99	204	QSRGPNNRS	0.91/-0.52645022
100	43	KPQPKFTVS	0.67/-0.31219191
101	62	YSWFSGITQ	0.77/-0.41601612
102	59	IPHYSWFSG	0.98/-0.63268884
103	376	NSNQNTVSG	0.86/-0.52848642
104	372	DAYVNSNQN	0.92/-0.60470553
105	128	GTGPYANAS	0.69/-0.38814964
106	410	ADTQHISND	0.96/-0.66010948
107	330	KRDSDADSP	0.87/-0.57057488
108	255	QNAKEIRHK	0.53/-0.25034625
109	278	CNVQQCFGK	0.56/-0.30105439
110	30	QSERSHQTY	0.53/-0.2890537
111	408	LSADTQHIS	0.56/-0.36127016
112	186	YVEGSGRSA	0.98/-0.79492794
113	347	YSGSIRFDS	0.98/-0.79775431
114	409	SADTQHISN	0.55/-0.37989788
115	351	IRFDSTLPG	0.93/-0.76138085
116	80	PDGQGVPIA	0.61/-0.45241142
117	176	SPGTILPQG	0.98/-0.82963646
118	87	IAYGIPPSE	0.80/-0.65853321
119	188	EGSGRSASN	0.80/-0.67107022
120	397	KQSPESFDS	0.96/-0.90646967
121	78	KFPDGQGVp	0.87/-0.87581571
122	350	SIRFDSTLP	0.95/-0.97054987
123	85	VPIAYGIPP	0.94/-0.97212355
124	96	AKGYWYKHN	0.65/-0.711482

	125	399	SPESFDSL N	0.56/-0.62619157
	126	325	KLELFKRDS	0.69/-0.78342076
	127	280	VQQCFGKRG	0.86/-0.96094544
	128	38	YNRGRKPQP	0.77/-0.9033711
	129	340	KDTFELRYS	0.57/-0.7253276
	130	131	PYANASYGE	0.99/-1.1726272
	131	293	FGNEM LKL G	0.63/-0.81381149
	132	6	GHHAGSRSS	0.81/-1.0943574
	133	283	CFGKRGPLQ	0.94/-1.2487202
	134	284	FGKRGPLQN	0.78/-1.0955521
	135	383	SGSLSPKPQ	0.91/-1.260197
	136	247	SKPQQVTKQ	0.83/-1.1942034
	137	329	FKRDS DADS	0.90/-1.2730956
	138	262	HKILMKPRQ	0.98/-1.3900266
	139	382	VSGSLSPKP	0.76/-1.3047531
	140	28	VDQSERSHQ	0.57/-1.1424233
	141	404	DSL NLSADT	0.69/-1.2804554
	142	123	YFY YLGTGP	0.90/-1.5129985
	143	225	SIVKPD MAD	0.53/-1.1861916
	144	5	PGHHAGSRS	0.73/-1.389111
	145	308	PILAE LAPT	0.78/-1.5495611
	146	375	VNSNQNTVS	0.65/-1.4984255
	147	245	KDSKPQQVT	0.52/-1.3880026
	148	22	LKKT SWVDQ	0.57/-1.4740862
	149	72	QKGRDFKFP	0.66/-1.651665
<b>SARS-COV-2</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	<b>1</b>	<b>1192</b>	<b>NL NESLIDL</b>	<b>0.92/1.5879094</b>
	2	236	TRFQTLLAL	0.99/1.4877641
	3	983	RLDKVEAEV	0.71/1.6892505
	4	679	NSPRRARSV	0.87/1.0916413
	5	292	ALDPLSETK	0.92/1.0117189
	6	204	YSKHTPINL	0.47/1.4169218
	<b>7</b>	<b>820</b>	<b>DLLFNKVTL</b>	<b>0.73/1.1189541</b>
	8	972	AISSVLNDI	0.99/0.84037452
	9	1125	NCDVVIGIV	0.93/0.83215588
	10	97	KSNIIRGWI	0.89/0.86720593
	<b>11</b>	<b>846</b>	<b>ARDLICAQK</b>	<b>0.82/0.93408287</b>

12	618	TEVPVAIHA	0.98/0.76924266
13	852	AQKFNGLTV	0.98/0.73319244
14	115	QSLIVNNA	0.95/0.76004022
15	123	ATNVVIKVC	0.84/0.8570027
16	612	YQGVNCTEV	0.69/0.96038564
17	535	KNKCVNFNF	0.97/0.64771057
18	54	LFLPFFSNV	0.83/0.76135348
19	464	FERDISTEI	0.67/0.92124621
20	735	SVDCTMYIC	0.99/0.56080212
21	706	AYSNNSIAI	0.71/0.83826283
22	814	KRSFIEDLL	0.84/0.70530797
23	958	ALNTLVKQL	0.14/1.3939207
24	1185	RLNEVAKNL	0.23/1.2586948
25	345	TRFASVYAW	0.37/1.1119585
26	28	YTNSFTRGV	0.99/0.4496507
27	350	VYAWNRKRI	0.93/0.49736819
28	171	VSQPFLMDL	0.64/0.78489273
29	233	INITRFQTL	0.88/0.51843698
30	119	IVNNATNVV	0.62/0.75028555
<b>31</b>	<b>1060</b>	<b>VVFLHVTYV</b>	<b>0.59/0.77948535</b>
32	333	TNLCPFGEV	0.87/0.49897438
33	163	ANNCTFEYV	0.67/0.68976923
34	336	CPFGEVFNA	0.86/0.49893355
35	976	VLNDILSRL	0.19/1.1679864
36	634	RVYSTGSNV	0.61/0.74770151
37	725	EILPVSMTK	0.85/0.5027134
38	810	SKPSKRSFI	0.98/0.37244882
39	448	NYNYLYRLF	1.00/0.34772566
40	502	GVGYQPYRV	0.86/0.48701691
41	833	FIKQYGDCL	0.76/0.57526737
42	880	GTITSGWTF	0.75/0.57483752
43	319	RVQPTESIV	0.38/0.94410253
44	433	VIAWNSNNL	0.82/0.4956948
45	456	FRKSNLKPF	0.96/0.34990342
46	836	QYGDCLGDI	0.68/0.61787268
47	1006	TYVTQQLIR	0.99/0.29995915
48	797	FGGFNFSQI	0.45/0.83156397
49	280	NENGTITDA	0.99/0.2889406
50	59	FSNVTWFHA	0.53/0.74877447



51	399	SFVIRGDEV	0.97/0.30742774
52	395	VYADSFVIR	0.01/1.2592861
53	780	EVFAQVKQI	0.79/0.4771388
54	416	GKIADYNYK	0.99/0.27547906
55	62	VTWFHAIHV	0.41/0.85297789
56	827	TLADAGFIK	0.26/0.99225063
57	298	ETKCTLKSF	0.90/0.34635795
58	318	FRVQPTESI	1.00/0.24226172
59	925	NQFNSAIGK	0.76/0.48116482
60	609	AVLYQGVNC	0.96/0.27961479
61	892	AALQIPFAM	0.30/0.936978
62	1000	RLQSLQTYV	0.12/1.1162891
63	516	ELLHAPATV	0.39/0.84172953
64	1010	QQLIRAAEI	0.91/0.3197616
<b>65</b>	<b>903</b>	<b>AYRFNGIGV</b>	<b>0.19/1.0390298</b>
66	1171	GINASVVNI	0.00/1.2240296
67	1127	DVVIGIVNN	0.95/0.2685717
68	677	QTNSPRRAR	0.66/0.55678436
69	764	NRALTGIAV	0.34/0.86148546
70	1026	ATKMSECVL	0.97/0.2116657
<b>71</b>	<b>1054</b>	<b>QSAPHGVVF</b>	<b>0.44/0.73864024</b>
72	460	NLKPFERDI	0.41/0.75730112
73	973	ISSVLNDIL	0.95/0.20493016
74	1124	GNCDVVIGI	0.04/1.1035891
75	1055	SAPHGVVFL	0.35/0.78352262
76	504	GYQPYRVVV	0.23/0.90122156
77	1099	GTHWFVTQR	0.27/0.85459551
78	382	VSPTKLNDL	0.82/0.30409279
79	121	NNATNVVIK	0.71/0.41011789
80	759	FCTQLNRAL	0.85/0.26316347
81	122	NATNVVIKV	0.69/0.41830926
82	98	SNIIRGWIF	0.09/1.0182657
83	954	QNAQALNTL	0.70/0.40665469
84	926	QFNSAIGKI	0.58/0.5145498
85	386	KLNDLCFTN	0.42/0.67438404
86	327	VRFPNITNL	0.04/1.0522648
87	444	KVGGNYNYL	0.01/1.0792264
88	458	KSNLKPFER	0.01/1.0757571
89	258	WTAGAAAYY	0.54/0.5440791

90	1164	VDLGDISGI	0.95/0.12980383
91	757	GSFCTQLNR	0.83/0.23591759
92	1095	FVSNGTHWF	0.61/0.45070133
93	396	YADSFVIRG	0.00/1.0606061
94	478	TPCNGVEGF	0.84/0.22053613
95	392	FTNVYADSF	0.86/0.18825795
96	1062	FLHVTYVPA	0.95/0.094842726
97	781	VFAQVKQIY	0.71/0.33134329
98	361	CVADYSVLY	0.83/0.20890758
99	798	GGFNFSQIL	0.82/0.21884931
100	1144	ELDSFKEEL	0.27/0.76364207
101	529	KSTNLVKNK	0.99/0.032044011
102	1163	DVDLGDISG	0.89/0.13059576
103	324	ESIVRFPNI	0.18/0.84007409
104	1086	KAHFPREGV	0.94/0.076210395
105	923	IANQFNSAI	0.93/0.085106375
106	989	AEVQIDRLI	0.86/0.1550506
107	102	RGWIFGTTL	0.80/0.21354425
108	685	RSVASQSII	0.57/0.43730301
109	449	YNYLYRLFR	0.84/0.16648622
110	1190	AKNLNESLI	0.04/0.96099358
111	584	ILDITPCSF	0.69/0.31041717
112	15	CVNLTTTRTQ	0.98/0.019925455
113	285	ITDAVDCAL	0.75/0.23848008
114	89	GVYFASTEK	0.00/0.98684427
115	55	FLPFFSNVT	0.72/0.26607576
116	538	CVNFNFNGL	0.00/0.9787292
117	93	ASTEKSNII	0.49/0.48581774
118	487	NCYFPLQSY	0.74/0.22240704
119	494	SYGFQPTNG	0.98/-0.022026824
120	138	DPFLGVYYH	0.41/0.54564184
121	1181	KEIDRLNEV	0.00/0.95542235
122	208	TPINLVRDL	0.01/0.93604158
<b>123</b>	<b>951</b>	<b>VVNQNAQAL</b>	<b>0.16/0.78381232</b>
124	135	FCNDPFLGV	0.00/0.94369976
125	252	GDSSSGWTA	0.79/0.15356838
126	153	MESEFRVYS	0.90/0.042492164
127	387	LNDLCFTNV	0.31/0.63039809
128	268	GYLQPRTFL	0.25/0.68904137

129	13	SQCVNLTTR	0.71/0.22569446
130	255	SSGWTAGAA	0.82/0.10844838
131	198	DGYFKIYSK	0.75/0.17701473
132	847	RDLICAQKF	0.93/-0.0040749685
133	342	FNATRFASV	0.18/0.74271004
134	1067	YVPAQEKNF	0.94/-0.021668534
135	509	RVVVLSEFEL	0.54/0.37790287
136	869	MIAQYTSAL	0.41/0.50570354
137	413	GQTGKIADY	0.95/-0.036742046
138	474	QAGSTPCNG	0.73/0.17553918
139	530	STNLVKNKC	0.77/0.13357426
140	295	PLSETKCTL	0.88/0.023422332
141	139	PFLGVYYHK	0.56/0.3361685
142	155	SEFRVYSSA	0.44/0.44339405
143	987	VEAEVQIDR	0.34/0.53761645
144	1076	TTAPAICHD	0.46/0.41720267
145	675	QTQTNSPRR	0.90/-0.025268287
146	554	ESNKKFLPF	0.99/-0.11633723
147	787	QIYKTPPIK	0.71/0.16240686
148	525	CGPKKSTNL	0.81/0.06208729
149	536	NKCVNFNFN	0.99/-0.11902214
150	576	VRDPQTLEI	0.06/0.81001705
151	940	STASALGKL	0.98/-0.11279199
152	559	FLPFQQFGR	0.80/0.066311857
153	400	FVIRGDEV	0.89/-0.02750081
154	238	FQTLLALHR	0.85/0.0028903032
155	30	NSFTRGVYY	0.02/0.83054514
156	1007	YVTQQLIRA	0.67/0.18003803
157	533	LVKNKCVNF	0.00/0.85001899
158	1118	DNTFVSGNC	0.95/-0.1022437
159	409	QIAPGQTGK	0.00/0.83303707
<b>160</b>	<b>901</b>	<b>QMAYRFNGI</b>	<b>0.13/0.69832921</b>
<b>161</b>	<b>1208</b>	<b>QYIKWPWYI</b>	<b>/0.82786365</b>
162	313	YQTSNFRVQ	0.99/-0.16260307
163	1123	SGNCDVVIG	0.87/-0.044511101
164	912	TQNVLYENQ	0.95/-0.12643186
165	1115	ITTDNTFVS	0.79/0.031855701
166	378	KCYGVSPK	0.99/-0.17129001

167	532	NLVKNKCVN	0.85/-0.03143559
168	189	LREFVFKNI	0.13/0.68287982
169	348	ASVYAWN RK	0.41/0.4028606
170	568	DIADTTDAV	0.01/0.79165536
171	1016	AEIRASANL	0.78/0.017047345
172	379	CYGVSP TKL	0.00/0.79227611
173	270	LQPRTFLLK	0.00/0.78938338
174	777	NTQEVFAQV	0.10/0.68271776
175	77	KRFDNPVLP	0.72/0.060755904
176	965	QLSSNFGAI	0.56/0.21791063
177	84	LPFNDGVYF	0.80/-0.022575978
178	353	WNRKRISNC	0.62/0.14927305
179	360	NCVADYSVL	0.94/-0.17191863
180	1096	VSNGTHW FV	0.32/0.44642728
181	687	VASQSI IAY	0.52/0.24420376
182	883	TSGWTFGAG	0.67/0.087730892
183	975	SVLNDILSR	0.90/-0.14367289
184	755	QYGSFCTQL	0.02/0.73586609
<b>185</b>	<b>1041</b>	<b>DFCGKGYHL</b>	<b>0.02/0.73481455</b>
186	1121	FVSGNCDVV	0.18/0.57456887
187	1051	SFPQSAPHG	0.80/-0.045461664
188	206	KHTPINLVR	0.94/-0.18910122
189	676	TQTNSPRRA	0.90/-0.15761293
190	417	KIADYNYKL	0.00/0.74005063
191	697	MSLGAENSV	0.91/-0.17159345
192	447	GNYNLYRL	0.01/0.72750718
193	782	FAQVKQIYK	0.22/0.51645801
194	23	QLPPAYTNS	0.91/-0.1743011
195	999	GRLQSLQTY	0.56/0.17547758
196	1137	VYDPLQPEL	0.14/0.59484561
197	1226	AIVMVTIML	/0.73431223
198	169	EYVSQPFLM	0.00/0.7324723
199	264	AYYVGYLQP	0.95/-0.21896143
200	630	TPTWRVYST	0.64/0.090783366
201	614	GVNCTEVPV	0.00/0.72983584
202	227	VDLPIGINI	0.13/0.59776359
203	1150	EELDKYFKN	0.91/-0.18583258
<b>204</b>	<b>1056</b>	<b>APHGVVFLH</b>	<b>0.14/0.57504297</b>
205	592	FGGVSVITP	0.97/-0.25743026

206	894	LQIPFAMQM	0.74/-0.037819834
207	234	NITRFQTLL	0.57/0.13034199
208	664	IPIGAGICA	0.78/-0.081495116
209	321	QPTESIVRF	0.14/0.55765263
210	253	DSSSGWTAG	0.72/-0.023444227
211	27	AYTNSFTRG	0.56/0.13432448
212	550	GVLTESNKK	0.67/0.022718272
213	193	VFKNIDGYF	0.84/-0.14767335
214	660	YECDIPIGA	0.07/0.62181879
215	288	AVDCALDPL	0.53/0.15420622
<b>216</b>	<b>1209</b>	<b>YIKWPWYIW</b>	<b>/0.68245283</b>
217	1204	GKYEQYIKW	0.01/0.6658816
218	1085	GKAHFPREG	0.94/-0.27137105
219	734	TSVDCTMYI	0.07/0.59295884
220	312	IYQTSNFRV	0.27/0.39102148
221	46	SVLHSTQDL	0.59/0.070628293
222	389	DLCFTNVYA	0.06/0.59541601
223	501	NGVGYPYR	0.63/0.025251303
<b>224</b>	<b>919</b>	<b>NQKLIANQF</b>	<b>0.00/0.65422365</b>
225	269	YLQPRTFLL	0.01/0.64342282
226	354	NRKRISNCV	0.01/0.63952346
227	627	DQLTPTWRV	0.00/0.6487372
228	446	GGNYNYLYR	0.00/0.64830171
229	394	NVYADSFVI	0.76/-0.11532989
230	32	FTRGVYYPD	0.91/-0.26674456
231	370	NSASFSTFK	0.01/0.63262149
232	832	GFIKQYGDC	0.85/-0.20761876
233	1196	SLIDLQELG	0.75/-0.11355354
234	1210	IKWPWYIWL	/0.63616187
235	907	NGIGVTQNV	0.03/0.60456213
236	956	AQALNTLVK	0.00/0.62945518
237	712	IAIPTNFTI	0.01/0.6169805
238	450	NYLYRLFRK	0.00/0.62171566
239	1262	EPVLKGVKL	/0.61158557
240	1165	DLGDISGIN	0.84/-0.23244334
241	505	YQPYRVVVL	0.01/0.5965429
242	688	ASQSIIAYT	0.99/-0.3882286
243	1183	IDRLNEVAK	0.01/0.59035876
244	1122	VSGNCDVVI	0.68/-0.07996517

245	149	NKSWMESEF	0.77/-0.1756724
246	626	ADQLTPTWR	0.93/-0.33882418
247	1194	NESLIDLQE	0.77/-0.17943301
248	334	NLCPFGEVF	0.00/0.58858915
249	95	TEKSNIIRG	0.07/0.51324547
250	86	FNDGVYFAS	0.84/-0.2570983
251	136	CNDPFLGVY	0.00/0.58175107
252	221	SALEPLVDL	0.10/0.47975488
253	686	SVASQSIIA	0.12/0.45958209
254	997	ITGRLQSLQ	0.82/-0.24138811
255	38	YPDKVFRSS	0.90/-0.32638397
256	1169	ISGINASVV	0.00/0.56782952
257	186	FKNLREFVF	0.19/0.37421466
258	1044	GKGYHLMSF	0.80/-0.23905486
259	402	IRGDEVROI	0.02/0.53867845
260	410	IAPGQTGKI	0.12/0.43746731
261	3	VFLVLLPLV	0.95/-0.39272787
262	993	IDRLITGRL	0.96/-0.40288235
263	477	STPCNGVEG	0.68/-0.12441887
264	1221	IAGLIAIVM	/0.5530776
265	83	VLPFNDGVY	0.90/-0.34931234
266	1133	VNNTVYDPL	0.91/-0.36386937
267	1175	SVVNIQKEI	0.92/-0.37518574
268	1195	ESLIDLQEL	0.04/0.50136368
269	112	SKTQSLIV	0.65/-0.11336405
270	1173	NASVVNIQK	0.04/0.48923419
271	42	VFRSSVLHS	0.78/-0.25143064
272	60	SNVTWFHAI	0.00/0.52175752
273	1028	KMSECVLGQ	0.86/-0.33830367
274	691	SIIAYTMSL	0.08/0.4387551
275	125	NVVIKVCEF	0.08/0.43765697
276	916	LYENQKLIA	0.98/-0.46793709
277	250	TPGDSSSGW	0.72/-0.20818618
278	187	KNLREFVFK	0.03/0.48025447
279	551	VLTESNKKF	0.91/-0.40127646
280	754	LQYGSFCTQ	0.73/-0.22195096
281	708	SNNSIAIPT	0.97/-0.46238529
282	50	STQDLFLPF	0.96/-0.45381589
283	563	QQFGRDIAD	0.97/-0.46484265

284	75	GTKRFDNPV	0.02/0.48077201
285	813	SKRSFIEDL	0.55/-0.054529485
286	644	QTRAGCLIG	0.94/-0.44643291
287	748	ECSNLLLQY	0.86/-0.36907603
288	701	AENSVAYSN	0.06/0.42092325
289	710	NSIAIPTNF	0.00/0.48057058
290	215	DLPQGFSAL	0.00/0.47967982
291	195	KNIDGYFKI	0.04/0.43907249
292	465	ERDISTEY	0.04/0.43514758
293	1224	LIAIVMTI	/0.47352128
294	1081	ICHDGKAHF	0.00/0.47336462
295	678	TNSPRRARS	0.00/0.46942667
296	962	LVKQLSSNF	0.62/-0.15274303
297	986	KVEAEVQID	0.99/-0.52449453
298	85	PFNDGVYFA	0.02/0.44528067
299	1106	QRNFYEPQI	0.00/0.46521579
300	1129	VIGIVNNTV	0.00/0.45624515
301	655	HVNNSYECD	0.98/-0.52703499
302	898	FAMQMAYRF	0.59/-0.13988547
303	783	AQVKQIYKT	1.00/-0.55131466
304	1214	WYIWLGFIA	/0.44831422
305	645	TRAGCLIGA	0.01/0.4360915
306	88	DGVYFASTE	0.88/-0.43436613
307	1102	WFVTQRNFY	0.01/0.43437486
308	403	RGDEVQRQA	0.01/0.41897524
309	63	TWFHAIHVS	0.93/-0.51246264
310	1078	APAICHDGK	0.65/-0.23312101
311	239	QTLLALHRS	0.89/-0.47398884
312	182	KQGNFKNLR	0.03/0.38099958
313	2	FVFLVLLPL	0.00/0.40759495
314	704	SVAYSNNNSI	0.03/0.37331941
315	937	SLSTASAL	0.00/0.40309069
316	260	AGAAAYYVG	0.91/-0.51126797
317	499	PTNGVGYQP	0.91/-0.51404835
<b>318</b>	<b>1219</b>	<b>GFIAGLIAI</b>	<b>/0.38830748</b>
319	235	ITRFQTLLA	0.00/0.38611895
320	1264	VLKGVKLHY	/0.38501663
321	1119	NTFVSGNCD	0.97/-0.59392325
322	596	SVITPGTNT	0.58/-0.22309952

323	1167	GDISGINAS	0.90/-0.55720895
324	854	KFNGLTVLP	0.53/-0.18989855
325	776	KNTQEVFAQ	0.92/-0.58719275
326	917	YENQKLIAN	0.54/-0.20739899
327	263	AAYYVGYLQ	0.90/-0.5712202
328	214	RDLPQGFSA	0.84/-0.5154096
329	1203	LGKYEQYIK	0.86/-0.5409742
330	473	YQAGSTPCN	0.95/-0.63128422
331	913	QNVLYENQK	0.52/-0.20458317
332	423	YKLPDDFTG	0.81/-0.49564959
333	950	DVVNQNAQA	0.68/-0.36599599
334	484	EGFNCYFPL	0.68/-0.36825199
335	768	TGIAVEQDK	0.55/-0.24021556
336	896	IPFAMQMAY	0.97/-0.66172733
337	352	AWNRRKRISN	0.63/-0.32440239
338	226	LVDLPIGIN	0.70/-0.39579694
339	517	LLHAPATVC	0.85/-0.54749701
340	1161	SPDVDLGDI	0.59/-0.2922528
341	81	NPVLPFNDG	0.63/-0.348098
342	904	YRFNGIGVT	0.85/-0.56977742
343	763	LNRALTGIA	0.95/-0.67111733
344	190	REFVFKNID	0.72/-0.44130943
345	695	YTMSLGAEN	0.98/-0.70186504
346	70	VSGTNGTKR	0.63/-0.35311067
347	693	IAYTMSLGA	0.82/-0.54699421
348	161	SSANNCTFE	0.63/-0.36027531
349	1075	FTTAPAICH	0.92/-0.65180223
350	796	DFGGFNFSQ	0.68/-0.41568299
351	340	EVFNATRFA	0.75/-0.49056096
352	909	IGVTQNVLY	0.93/-0.67251748
353	146	HKNNKSWME	0.98/-0.72398028
354	673	SYQTQTNSP	0.99/-0.7400553
355	337	PFGEVFNAT	0.63/-0.38873589
356	835	KQYGDCLGD	0.98/-0.7445586
357	571	DTTDAVRDP	0.68/-0.44612401
358	66	HAIHVSGTN	0.93/-0.70085024
359	362	VADYSVLYN	0.67/-0.44691471
360	212	LVRDLPQGF	0.90/-0.68007362
361	1023	NLAATKMSE	0.53/-0.31242396



362	598	ITPGTNTSN	0.93/-0.73041048
363	586	DITPCSFGG	0.97/-0.77631283
364	126	VVIKVCEFQ	0.61/-0.42185546
365	930	AIGKIQDSL	0.58/-0.3943784
366	430	TGCVIAWNS	0.76/-0.5860783
367	1029	MSECVLGQS	0.93/-0.75851111
368	581	TLEILDITP	0.63/-0.46402363
369	597	VITPGTNTS	0.75/-0.5957304
370	113	KTQSL LIVN	0.53/-0.38088275
371	955	NAQALNTLV	0.73/-0.58185384
372	1097	SNGTHW FVT	0.80/-0.65699727
373	893	ALQIPFAMQ	0.65/-0.5129913
374	64	WFHAIHVSG	0.57/-0.43580992
375	1017	EIRASANLA	0.83/-0.69786445
376	1112	PQIITTDNT	0.93/-0.79950416
377	110	LDSKTQ SLL	0.77/-0.64316734
378	247	SYLTPGDSS	0.95/-0.82460435
379	611	LYQGVNCTE	0.98/-0.86124867
380	48	LHSTQDLFL	0.80/-0.70004985
381	1117	TDNTFVSGN	0.55/-0.45005215
382	713	AIPTNFTIS	0.81/-0.71497488
383	885	GWTFGAGAA	0.58/-0.48541337
384	621	PVAIHADQL	0.68/-0.59137254
385	1187	NEVAKNLNE	0.62/-0.53369454
386	426	PDDFTGCVI	0.65/-0.5747398
387	291	CALDPLSET	0.56/-0.48548595
388	741	YICGDSTEC	0.59/-0.51750479
389	231	IGINTRFQ	0.87/-0.80177053
390	726	ILPVSMTKT	0.63/-0.5635257
391	633	WRVYSTGSN	0.80/-0.74120623
392	276	LLKYNENGT	0.98/-0.9406483
393	657	NNSYECDIP	0.99/-0.962859
394	74	NGTKRFDNP	0.59/-0.56674644
395	5	LVLLPLVSS	0.97/-0.947971
396	1158	NHTSPDV DL	0.86/-0.84283319
397	225	PLVDLP IGI	0.54/-0.52759395
398	573	TDAVRDPQT	0.70/-0.70467958
399	1025	AATKMSECV	0.70/-0.70789169
400	566	GRDIADTTD	0.68/-0.70104485

401	1159	HTSPDVDLG	0.57/-0.59425387
402	580	QTLEILDIT	0.99/-1.0148329
403	730	SMTKTSVDC	0.93/-0.96031133
404	547	TGTGVLTES	0.63/-0.66250487
405	245	HRSYLTPGD	0.94/-0.97707267
406	391	CFTNVYADS	0.61/-0.64858188
407	911	VTQNVLYEN	0.89/-0.93287129
408	863	PLLTDEMIA	0.88/-0.93527199
409	522	ATVCGPKKS	0.93/-0.99606588
410	498	QPTNGVGyQ	0.60/-0.66984026
411	6	VLLPLVSSQ	0.69/-0.76803716
412	750	SNLLLQYGS	0.51/-0.61301734
413	756	YGSFCTQLN	0.87/-0.97427523
414	665	PIGAGICAS	0.85/-0.95602426
415	49	HSTQDLFLP	0.91/-1.0195034
416	646	RAGCLIGAE	0.68/-0.80187557
417	770	IAVEQDKNT	0.94/-1.0740015
418	981	LSRLDKVEA	0.60/-0.74646517
419	132	EFQFCNDPF	0.51/-0.67539709
420	80	DNPVLPFND	0.77/-0.93836403
421	539	VNFNFNGLT	0.85/-1.0241041
422	52	QDLFLPFFS	0.69/-0.88010274
423	4	FLVLLPLVS	0.69/-0.88344636
424	19	TTRTQLPPA	0.56/-0.75895321
425	463	PFERDISTE	0.85/-1.0494737
426	471	EIYQAGSTP	0.56/-0.80393159
427	578	DPQTLEILD	0.77/-1.0158216
428	938	LSSTASALG	0.83/-1.0760172
429	440	NLDSKVGGN	0.96/-1.2071085
430	789	YKTPPIKDF	0.51/-0.76410238
431	1047	YHLMSFPQS	0.97/-1.2266547
432	124	TNVVIKVCE	0.89/-1.1756177
433	873	YTSALLAGT	0.79/-1.0772384
434	545	GLTGTGVLt	0.68/-0.97105971
435	952	VNQNAQALN	0.87/-1.1959752
436	108	TTLDSKTQS	0.55/-0.8761585
437	492	LQSYGFQPT	0.95/-1.2840322
438	7	LLPLVSSQC	0.91/-1.2472776
439	224	EPLVDLPIG	0.88/-1.2198089

	440	375	STFKCYGVS	0.87/-1.2291059
	441	438	SNNLDSKVG	0.74/-1.1128457
	442	858	LTVLPPLLT	0.61/-0.98417306
	443	173	QPFLMDLEG	0.65/-1.0346008
	444	931	IGKIQDSLS	0.54/-0.93081472
	445	544	NGLTGTGVL	0.80/-1.2226609
	446	414	QTGKIADYN	0.68/-1.1328443
	447	1148	FKEELDKYF	0.55/-1.0106407
	448	385	TKLNDLCFT	0.74/-1.2190256
	449	541	FNFNGLTGT	0.59/-1.0862641
	450	1069	PAQEKNFTT	0.57/-1.0798732
	451	1186	LNEVAKNLN	0.99/-1.5031759
	452	932	GKIQDSLSS	0.91/-1.4344021
	453	282	NGTITDAVD	0.81/-1.3404228
	454	792	PPIKDFGGF	0.64/-1.1739676
	455	632	TWRVYSTGS	0.97/-1.5043721
	456	650	LIGAEHVNN	0.63/-1.1938114
	457	216	LPQGFSALE	0.58/-1.1812157
	458	244	LHRSYLTPG	0.68/-1.3189648
	459	25	PPAYTNSFT	0.60/-1.269398
	460	309	EKGIYQTSN	0.66/-1.3328727
	461	963	VKQLSSNFG	0.76/-1.4346284
	462	1138	YDPLQPELD	0.96/-1.7567044
	463	727	LPVSMTKTS	0.94/-1.8342409
	464	1162	PDVDLGDIS	0.75/-1.685451
	465	1145	LDSFKEELD	0.75/-1.6857213
	466	1142	QPELDSFKE	0.82/-1.796165
	467	491	PLQSYGFQP	0.55/-1.5478997
	468	802	FSQILPDPS	0.60/-1.6421824
	469	945	LGKLQDVVN	0.60/-1.7300742
	470	613	QGVNCTEVP	0.59/-1.7221469
<b>SARS-CoV</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	1	1153	GINASVVNI	0.97/1.2240296
	2	982	RLQSLQTYV	0.84/1.1162891
	3	188	KNKDGFLYV	0.74/1.2048106
	4	447	KLRPFERDI	0.87/1.0022281
	5	162	EYISDAFSL	0.79/1.0588161

6	1071	FPREGVFVF	0.83/1.013369
7	250	AAYFVGYLK	0.99/0.80755975
<b>8</b>	<b>1042</b>	<b>VVFLHVTYV</b>	<b>0.99/0.77948535</b>
9	382	VYADSFVVK	0.76/0.94660302
10	965	RLDKVEAEV	0.00/1.6892505
<b>11</b>	<b>883</b>	<b>QMAYRFNGI</b>	<b>0.98/0.69832921</b>
<b>12</b>	<b>1023</b>	<b>DFCGKGYHL</b>	<b>0.94/0.73481455</b>
13	834	AQKFNGLTV	0.93/0.73319244
<b>14</b>	<b>1174</b>	<b>NLNESLIDL</b>	<b>0.06/1.5879094</b>
15	403	GVIADYNYK	0.90/0.72832925
16	1084	WFITQRNFF	0.89/0.73497574
17	366	CYGVSATKL	0.92/0.68821391
18	1186	GKYEQYIKW	0.94/0.6658816
19	439	KYRYLRHGK	0.98/0.60308771
20	958	VLNDILSRL	0.41/1.1679864
21	818	QYGECLGDI	0.96/0.54398657
22	502	ELLNAPATV	0.77/0.7335387
23	809	TLADAGFMK	0.80/0.7032719
24	115	IIINNSTNV	0.95/0.54319162
25	1063	ICHEGKAYF	0.98/0.49852803
26	311	GDVVRFPNI	0.87/0.60715172
27	314	VRFPNITNL	0.42/1.0522648
28	320	TNLCPFGEV	0.97/0.49897438
29	156	AFNCTFEYI	0.55/0.88025204
30	356	YNSTFFSTF	0.86/0.56550332
31	175	KSGNFKHLR	0.80/0.62538011
32	796	KRSFIEDLL	0.72/0.70530797
33	415	DFMGCVLAW	0.99/0.43251469
34	940	ALNTLVKQL	0.02/1.3939207
35	717	SVDCNMYIC	0.58/0.83015798
36	726	GDSTECANL	0.97/0.43466714
37	236	TAFSPAQDI	0.98/0.40751572
38	562	VRDPKTSEI	0.26/1.1009364
39	119	NSTNVVIRA	0.85/0.50140518
40	954	AISSVLNDI	0.50/0.84037452
41	1088	QRNFFSPQI	0.41/0.92732642
42	1054	ERNFTTAPA	0.98/0.34762297
43	390	KGDDVRQIA	0.90/0.42722844
44	171	DVSEKSGNF	0.99/0.33539521

45	755	EQDRNTREV	0.69/0.63416129
46	404	VIADYNYKL	1.00/0.31666138
47	55	TQDLFLPFY	0.84/0.43933781
48	1167	RLNEVAKNL	0.02/1.2586948
49	267	DENGTITDA	0.88/0.38987142
50	1070	YFPREGVHV	0.98/0.28571274
51	201	QPIDVVRDL	0.09/1.1565693
52	604	TDVSTAIHA	0.84/0.40572419
53	435	NYNYKYRYL	0.03/1.2104636
54	321	NLCPFGEVF	0.65/0.58858915
55	440	YRYLRHGKL	0.95/0.28440489
56	282	PLAELKCSV	0.99/0.23875545
57	919	SLTTTSTAL	0.71/0.50881171
58	108	NNKSQSVII	0.94/0.27236091
59	947	QLSSNFGAI	0.99/0.21791063
60	874	AALQIPFAM	0.25/0.936978
61	369	VSATKLNDL	0.79/0.39639835
62	223	PLGINITNF	0.98/0.20545682
63	120	STNVVIRAC	0.38/0.80215202
64	374	LNDLCFSNV	0.45/0.73166267
65	671	QKSIVAYTM	0.90/0.27893316
66	418	GCVLAWNTR	0.97/0.2030597
67	616	TPAWRIYST	0.74/0.42093465
68	590	ASSEVAVLY	0.34/0.81690548
69	2	FIFLLFLTL	0.59/0.56257776
70	1017	GQSKRVDFC	0.99/0.15997079
<b>71</b>	<b>802</b>	<b>DLLFNKVTI</b>	<b>0.01/1.1189541</b>
72	226	INITNFRAI	0.81/0.31891554
73	658	ASYHTVSLL	0.01/1.1156833
74	554	DVSDFTDSV	0.11/1.0096042
75	667	RSTSQKSIV	0.98/0.13520779
<b>76</b>	<b>828</b>	<b>ARDLICAQK</b>	<b>0.18/0.93408287</b>
77	1106	GNCDVVIGI	0.01/1.1035891
78	764	FAQVKQMYK	0.80/0.30544036
79	1177	ESLIDLQEL	0.60/0.50136368
80	431	TSTGNYNYK	0.48/0.61544296
81	1069	AYFPREGVF	0.44/0.64684427
82	981	GRLQSLQTY	0.91/0.17547758
83	1073	REGVVFVNG	0.79/0.29511842

84	323	CPFGEVFNA	0.58/0.49893355
85	1002	ASANLAATK	0.81/0.26402875
86	953	GAISSVLND	1.00/0.06918826
87	122	NVVIRACNF	0.26/0.80007167
88	40	VYYPDEIFR	0.76/0.29961556
89	759	NTREVFAQV	0.06/0.9893323
90	420	VLAWNTRNI	0.23/0.81608829
91	490	GYQPYRVVV	0.14/0.90122156
92	763	VFAQVKQMY	0.99/0.049718937
93	389	VKGDDVRQI	0.78/0.25960269
94	938	AQALNTLVK	0.41/0.62945518
<b>95</b>	<b>885</b>	<b>AYRFNGIGV</b>	<b>0.00/1.0390298</b>
96	125	IRACNFELC	0.80/0.23153875
97	68	GFHTINHTF	0.47/0.55733551
98	1119	VYDPLQPEL	0.43/0.59484561
99	753	AAEQDRNTR	0.46/0.55838722
100	760	TREVFAQVK	0.60/0.41458855
101	998	AEIRASANL	0.99/0.017047345
102	733	NLLQYGSF	0.93/0.063908892
103	289	SVKSFEIDK	0.17/0.82158984
104	524	CVNFNFNGL	0.01/0.9787292
105	1163	KEIDRLNEV	0.03/0.95542235
106	1103	FVSGNCDVV	0.41/0.57456887
107	306	RVVPSGDVV	0.71/0.26777445
<b>108</b>	<b>1190</b>	<b>QYIKWPWYV</b>	<b>/0.975564</b>
109	332	TKFPSVYAW	0.09/0.88229049
110	646	YECDIPIGA	0.35/0.62181879
111	255	GYLKPTTFM	0.70/0.27103346
112	1083	SWFITQRNF	0.99/-0.020811536
113	832	ICAQKFENGL	1.00/-0.033742263
114	829	RDLICAQKF	0.97/-0.0040749685
115	897	VLZENQKQI	0.78/0.18363948
116	1172	AKNLNESLI	0.00/0.96099358
117	799	FIEDLLFNK	0.95/0.0075979248
118	1033	SFPQAAPHG	0.86/0.095792957
119	101	WVFGSTMNN	0.81/0.14346986
120	109	NKSQSVIII	0.55/0.39016662
121	688	AYSNNTIAI	0.19/0.74549251

122	82	PFKDGIFYFA	0.90/0.029161163
123	780	GGFNFSQIL	0.71/0.21884931
124	707	EVMPVSMAM	0.22/0.70548911
<b>125</b>	<b>933</b>	<b>VVNQNAQAL</b>	<b>0.14/0.78381232</b>
126	1020	KRVDFCGKG	0.98/-0.059610779
127	437	NYKYRYLRH	0.27/0.64150725
128	595	AVLYQDVNC	0.87/0.038606135
129	229	TNFRAILTA	0.44/0.46188632
130	132	LCDNPFFAV	0.44/0.46102711
131	76	FGNPVIPFK	0.78/0.12097698
132	1039	PHGVVFLHV	0.93/-0.030721353
133	737	QYGSFCTQL	0.16/0.73586609
134	174	EKSGNFKHL	0.00/0.89543681
135	908	QFNKAISQI	0.47/0.4236368
136	1165	IDRLNEVAK	0.30/0.59035876
137	436	YNYKYRYLR	0.88/0.0067398286
138	457	NVPFSPDGK	0.80/0.08543008
139	971	AEVQIDRLI	0.73/0.1550506
140	371	ATKLNDLCF	0.57/0.31481201
141	300	YQTSNFRVV	0.06/0.8233986
142	716	TSVDCNMYI	0.23/0.65244227
143	555	VSDFTDSVR	0.87/0.0086210083
144	543	KRFQPFQQF	0.35/0.52788471
145	308	VPSGDVVRF	0.35/0.52506127
146	1089	RNFFSPQII	0.41/0.46333694
147	884	MAYRFNGIG	0.88/-0.0076877628
148	1166	DRLNEVAKN	0.98/-0.11756115
149	996	RAAEIRASA	0.72/0.13858567
150	703	SITTEVMPV	0.57/0.28665971
151	693	TIAIPTNFS	0.93/-0.07363762
<b>152</b>	<b>1036</b>	<b>QAAPHGVVF</b>	<b>0.18/0.67508934</b>
153	52	LYLTQDLFL	0.70/0.14707418
154	1154	INASVVNIQ	0.79/0.056617765
155	117	INNSTNVVI	0.65/0.1916925
156	519	LIKNQCVNF	0.55/0.28881045
157	353	SVLYNSTFF	0.88/-0.044011066
158	807	KVTLADAGF	0.99/-0.15789446
159	779	FGGFNFSQI	0.00/0.83156397

160	862	GTATAGWTF	0.39/0.44146214
161	727	DSTECANLL	0.25/0.58096964
162	969	VEAEVQIDR	0.29/0.53761645
163	944	LVKQLSSNF	0.97/-0.15274303
164	44	DEIFRSDTL	0.28/0.53527504
165	589	NASSEVAVL	0.31/0.50025079
166	625	GNNVFQTQA	0.94/-0.13078131
167	686	SIAYSNNTI	0.77/0.038400848
168	279	SQNPLAELK	0.00/0.80511986
169	58	LFLPFYSNV	0.29/0.51471164
170	182	LREFVFKNK	0.28/0.52209339
171	1176	NESLIDLQE	0.98/-0.17943301
172	297	KGIYQTSNF	0.66/0.13239
173	256	YLKPTTFML	0.00/0.78015806
174	889	NGIGVTQNV	0.17/0.60456213
175	668	STSQKSIVA	0.94/-0.16673334
176	208	DLPSGFNTL	0.05/0.72145971
177	94	KSNVVRGWV	0.09/0.68135937
178	1150	DISGINASV	0.80/-0.028901047
179	433	TGNYNYKYR	0.02/0.74372612
180	1126	ELDSFKEEL	0.00/0.76364207
181	1026	GKGYHLMSF	1.00/-0.23905486
182	769	QMYKTPTLK	0.73/0.028274204
183	1145	DVDLGDISG	0.62/0.13059576
184	358	STFFSTFKC	0.81/-0.061051474
185	230	NFRAILTAF	0.18/0.56811097
186	443	LRHGKLRPF	0.22/0.51348983
187	1037	AAPHGVVFL	0.00/0.73228057
188	620	RIYSTGNNV	0.00/0.73177713
189	1024	FCGKGYHLM	0.70/0.029800619
190	95	SNVVRGWVF	0.57/0.15189833
191	791	PLKPTKRSF	0.93/-0.21777967
192	1151	ISGINASVV	0.14/0.56782952
193	1159	VNIQKEIDR	0.97/-0.26534154
194	880	FAMQMAYRF	0.84/-0.13988547
195	501	FELLNAPAT	0.51/0.1899194
196	785	SQILPDPLK	0.97/-0.27104059
197	494	YRVVVSFE	0.87/-0.17477017
198	621	IYSTGNNVF	0.21/0.48460579



199	1056	NFTTAPAIC	0.99/-0.29928407
200	131	ELCDNPFFA	0.02/0.66894397
201	761	REVFAQVKQ	0.80/-0.11119255
202	149	HTMIFDNAF	0.02/0.66053968
203	935	NQNAQALNT	0.99/-0.31166887
204	377	LCFSNVYAD	0.97/-0.29241058
205	407	DYNYKLPDD	0.96/-0.28377689
206	921	TTTSTALGK	0.60/0.073904519
207	839	GLTVLPPLL	0.67/0.0030849041
208	294	EIDKGIYQT	0.99/-0.32034458
209	86	GIYFAATEK	0.00/0.66826612
210	47	FRSDTLYLT	0.94/-0.27964059
211	503	LLNAPATVC	0.93/-0.2705496
212	118	NNSTNVVIR	0.00/0.6543242
213	441	RYLRHGKLR	0.68/-0.033318088
214	475	YWPLNDYGF	0.99/-0.34362042
215	34	TSSMRGVYY	0.00/0.64601526
216	1081	GTSWFITQR	0.01/0.63397739
217	694	IAIPTNFSI	0.00/0.64266886
218	153	FDNAFNCTF	0.71/-0.0684593
219	805	FNKVTLADA	0.95/-0.30930826
220	903	KQIANQFNK	0.06/0.58043944
221	498	VLSFELLNA	0.02/0.61958251
222	17	DRCTTFDDV	0.22/0.41115954
223	278	CSQNPLAEL	0.00/0.62904085
224	1049	YVPSQERNF	0.00/0.62810603
225	577	SFGGVSVIT	0.72/-0.092513133
226	1107	NCDVVIGII	0.04/0.58602062
227	331	ATKFPSVYA	0.03/0.59209716
228	315	RFPNITNLC	0.61/0.010634901
229	1109	DVVIGIINN	0.25/0.3701283
230	1100	DNTFVSGNC	0.72/-0.1022437
231	491	YQPYRVVVL	0.02/0.5965429
232	1164	EIDRLNEVA	0.97/-0.35660241
233	1244	EPVLKGVKL	/0.61158557
234	729	TECANLLLQ	0.88/-0.27126781
235	473	NCYWPLNDY	0.17/0.43703774
236	31	TQHTSSMRG	0.85/-0.24366779
237	858	ALVSGTATA	0.06/0.54390801

238	373	KLNDLCFSN	0.00/0.59853344
239	746	NRALSGIAA	0.04/0.5554449
240	464	GKPCTPPAL	0.02/0.56591254
241	516	STDLIKNQC	0.08/0.50549987
<b>242</b>	<b>1038</b>	<b>APHGVVFLH</b>	<b>0.01/0.57504297</b>
243	613	DQLTPAWRI	0.15/0.43459649
244	220	FKLPLGINI	0.00/0.57760919
245	106	TMNNKSQSV	0.83/-0.25421526
246	705	TTEVMPVSM	0.62/-0.045282362
247	815	FMKQYGECL	0.03/0.54054092
248	598	YQDVNCTDV	0.00/0.56479598
249	1067	GKAYFPREG	0.93/-0.36672025
250	154	DNAFNCTFE	0.98/-0.41906956
251	887	RFNGIGVTQ	0.76/-0.2020951
252	1115	INNTVYDPL	0.73/-0.17409328
253	285	ELKCSVKSF	0.00/0.55424931
254	1203	IAGLIAIVM	/0.5530776
255	584	ITPGTNASS	0.99/-0.44178811
256	573	ISPCSFGGV	0.74/-0.19370372
257	412	LPDDFMGCV	0.11/0.43347787
258	397	IAPGQTGVI	0.03/0.5118819
259	522	NQCVNFNFN	0.52/0.021830229
260	396	QIAPGQTV	0.00/0.54060927
261	657	CASYHTVSL	0.57/-0.03478087
262	352	YSVLYNSTF	0.86/-0.32544177
263	383	YADSFVVKG	0.08/0.4540279
264	1000	IRASANLAA	0.60/-0.067413399
265	1018	QSKRVDFCG	0.99/-0.45968019
266	699	NFSISITTE	0.77/-0.24155566
267	1095	QIITTDNTF	0.83/-0.30217935
268	363	TFKCYGVSA	0.66/-0.13507416
269	398	APGQTGVIA	0.02/0.50481136
270	489	IGYQPVRVV	0.02/0.4999939
271	176	SGNFKHLRE	0.94/-0.42435894
272	1208	AIVMVTILL	/0.51235881
273	357	NSTFFSTFK	0.05/0.46101123
274	376	DLCFSNVYA	0.10/0.40826713
275	495	RVVVLSEFEL	0.13/0.37790287
276	134	DNPFFAVSK	0.00/0.50568662

277	523	QCVNFNFNG	0.88/-0.3771993
<b>278</b>	<b>1191</b>	<b>YIKWPWYVW</b>	<b>/0.50188623</b>
279	808	VTLADAGFM	0.79/-0.28951613
280	1078	VFNGTSWFI	0.01/0.48775526
281	1197	YVWLGFIAG	/0.49768537
282	778	YFGGFNFSQ	0.82/-0.32690109
283	172	VSEKSGNFK	0.80/-0.30778547
284	704	ITTEVMPVS	0.86/-0.36799302
285	1155	NASVVNIQK	0.00/0.48923419
286	732	ANLLQYGS	0.73/-0.24101658
287	736	LQYGSFCTQ	0.71/-0.22195096
288	505	NAPATVCGP	0.77/-0.28444478
289	324	PFGEVFNAT	0.87/-0.38873589
290	875	ALQIPFAMQ	0.99/-0.5129913
291	1206	LIAIVMTI	/0.47352128
292	92	TEKSNVVRG	0.02/0.44950188
293	849	DDMIAAYTA	0.54/-0.071829492
294	32	QHTSSMRGV	0.86/-0.39185733
295	214	NTLKPIFKL	0.05/0.4134994
296	430	ATSTGNVNY	0.00/0.46341048
297	1207	IAIVMVTIL	/0.45549214
<b>298</b>	<b>901</b>	<b>NQKQIANQF</b>	<b>0.05/0.40289272</b>
299	860	VSGTATAGW	0.88/-0.43015748
300	13	GSDLDRCTT	0.95/-0.50143335
301	852	IAAYTAALV	0.60/-0.16000632
302	302	TSNFRVPS	0.59/-0.15349161
303	895	QNVLYENQK	0.64/-0.20458317
304	820	GECLGDINA	0.00/0.42754106
305	497	VVLSFELLN	0.76/-0.33640163
306	334	FPSVYAWER	0.02/0.40036043
307	936	QNAQALNTL	0.01/0.40665469
308	520	IKNQCVNFN	0.81/-0.39497434
309	626	NNVFQTQAG	0.97/-0.55874967
310	455	ISNVPFSPD	0.99/-0.57920428
311	152	IFDNAFNCT	0.65/-0.24184003
312	676	AYTMSLGAD	0.96/-0.55566783
313	846	LLTDDMIAA	0.54/-0.14057689
314	649	DIPIGAGIC	0.80/-0.405829
315	299	IYQTSNFRV	0.00/0.39102148

316	1032	MSFPQAAPH	0.83/-0.43993695
<b>317</b>	<b>1201</b>	<b>GFIAGLIAI</b>	<b>/0.38830748</b>
318	673	SIVAYTMSL	0.01/0.37599299
319	1246	VLKGVKLHY	/0.38501663
320	200	YQPIDVVRD	0.92/-0.54851618
321	212	GFNTLKPIF	0.59/-0.22225368
322	15	DLDRCTTFD	0.99/-0.62317732
323	957	SVLNDILSR	0.51/-0.14367289
324	838	NGLTVLPPL	0.73/-0.36548633
325	1019	SKRVDFCGK	0.56/-0.19627116
326	178	NFKHLREFV	0.00/0.36348113
327	251	AYFVGYLKP	0.79/-0.43021312
328	979	ITGRLQSLQ	0.60/-0.24138811
329	446	GKLRPFERD	0.97/-0.62008704
330	1096	IITDNTFV	0.58/-0.23155762
331	892	GVTQNVLYE	0.66/-0.31273301
332	183	REFVFKNKD	0.97/-0.62576758
333	163	YISDAFSLD	0.69/-0.34672464
334	803	LLFNKVTLA	0.64/-0.29799872
335	601	VNCTDVSTA	0.85/-0.50938353
336	653	GAGICASYH	0.64/-0.30590714
337	781	GFNFSQILP	0.78/-0.45454639
338	193	FLYVYKGYQ	0.97/-0.64488319
339	343	KKISNCVAD	0.81/-0.48534288
340	186	VFKNKDGFL	0.74/-0.42051083
341	215	TLKPIFKLP	0.87/-0.5524945
342	3	IFLLFLTTLT	0.99/-0.6749784
343	257	LKPTTFMLK	0.52/-0.21193809
344	587	GTNASSEVA	0.89/-0.58466495
345	59	FLPFYSNVT	0.56/-0.25567868
346	962	ILSRDKVE	0.98/-0.67803869
347	662	TVSLLRSTS	0.61/-0.31640394
348	123	VVIRACNFE	0.92/-0.62889917
349	1113	GIINNTVYD	0.79/-0.50753657
350	731	CANLLLQYG	0.63/-0.35212059
351	812	DAGFMKQYG	0.91/-0.63706258
352	721	NMYICGDST	0.94/-0.66739372
353	976	DRLITGRLQ	0.95/-0.68422966
354	270	GTITDAVDC	0.90/-0.63452242

355	652	IGAGICASY	0.96/-0.69611784
356	1182	LQELGKYEQ	0.91/-0.64746161
357	881	AMQMAYRFN	0.89/-0.62842562
358	1074	EGVVFVNGT	0.81/-0.55295901
359	712	SMAKTSVDC	0.97/-0.73165598
360	750	SGIAAEQDR	0.97/-0.73414792
361	641	HVDTSYECD	0.79/-0.55453632
362	470	PALNCYWPL	0.98/-0.74879714
363	772	KTPTLKYFG	0.87/-0.63912731
364	169	SLDVSEKSG	0.80/-0.56984781
365	1048	TYVPSQERN	0.60/-0.37130228
366	140	VSKPMGTQT	0.59/-0.36264949
367	844	PPLLTDDMI	0.93/-0.7035536
368	968	KVEAEVQID	0.72/-0.52449453
369	85	DGIYFAATE	0.81/-0.61792411
370	1011	MSECVLGQS	0.95/-0.75851111
371	480	DYGFYTTTG	0.65/-0.45976869
372	571	LDISPCSFG	0.90/-0.71313511
373	605	DVSTAIHAD	0.73/-0.55196068
374	61	PFYSNVTGF	0.58/-0.40289454
375	428	IDATSTGNY	0.73/-0.55787431
376	878	IPFAMQMAY	0.83/-0.66172733
377	137	FFAVSKPMG	0.98/-0.8140195
378	856	TAALVSGTA	0.62/-0.4559563
379	542	SKRFQPFQQ	0.77/-0.60706626
380	196	VYKGYQPID	0.87/-0.71193255
381	1064	CHEGKAYFP	0.98/-0.82245574
382	394	VRQIAPGQT	0.97/-0.83689108
383	675	VAYTMSLGA	0.70/-0.57061653
384	546	QPFQQFGRD	0.97/-0.84322976
385	456	SNVPFSPDG	0.69/-0.56400599
386	709	MPVSMAKTS	0.93/-0.81254283
387	752	IAAEQDRNT	0.63/-0.51852776
388	633	AGCLIGAEH	0.63/-0.52021926
389	775	TLKYFGGFN	0.80/-0.70419167
390	816	MKQYGECLG	0.91/-0.81836769
391	88	YFAATEKSN	0.86/-0.77045812
392	191	DGFLYVYKG	0.54/-0.45379858
393	166	DAFSLDVSE	0.98/-0.89696565

394	1161	IQKEIDRLN	0.73/-0.6602502
395	654	AGICASYHT	0.78/-0.72217748
396	114	VIIINNSTN	0.75/-0.69732738
397	931	QDVVNQNAQ	0.68/-0.62903621
398	264	LKYDENGTI	0.94/-0.89676427
399	585	TPGTNASSE	0.79/-0.74784998
400	661	HTVSLLRST	0.73/-0.69219799
401	317	PNITNLCPF	0.75/-0.71932082
402	484	YTTTGIGYQ	0.93/-0.90376755
403	408	YNYKLPDDF	0.85/-0.83580025
404	143	PMGTQTHTM	0.99/-0.99044343
405	630	QTQAGCLIG	0.76/-0.76196468
406	204	DVVRDLPSG	0.85/-0.85876101
407	23	DDVQAPNYT	0.78/-0.78924934
408	271	TITDAVDCS	0.64/-0.65928597
409	783	NFSQILPDP	0.95/-0.97250767
410	855	YTAALVSGT	0.54/-0.56296968
411	643	DTSYECDIP	0.95/-0.97409968
412	385	DSFVVKGDD	0.97/-1.0069969
413	5	LLFLTLTSG	0.63/-0.67296897
414	720	CNMYICGDS	0.83/-0.88885802
415	26	QAPNYTQHT	0.74/-0.80155471
416	24	DVQAPNYTQ	0.60/-0.68396787
417	135	NPFFAVSKP	0.60/-0.6988653
418	532	LTGTGVLTP	0.65/-0.77063935
419	392	DDVRQIAPG	0.62/-0.75249678
420	339	AWERKKISN	0.93/-1.0690283
421	69	FHTINHTFG	0.67/-0.81235071
422	948	LSSNFGAIS	0.66/-0.81528605
423	549	QQFGRDVSD	0.93/-1.0948516
424	680	SLGADSSIA	0.96/-1.1350953
425	786	QILPDPLKP	0.73/-0.90576926
426	527	FNFNGLTGT	0.91/-1.0862641
427	1094	PQIITDNT	0.59/-0.79950416
428	813	AGFMKQYGE	0.64/-0.85601039
429	485	TTTGIGYQP	0.73/-0.97038176
430	128	CNFELCDNP	0.68/-0.92957098
431	1051	PSQERNFTT	0.56/-0.81950656
432	564	DPKTSEILD	0.55/-0.8202666

433	893	VTQNVLYEN	0.66/-0.93287129
434	1045	LHVTYVPSQ	0.99/-1.2654502
435	1140	NHTSPDVDL	0.55/-0.84283319
436	87	IYFAATEKS	0.54/-0.84065819
437	1173	KNLNEIDL	0.77/-1.0715629
438	290	VKSFEIDKG	0.76/-1.0690506
439	103	FGSTMNNKS	0.96/-1.2927739
440	327	EVFNATKFP	0.66/-0.99992216
441	259	PTTFMLKYD	0.90/-1.241965
442	782	FNFSQILPD	0.62/-0.9620474
443	273	TDAVDCSQN	0.60/-0.9602398
444	788	LPDPLKPTK	0.66/-1.0655041
445	1004	ANLAATKMS	0.87/-1.288379
446	606	VSTAIHADQ	0.79/-1.2439519
447	9	TLTSGSDL	0.52/-0.97728736
448	170	LDVSEKSGN	0.85/-1.3174187
449	318	NITNLCPPG	0.94/-1.4393988
450	945	VKQLSSNFG	0.93/-1.4346284
451	203	IDVVRDLPS	0.96/-1.4851088
452	184	EFVFKNKDG	0.68/-1.2268172
453	599	QDVNCTDVS	0.97/-1.5251249
454	458	VPFSPDGKP	0.94/-1.4954966
455	1028	GYHLMSFPQ	0.61/-1.2157184
456	7	FLTLTSGSD	0.87/-1.481056
457	580	GVSVITPGT	0.69/-1.3218354
458	1122	PLQPELDSF	0.55/-1.1856682
459	6	LFLTLTSGS	0.94/-1.5786607
460	1116	NNTVYDPLQ	0.85/-1.492597
461	513	PKLSTDLIK	0.60/-1.2689348
462	636	LIGAEHVDI	0.86/-1.5903752
463	916	IQESLTTTS	0.80/-1.5617904
464	367	YGVSA TKLN	0.96/-1.7341912
465	663	VSLLRSTSQ	0.53/-1.364465
466	966	LDKVEAEVQ	0.67/-1.5533956
467	767	VKQMYKTPT	0.96/-2.0682375
468	1120	YDPLQPELD	0.59/-1.7567044
469	28	PNYTQHTSS	0.66/-2.1108841
MERS-CoV			

Viral protein name	Rank	Start position	Sequence	Score ANN/SVM
<b>Spike</b>	1	751	RSVPGEMRL	0.95/1.0687206
	2	336	AIDCGFNDL	0.98/0.90784894
	3	1	MIHSVFLLM	0.97/0.91473499
	4	1032	ALSKLASEL	0.92/0.95686386
	5	1278	ESYIDLKEL	0.91/0.94986528
	6	630	FVYDAYQNL	0.84/0.98684097
	7	498	SYINKCSRL	0.47/1.3067989
	8	270	YVDLYGGNM	0.95/0.79768027
	9	112	FANGFVVRI	0.45/1.2939448
	10	1244	TSIPNFGSL	0.92/0.77792369
	11	845	SVRNLFASV	0.97/0.70106372
	12	772	QLNSSYFKL	0.97/0.67687168
	13	823	EYGQFCSKI	0.93/0.69441063
	14	473	FSNPTCLIL	0.87/0.74637331
	15	307	RKAWAAFYV	0.84/0.7381001
	16	292	YYSIIPHSI	0.79/0.77662825
	17	282	ATLPVYDTI	0.07/1.4867292
	18	1065	AQIDRLING	0.77/0.72729886
	19	1070	LINGRLTTL	0.72/0.7584431
	20	1074	RLTTLNAFV	0.20/1.2783667
	21	1201	NTKYVAPQV	0.91/0.55793045
	22	97	PQKLFVANY	0.70/0.7543466
	23	1254	QINTTLLDL	0.81/0.63927539
	<b>24</b>	<b>975</b>	<b>SIFYRLNGV</b>	<b>0.98/0.45706474</b>
	25	1166	APVNGYFIK	0.99/0.43439606
	<b>26</b>	<b>1025</b>	<b>AVNNNAQAL</b>	<b>0.33/1.0919338</b>
	27	595	KIASQLGNC	0.94/0.48078322
	28	317	KLQPLTFL	0.21/1.1941265
	29	434	ASNCYSSLI	0.79/0.61355361
	30	1197	ITSLNTKYV	0.92/0.48237559
	31	31	IEVDIQQTF	0.96/0.44123869
	32	346	QLHCSYESF	0.98/0.41824276
	33	278	MFQFATLPV	0.94/0.45179384
	34	303	IQSDRKAWA	1.00/0.3750027
	35	226	NSFKEYFNL	0.55/0.81163751
	36	737	ALPDTPSTL	0.06/1.2937612
	<b>37</b>	<b>1128</b>	<b>VNAPNGLYF</b>	<b>0.60/0.74121655</b>



38	1141	YYPSNHIEV	0.12/1.2183088
<b>39</b>	<b>1275</b>	<b>ALNESYIDL</b>	<b>0.00/1.3214488</b>
40	3	HSVFLLMFL	0.33/0.97938501
41	242	TYNITEDEI	0.52/0.78788058
42	462	GPISQFNYK	0.73/0.57779357
43	624	GVRQQRFBVY	0.96/0.33001584
44	1073	GRLTTLNAF	0.82/0.46949067
45	972	FAQSIFYRL	0.31/0.97721311
46	1002	NQALGAMQT	0.86/0.4220723
47	42	KTWPRPIDV	0.30/0.97309465
48	419	SVNDFTCSQ	0.64/0.62272262
49	1215	STNLPPPLL	0.72/0.53785544
50	156	FSDGKMGRF	0.88/0.37258124
51	703	TYGPLQTPV	0.73/0.51648442
52	631	VYDAYQNLV	0.75/0.46216708
53	787	SFGVTQEYI	0.83/0.36450572
54	1258	TLLDLTYEM	0.13/1.0607423
55	141	RKIYPAFML	0.93/0.26058818
56	433	IASNCYSSL	0.99/0.18701988
57	900	IADPGYMQG	0.90/0.27478176
58	164	FFNHTLVLL	0.00/1.1687357
59	221	RNASLNSFK	0.88/0.28872588
60	881	TGSRARSAS	0.96/0.20003002
61	94	GTTPQKLFV	0.85/0.30586214
62	328	SVDGYIRRA	0.62/0.52897679
63	576	QYGTDTNSV	0.99/0.15513332
64	395	QVYNFKRLV	0.65/0.49190588
65	25	SVKSACIEV	0.14/1.0016961
66	826	QFCSKINQA	0.94/0.19846214
67	641	YYSDDGNYV	0.00/1.1359411
68	918	ASARDLICA	0.77/0.36455438
69	1049	ASIGDIIQR	0.92/0.21269236
70	699	RRDSTYGPL	0.86/0.26840132
71	730	PLGQSLCAL	0.82/0.30213022
72	841	RQDDSVRNL	0.06/1.0596324
73	570	GFGITVQYG	0.70/0.41848594
74	220	NRNASLNSF	0.38/0.73628472
<b>75</b>	<b>920</b>	<b>ARDLICAQY</b>	<b>0.14/0.97234577</b>
76	92	ATGTTPQKL	0.67/0.43882611

77	655	VSVPVSVIY	0.50/0.6044505
78	1000	KFNQALGAM	0.23/0.86456217
79	1268	SLQQVVKAL	0.01/1.0736436
80	266	FSSRYVDLY	0.67/0.40983872
81	1309	ALALCVFFI	/1.0797517
82	1050	SIGDIIQRL	0.72/0.35948984
83	1116	FCGQGTHIV	0.85/0.22624257
84	331	GYIRRAIDC	0.45/0.62562951
85	136	TSATIRKIY	0.85/0.22399459
86	997	IANKFNQAL	0.55/0.5224597
87	710	PVGCVLGLV	0.78/0.29204099
88	480	ILATVPHNL	0.32/0.7473596
89	580	DTNSVCPKL	0.53/0.53162014
90	78	QGDHGDMYV	0.64/0.41941369
<b>91</b>	<b>892</b>	<b>DLLFDKVTI</b>	<b>0.56/0.49242704</b>
92	933	KVLPPLMDV	0.69/0.35876353
93	535	WEDGDYYRK	0.81/0.22653775
94	627	QQRFBYDAY	0.84/0.19222925
95	163	RFFNHTLVL	0.91/0.1208001
96	404	FTNCNYNLT	0.98/0.046119296
97	232	FNLRNCTFM	0.05/0.97291655
98	11	LLTPTESYV	0.71/0.3081859
99	263	VHLFSSRYV	0.90/0.10626798
100	310	WAAFYVYKL	0.13/0.87561177
101	698	KRRDSTYGP	0.97/0.031699774
102	949	SSLLGSIAG	0.97/0.023813555
<b>103</b>	<b>1291</b>	<b>YYNKWPWYI</b>	<b>/0.98943415</b>
104	827	FCSKINQAL	0.82/0.1685718
105	969	AIPFAQSIF	0.72/0.26687558
106	262	GVHLFSSRY	0.43/0.55464322
107	786	FSFGVTQEY	0.97/0.0096269755
108	968	AAIPFAQSI	0.10/0.8750469
109	483	TVPHNLTTI	0.22/0.75473192
110	866	GGDFNLTL	0.31/0.66065832
111	108	DVKQFANGF	0.00/0.96649715
112	459	SSAGPISQF	0.65/0.31636956
113	444	DYFSYPLSM	0.18/0.78615375
114	1122	HIVSFVVNA	0.15/0.81293353
115	50	VSKADGHIY	0.03/0.9329199

116	1163	NCIAPVNGY	0.73/0.23244441
117	453	KSDLVSSA	0.84/0.12022072
118	522	QYSPCVSIV	0.01/0.94906953
119	1047	ISASIGDII	0.99/-0.036092964
120	179	LLRAFYCIL	0.01/0.94087367
121	1185	SYTGSSFYA	0.87/0.080057043
122	391	GTPPQVYNF	0.05/0.8975718
123	59	PQGRTYSNI	0.91/0.035446458
124	1171	YFIKTNNTR	0.95/-0.0050865908
125	887	RSAIEDLLF	0.90/0.04378359
126	355	DVESGVYSV	0.03/0.90428769
127	716	GLVNSSLFV	0.52/0.40571098
128	1230	DFQDELDEF	0.98/-0.059229733
129	1284	KELGNYTTY	/0.92016098
130	526	CVSIVPSTV	0.00/0.91874491
131	213	DCSDGNYNR	0.57/0.34873258
132	1179	RIVDEWSYT	0.76/0.15226026
133	1234	ELDEFFKNV	0.02/0.8921197
134	752	SVPGEMRLA	0.97/-0.063279431
135	674	FGSVACEHI	0.98/-0.081708659
136	982	GVGITQQVL	0.39/0.50294203
137	1332	KCNRCCDRY	/0.89207171
138	304	QSDRKAWAA	0.50/0.38817415
139	813	GFQKCEQLL	0.56/0.32803925
140	4	SVFLLMFL	0.01/0.87744856
<b>141</b>	<b>1115</b>	<b>GFCGQGTHI</b>	<b>0.97/-0.086697044</b>
<b>142</b>	<b>1302</b>	<b>GFIAGLVAL</b>	<b>/0.88200538</b>
143	857	QSSPIIPGF	0.98/-0.10086419
144	178	TLLRAFYCI	0.01/0.86649455
145	1012	FTTTNEAFR	0.85/0.02424523
146	582	NSVCPKLEF	0.65/0.22269833
147	1241	NVSTSIPNF	0.47/0.40157321
148	548	LEGGGWLVA	0.37/0.49860677
149	1035	KLASELSNT	0.96/-0.092672325
150	1120	GTHIVSFVV	0.70/0.16135788
151	56	IIYPQGRTY	0.86/0.00072868393
152	506	LLSDDRTEV	0.00/0.85876251

153	617	FQNCTAVGV	0.32/0.53059444
<b>154</b>	<b>1134</b>	<b>LYFMHVGYY</b>	<b>0.99/-0.14427586</b>
155	1126	FVVNAPNGL	0.90/-0.06198942
156	113	ANGFVVRIG	0.02/0.81652064
157	114	NGFVVRIGA	0.62/0.21548528
158	534	VWEDGDYYR	0.77/0.061463248
159	926	AQYVAGYKV	0.68/0.14881084
160	648	YYCLRACVS	0.99/-0.16373158
161	276	GNMFQFATL	0.17/0.65487679
162	18	YVDVGPDSV	0.51/0.31472512
163	540	YYRKQLSPL	0.45/0.36831902
164	1038	SELSNTFGA	0.44/0.3777823
165	499	YINKCSRL	0.83/-0.012687027
166	313	FYVYKLQPL	0.08/0.73686339
167	686	MSQYSRSTR	0.37/0.44470934
168	1165	IAPVNGYFI	0.01/0.80325124
169	49	DVSKADGII	0.83/-0.020093876
170	825	GQFCSKINQ	0.99/-0.18087676
171	563	MTEQLQMGF	0.91/-0.10586779
172	653	ACVSVPVSV	0.34/0.45748738
<b>173</b>	<b>977</b>	<b>FYRLNGVGI</b>	<b>0.00/0.79279512</b>
174	296	IPHSIRSIQ	0.90/-0.10977856
175	1147	IEVVSAYGL	0.05/0.73879841
176	690	SRSTRSMLK	0.51/0.27723927
177	1053	DIIQRLDVL	0.10/0.6839985
178	229	KEYFNLRNC	0.03/0.75375487
179	954	SIAGVGWTA	0.67/0.10986263
180	1227	TGIDFQDEL	0.07/0.70964031
181	692	STRSMLKRR	0.80/-0.021814712
182	1014	TTNEAFRKV	0.00/0.77732566
183	671	ATLFGSVAC	0.17/0.59396847
184	1236	DEFFKNVST	0.99/-0.22718534
185	327	FSVDGYIRR	0.00/0.76216554
186	491	ITKPLKYSY	0.52/0.24070756
187	1112	KRSGFCGQG	0.92/-0.1603229
188	442	ILDYFSYPL	0.88/-0.12223742
189	366	FEAKPSGSV	0.67/0.074974423
190	176	CGTLLRAFY	0.95/-0.2078755
191	231	YFNLRNCTF	0.12/0.62145635

192	295	IIPHSIRSI	0.53/0.21101408
193	706	PLQTPVGCV	1.00/-0.26141429
194	392	TPPQVYNFK	0.03/0.70677111
195	890	IEDLLFDKV	0.84/-0.10481566
196	1310	LALCVFFIL	/0.73482458
197	567	LQMGFGITV	0.71/0.02301018
198	1206	APQVTYQNI	0.03/0.70174242
199	1160	NPTNCIAPV	0.71/0.020766975
200	893	LLFDKVTIA	0.35/0.37940735
201	721	SLFVEDCKL	0.05/0.67843327
202	882	GSRSAISAI	0.29/0.43802917
203	626	RQQRFBVYDA	0.09/0.63256432
204	970	IPFAQSIFY	0.79/-0.068162709
205	1247	PNFGSLTQI	0.88/-0.16010176
206	922	DLICAQYVA	0.97/-0.27332673
<b>207</b>	<b>1130</b>	<b>APNGLYFMH</b>	<b>0.30/0.39359148</b>
208	299	SIRSIQSDR	0.82/-0.12991588
209	1293	NKWPWYIWL	/0.68175091
210	600	LGNCVEYSL	0.66/0.019552802
211	668	KTHATLFGS	0.88/-0.20124114
212	921	RDLICAQYV	0.01/0.66830014
213	1344	DLEPHKVHV	/0.67725876
214	647	NYYCLRACV	0.29/0.38221191
<b>215</b>	<b>993</b>	<b>NQKLIANKF</b>	<b>0.02/0.65017867</b>
216	245	ITEDEILEW	0.58/0.082462128
217	553	WLVASGSTV	0.01/0.64985351
218	1145	NHIEVVSAY	0.13/0.52891408
219	1158	AANPTNCIA	0.02/0.63833265
220	642	YSDDGNYYC	0.18/0.47774429
221	1249	FGSLTQINT	0.80/-0.14549239
222	844	DSVRNLFAS	0.60/0.05024281
223	766	HPIQVDQLN	0.96/-0.31041729
224	782	IPTNFSFGV	0.65/-0.0019764399
225	177	GTLLRAFYC	0.02/0.6217205
226	305	SDRKAWAAF	0.75/-0.112341
227	38	TFFDKTWPR	0.05/0.58761966
228	1016	NEAFRKVQD	0.83/-0.19275131
229	562	AMTEQLQMG	0.80/-0.16549054

230	720	SSLFVEDCK	0.55/0.083770077
231	678	ACEHISSTM	0.09/0.54096983
232	865	FGGDFNLT	0.72/-0.092457905
233	915	QGPASARDL	0.72/-0.092768774
234	294	SIIPHSIRS	0.57/0.057063443
235	77	YQGDHGDMY	0.84/-0.21452308
236	85	YVYSAGHAT	0.98/-0.3570909
237	57	IYPQGRYS	0.89/-0.26710895
238	780	LSIPTNFSF	0.64/-0.017340059
239	801	KVTVDCKQY	0.98/-0.35759757
240	861	IIPGFGGDF	0.56/0.061012449
241	533	TVWEDGDYY	0.66/-0.045226686
<b>242</b>	<b>1292</b>	<b>YNKWPWYIW</b>	<b>/0.60806956</b>
243	985	ITQQVLSEN	0.97/-0.36321819
244	140	IRKIYPAFM	0.04/0.5665451
245	556	ASGSTVAMT	0.17/0.4348019
246	615	GVFQNCTAV	0.04/0.56373656
247	962	AGLSSFAAI	0.08/0.51323156
248	253	WFGITQTAQ	0.85/-0.25945579
249	513	EVPQLVNAN	0.51/0.072816261
250	846	VRNLFASVK	0.09/0.49084835
251	1046	AISASIGDI	0.04/0.53748172
252	1078	LNAFVAQQL	0.70/-0.12423014
253	234	LRNCTFMYT	0.00/0.57528912
254	115	GFVVRIGAA	0.91/-0.33511276
255	707	LQTPVGCVL	0.75/-0.17520072
256	1266	MLSLQQVVK	0.08/0.49193623
257	1152	AYGLCDAAN	0.54/0.025994238
258	1335	RCCDRYEEY	/0.56468276
259	1132	NGLYFMHVG	0.91/-0.34904069
260	643	SDDGNYYCL	0.02/0.53931673
261	863	PGFGGDFNL	0.82/-0.26686725
262	1246	IPNFGSLTQ	0.87/-0.32157967
263	309	AWAAFYVYK	0.00/0.54566796
264	246	TEDEILEWF	0.05/0.4922968
265	376	EQAEGVECD	0.97/-0.42865706
266	676	SVACEHISS	0.91/-0.37144384
267	1303	FIAGLVALA	/0.53558793
268	1216	TNLPPPLL	0.87/-0.33564407

269	1125	SFVVNAPNG	0.85/-0.31576137
270	70	TYQGLFPYQ	0.64/-0.10616507
271	330	DGYIRRAID	0.99/-0.45820288
272	105	YSQDVKQFA	0.58/-0.052124143
273	10	FLLTPTESY	0.99/-0.46327628
274	396	VYNFKRLVF	0.13/0.39443369
275	878	SISTGSRSA	0.93/-0.40687571
276	406	NCNYNLTKL	0.07/0.45311362
277	390	SGTPPQVYN	0.99/-0.46850976
278	621	TAVGVRQQR	0.02/0.4994245
279	271	VDLYGGNMF	0.86/-0.34166406
280	1164	CIAPVNGYF	0.00/0.51460498
281	1069	RLINGRLTT	0.68/-0.16814007
282	1193	APEPITSLN	0.63/-0.12071683
283	794	YIQTTIQKV	0.63/-0.1275968
284	258	QTAQGVHLF	0.02/0.48126671
285	614	RGVFQNCTA	0.96/-0.45959761
286	1024	DAVNNNAQA	0.66/-0.16084347
287	913	MQQGPASAR	0.55/-0.056978444
288	1154	GLCDAANPT	0.93/-0.43989785
289	135	STSATIRKI	0.01/0.47961503
290	622	AVGVRQQRF	0.01/0.47770348
291	708	QTPVGCVLG	0.98/-0.49560177
292	822	REYGQFCSK	0.05/0.43280774
293	122	AAANSTGTV	0.73/-0.26108703
294	1342	EYDLEPHKV	/0.46745524
295	663	YDKETKTHA	0.98/-0.51502941
296	1204	YVAPQVITYQ	0.00/0.46243144
297	300	IRSIQSDRK	0.03/0.4298579
298	660	SVIYDKETK	0.67/-0.21073164
299	1311	ALCVFFILC	/0.45721592
300	943	MEAAYTSSL	0.01/0.44420198
301	749	SVRSVPGEM	0.01/0.44105273
302	1297	WYIWLGFIA	/0.44831422
303	217	GNYNRNASL	0.01/0.43741602
304	197	AGNSYTSFA	0.74/-0.29348105
305	953	GSIAGVGWT	0.86/-0.41798702
306	998	ANKFNQALG	0.03/0.41171683
307	95	TTPQKLFVA	0.99/-0.55128102

308	955	IAGVGWTAG	0.58/-0.14139627
309	565	EQLQMGFGI	0.89/-0.45194789
310	509	DDRTEVPQL	0.51/-0.07652848
311	802	VTVDCKQYV	0.01/0.42310575
312	628	QRFVYDAYQ	0.82/-0.38776991
313	764	FNHPIQVDQ	0.93/-0.50039409
314	161	MGRFFNHTL	0.00/0.42042943
315	665	KETKTHATL	0.04/0.37890249
316	501	NKCSRLSD	0.97/-0.55271012
317	22	GPDSVKSAC	0.61/-0.19345945
318	512	TEVPQLVNA	0.03/0.38606041
319	1007	AMQTGFTTT	0.90/-0.48650257
320	1245	SIPNFGSLT	0.72/-0.30830238
321	1041	SNTFGAISA	0.05/0.36159342
322	65	SNITITYQG	0.87/-0.46133046
323	555	VASGSTVAM	0.00/0.40780761
324	1330	KLKCNRCCD	/0.4029762
325	224	SLNSFKEYF	0.00/0.3972556
326	432	AIASNCYSS	0.55/-0.15764841
327	39	FFDKTWPRP	0.96/-0.56834753
328	416	SLFSVNDFT	0.66/-0.26923111
329	1129	NAPNGLYFM	0.00/0.3903336
330	645	DGNYYCLRA	0.58/-0.19163244
331	1173	IKTNNTRIV	0.01/0.37834217
332	964	LSSFAAIPF	0.83/-0.44408461
333	935	LPPLMDVNM	0.64/-0.25524073
334	1251	SLTQINTTL	0.00/0.38367782
335	1059	DVLEQDAQI	0.00/0.38097157
336	696	MLKRRDSTY	0.00/0.38077499
337	649	YCLRACVSV	0.01/0.37036168
338	495	LKYSYINKC	0.54/-0.1608164
339	485	PHNLTTITK	0.75/-0.37712782
340	188	EPRSGNHCP	0.92/-0.54962202
341	872	TLLEPVSIS	0.77/-0.41449786
342	1159	ANPTNCIAP	0.55/-0.19517041
343	1044	FGAISASIG	0.94/-0.60011359
344	165	FNHTLVLLP	0.93/-0.59351837
345	461	AGPISQFNY	0.53/-0.200378
346	465	SQFNYKQSF	0.90/-0.57048416



347	239	FMYTYNITE	0.67/-0.34847386
348	502	KCSRLLSDD	0.95/-0.63288187
349	334	RRIDCGFN	0.91/-0.60643944
350	959	GWTAGLSSF	0.77/-0.46872835
351	603	CVEYSLYGV	0.72/-0.42265218
352	52	KADGHIYPQ	0.71/-0.41343896
353	662	IYDKETKTH	0.96/-0.66858006
354	564	TEQLQMGFG	0.79/-0.50194902
355	588	LEFANDTKI	0.94/-0.65250566
356	399	FKRLVFTNC	0.62/-0.33996511
357	1022	VQDAVNNNA	0.56/-0.28132524
358	858	SSPIPGFG	1.00/-0.72965374
359	518	VNANQYSPC	0.99/-0.71992597
360	869	FNLTLLEPV	0.91/-0.64120938
361	289	TIKYYSIIP	0.71/-0.44173822
362	1086	LVRSESAAL	0.55/-0.28200055
363	1089	SESAALSAQ	0.68/-0.42090549
364	904	GYMQGYDDC	0.64/-0.38370308
365	1198	TSLNTKYVA	0.61/-0.35433589
366	974	QSIFYRLNG	0.53/-0.27617988
367	507	LSDDRTEVP	0.98/-0.72719503
368	1168	VNGYFIKTN	0.62/-0.36813429
369	439	SSLILDYFS	0.97/-0.72620275
370	281	FATLPVYDT	0.78/-0.54298364
371	209	TPATDCSDG	0.96/-0.728034
372	1279	SYIDLKELG	0.77/-0.53928964
373	940	DVNMEAAYT	0.54/-0.3102788
374	279	FQFATLPVY	0.94/-0.7221058
375	372	GSVVEQAEG	0.95/-0.73672015
376	597	ASQLGNCVE	0.81/-0.59946363
377	1256	NTLLDLTY	0.99/-0.78138739
378	40	FDKTWPRPI	0.58/-0.37239834
379	369	KPSGSVVEQ	0.68/-0.50061853
380	208	HTPATDCSD	0.94/-0.76122135
381	742	PSTLTPRS	0.96/-0.78323864
382	436	NCYSSLILD	0.94/-0.77100125
383	496	KYSYINKCS	0.80/-0.65796787
384	45	PRPIDVSKA	0.83/-0.6902416
385	186	ILEPRSGNH	0.52/-0.38503946

386	670	HATLFGSVA	0.56/-0.43188367
387	746	TPRSVRSVP	0.60/-0.47314795
388	650	CLRACVSVP	0.73/-0.60447888
389	1243	STSIPNFGS	0.85/-0.72722733
390	130	VIISPSTSA	0.69/-0.57102461
391	72	QGLFPYQGD	0.82/-0.71369352
392	592	NDTKIASQL	0.66/-0.55518204
393	5	VFLLMFLLT	0.96/-0.85793243
394	260	AQGVHLFSS	0.84/-0.74222638
395	348	HCSYESFDV	0.72/-0.63106176
396	249	EILEWFGIT	0.93/-0.85377882
397	17	SYVDVGPDS	0.70/-0.63134303
398	322	TFLLDfsVD	0.84/-0.77261535
399	1023	QDAVNNNAQ	0.94/-0.87554894
400	1153	YGLCDAANP	0.98/-0.91728324
401	575	VQYGTDTNS	0.73/-0.66954552
402	873	LLEPVSIST	0.92/-0.86139275
403	256	ITQTAQGVH	0.97/-0.91150271
404	1087	VRSESAALS	0.71/-0.65321967
405	353	SFDVESGVY	0.66/-0.61260149
406	1257	TTLLDLTYE	0.81/-0.76817195
407	897	KVTIADPGY	0.53/-0.4890011
408	1133	GLYFMHVGy	0.64/-0.60141762
409	447	SYPLSMKSD	0.68/-0.64937752
410	1013	TTTNEAFRK	0.55/-0.51966562
411	1202	TKYVAPQVT	0.90/-0.86994577
412	365	SFEAKPSGS	0.83/-0.81594474
413	103	ANYSQDVkQ	0.76/-0.75077778
414	190	RSGNHCPAG	0.66/-0.65395516
415	1143	PSNHIEVVS	0.95/-0.94428207
416	212	TDCSDGNYN	0.94/-0.93562508
417	134	PSTSATIRK	0.80/-0.80034738
418	1169	NGYFIKTNN	1.00/-1.0070648
419	290	IKYYSIIPH	0.55/-0.560356
420	727	CKLPLGQSL	0.51/-0.52707663
421	520	ANQYSPCVS	0.95/-0.97060464
422	928	YVAGYKVLP	0.72/-0.76315835
423	820	LLREYGQFC	0.78/-0.83582724
424	55	GIYPQGRT	0.95/-1.0115031

425	244	NITEDEILE	0.91/-0.97500076
426	952	LGSIAGVGW	0.76/-0.82524351
427	1228	GIDFQDELD	0.97/-1.0387297
428	992	ENQKLIANK	0.72/-0.79125318
429	849	LFASVKSSQ	0.58/-0.65550029
430	741	TPSTLTPRS	0.86/-0.93821969
431	695	SMLKRRDST	0.74/-0.83179842
432	1055	IQRLDVLEQ	0.68/-0.77302756
433	1226	STGIDFQDE	0.72/-0.82938184
434	312	AFYVYKLQP	0.88/-0.98950297
435	789	GVTQEYIQT	0.53/-0.64904728
436	729	LPLGQSLCA	0.73/-0.85747062
437	1177	NTRIVDEWS	0.61/-0.75893742
438	1103	VNECVKAQS	0.74/-0.90425647
439	109	VKQFANGFV	0.51/-0.6836031
440	864	GFGGDFNLT	0.55/-0.72887075
441	1232	QDELDEFFK	0.72/-0.90314068
442	652	RACVSVVPS	0.58/-0.76755349
443	571	FGITVQYGT	0.60/-0.80617775
444	474	SNPTCLILA	0.52/-0.73311341
445	76	PYQGDHGM	0.87/-1.0831585
446	452	MKSDLVSS	0.95/-1.1763039
447	538	GDYYRKQLS	0.58/-0.80837331
448	1220	PPLGNSTG	0.76/-0.99012338
449	1031	QALSKLASE	0.53/-0.76319456
450	1271	QVVKALNES	0.82/-1.0623915
451	604	VEYSLYGVS	0.53/-0.77657278
452	14	PTESYVDVG	0.59/-0.85025294
453	74	LFPYQGDHG	0.55/-0.81621351
454	1253	TQINTLLD	0.56/-0.83021605
455	166	NHTLVLLPD	0.87/-1.1479338
456	835	LHGANLRQD	0.88/-1.1623996
457	517	LVNANQYSP	0.55/-0.8345149
458	144	YPAFMLGSS	0.54/-0.83135736
459	566	QLQMGFGIT	0.65/-0.9534416
460	810	VCNGFQKCE	0.52/-0.83107607
461	228	FKEYFNLRN	0.72/-1.0312241
462	73	GLFPYQGDH	0.71/-1.024888
463	875	EPVSISTGS	0.82/-1.1351898

	464	932	YKVLPLMD	0.86/-1.1806539
	465	733	QSLCALPDT	0.74/-1.0695994
	466	147	FMLGSSVGN	0.70/-1.0410759
	467	664	DKETKTHAT	0.93/-1.2779392
	468	797	TTIQKVTVD	0.95/-1.3074728
	469	379	EGVECDFSP	0.69/-1.0566543
	470	1096	AQLAKDKVN	0.65/-1.0263296
	471	119	RIGAAANST	0.57/-0.9525637
	472	88	SAGHATGTT	0.78/-1.2030449
	473	463	PISQFNYKQ	0.62/-1.0638477
	474	545	LSPLEGGGW	0.86/-1.3108022
	475	1196	PITSLNTKY	0.98/-1.4437418
	476	455	DLSVSSAGP	0.91/-1.3920107
	477	374	VVEQAEGVE	0.94/-1.4272455
	478	195	CPAGNSYTS	0.85/-1.3433177
	479	909	YDDCMQQGP	0.69/-1.2028043
	480	490	TITKPLKYS	0.65/-1.1637635
	481	37	QTFFDKTWP	0.77/-1.2887773
	482	210	PATDCSDGN	0.80/-1.3294758
	483	658	PVSVIYDKE	0.70/-1.2543317
	484	895	FDKVTIADP	0.67/-1.2474127
	485	731	LGQSLCALP	0.80/-1.3968119
	486	120	IGAAANSTG	0.51/-1.1441705
	487	581	TNSVCPKLE	0.58/-1.2526687
	488	1033	LSKLASELS	0.82/-1.5518681
	489	725	EDCKLPLGQ	0.94/-1.7100772
	490	836	HGANLRQDD	0.63/-1.41272
	491	363	VSSFEAKPS	0.71/-1.51183
	492	638	LVGYYSDDG	0.74/-1.6071945
	493	573	ITVQYGTDT	0.61/-1.567401
	494	735	LCALPDTPS	0.78/-1.8348734
	495	739	PDTPSTLTP	0.85/-1.9485422
<b>HCoV-NL63</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	1	647	SLAGGITYV	0.50/1.7485472
	2	1121	SAPDGLLFL	0.71/1.4701125
	3	1220	TLQEFAQNL	0.78/1.270963
	4	1165	YSDNGVFRV	0.98/1.0340139

5	1024	ALNHLTSQL	0.99/0.96059106
6	1154	GYVLRQPNL	0.99/0.94512873
7	1169	GVFRVTSRV	0.92/0.97829107
8	590	YTIVGALYV	0.90/0.96558479
9	178	NYSCVFSVV	0.55/1.3061226
10	1175	SRVMFQPRL	0.90/0.94576564
11	1066	RLAALNAFV	0.73/1.1011045
12	372	TVREIVVAR	0.98/0.83200893
13	1280	YVDLKLLNR	0.71/1.0833337
14	953	ALQARLNYV	0.28/1.4734208
15	1185	VLSDFVQIY	0.22/1.5251199
16	522	FSIRYIYNR	0.97/0.74513173
17	1001	AEAIHTVTI	0.76/0.93207468
18	150	RLHLYNVTR	0.94/0.74561912
19	785	IVVDCATYV	0.61/1.0353148
20	1155	YVLRQPNLV	0.96/0.66182292
21	1069	ALNAFVSQV	0.23/1.3344931
22	565	TICFSTVAV	0.94/0.61771094
23	588	TSYTIVGAL	0.92/0.62464212
24	666	NVSTGNIFI	0.94/0.60457205
25	825	ETNDVSSML	0.93/0.60718551
26	89	IYVTNEIGL	0.16/1.357696
27	709	LQLPNFYVYV	0.02/1.4941587
28	1076	QVLNKYTEV	0.76/0.73136175
29	519	TSHFSIRYI	0.53/0.95241857
30	491	ASFGGSCYV	0.96/0.51462509
31	415	VAFATFVDV	0.68/0.78532103
32	2	KLFLILLVL	0.09/1.3728741
33	1230	KYVKPNFDL	0.58/0.8803956
34	988	ASFSSVND A	0.79/0.64883649
35	196	THNGRVVNY	0.88/0.5477679
36	20	NSNANLSML	0.28/1.1461754
37	501	KPHQVNISL	0.01/1.414479
38	399	AVNFNVTTA	0.00/1.4215403
39	345	SSQPYYCFI	0.94/0.48121853
40	878	FSKVVTSG L	0.91/0.50924148
41	212	GYTDNIFSV	0.85/0.56238908
42	174	KCYFNYS CV	0.71/0.67844509
43	872	ALEDLLFSK	0.84/0.53717421

44	83	DVNQYYIYV	0.01/1.3630435
45	152	HLYNVTRTF	0.86/0.5089841
46	60	ANGFFYIDV	0.67/0.69532891
47	955	QARLNYVAL	0.16/1.2027791
48	419	TFVDVLVNV	0.00/1.3436307
49	468	VLPETYVAL	0.01/1.3326933
50	143	TISGETVRL	0.93/0.41008745
51	144	ISGETVRLH	0.88/0.44742436
52	1098	CVKSQSNRY	0.71/0.60557634
53	1213	DYVDVNKTL	0.65/0.66393519
54	69	GNHRSAFAL	0.61/0.70391144
55	795	NGNPRCKNL	0.58/0.72936105
56	969	QENQKILAA	0.95/0.35162883
57	1182	RLPVLSDFV	0.69/0.60633387
58	690	SIIGAMTAV	0.49/0.80587575
59	952	LALQARLNY	0.91/0.38221057
60	1156	VLRQPNLVL	0.99/0.28647426
61	899	GLSIADLAC	0.98/0.29253223
62	890	DVDYKSCTK	0.86/0.41155337
63	1192	IYNCNVTFV	0.24/1.0262973
64	197	HNGRVVNYT	0.94/0.32625754
65	1300	WLIISVVFV	/1.2571117
66	1228	LPKYVKPNF	0.94/0.31373832
67	308	DNLKSGVIV	0.94/0.30886148
68	122	SSFDCIVNL	0.39/0.8545901
69	625	NNCTKYNIY	0.77/0.46053662
70	646	QSLAGGITY	0.99/0.23858537
71	194	VTTHNGRVV	0.94/0.28631213
72	31	GVPDNSSTI	0.07/1.1530067
<b>73</b>	<b>1120</b>	<b>NSAPDGLLF</b>	<b>0.51/0.71103764</b>
74	909	QYYNGIMVL	0.39/0.82550391
75	1261	SLFQTTVEL	0.00/1.2047668
76	796	GNPRCKNLL	0.90/0.30205663
77	1087	SRRLAQQKI	0.72/0.4655759
78	87	YIYVTNEI	0.41/0.77168294
79	1002	EAIHTVTIA	0.86/0.31598044
80	902	IADLACAQY	0.96/0.21199025
81	686	VYQQSIIGA	0.79/0.38133482
82	1244	TYLNLSEL	0.43/0.73997232

83	1287	NRFENYIKW	/1.1620609
84	667	VSTGNIFIV	0.49/0.65592697
85	1168	NGVFRVTSR	0.78/0.3656689
86	946	AAIPFSLAL	0.45/0.69063352
87	437	YCDSPFEKL	0.82/0.31664561
88	697	AVNESRYGL	0.44/0.67509782
89	624	LNNCTKYNI	0.72/0.39111634
90	701	SRYGLQNLL	0.00/1.1082027
91	1042	SIQAIYDRL	1.00/0.1074006
92	1144	WSGICVDGI	0.52/0.58640681
93	413	WTVAFATFV	0.63/0.47556945
94	689	QSIIGAMTA	0.68/0.42525788
95	320	LQYDVLFYC	0.71/0.39428404
96	361	HVSTFVGVL	0.46/0.63670723
97	368	VLPPTVREI	0.34/0.75536966
98	96	GLNASVTLK	0.00/1.0926166
99	186	VNATVTVNV	0.18/0.91153623
100	228	NGFPFNNWF	0.72/0.36993696
101	93	NEIGLNASV	0.98/0.10922593
102	517	VRTSHFSIR	0.84/0.24562668
103	383	QFYINGFKY	0.79/0.28272963
104	1303	ISVVFVLL	/1.0702129
105	392	FDLGFIEAV	0.72/0.3446838
106	1073	FVSQVLNKY	0.57/0.48234715
107	409	ATDFWTVAF	0.29/0.76233728
108	1082	TEVRSSRRL	0.53/0.52013817
109	80	GYVDVNQYY	0.02/1.0291098
110	299	NLNFSANSV	0.06/0.98802186
111	1003	AIHTVTIAL	0.22/0.82698101
112	834	TFDSNAFSL	0.33/0.71582162
113	806	QYTSACKTI	0.99/0.055726595
114	67	DVGNHRSAF	0.98/0.060473689
115	611	YPVSGIREF	0.02/1.0179848
116	665	KNVSTGNIF	0.78/0.24818534
117	1265	TTVELQGLI	0.69/0.32803909
118	518	RTSHFSIRY	0.00/1.0179404
119	58	YSANGFFYI	0.01/1.0064755
120	1080	KYTEVRSSR	0.02/0.99452973
121	123	SFDCIVNLL	0.18/0.83170893

122	1216	DVNKTLQEF	0.94/0.069375114
123	309	NLKSGVIVF	0.03/0.97633635
124	7	LLVLPLASC	0.84/0.16380939
125	420	FVDVLNVVS	0.90/0.099772074
126	892	DYKSCTKGL	0.53/0.46727909
127	156	VTRTFYVPA	0.89/0.10463975
128	793	VCNGNPRCK	0.70/0.2923813
129	168	LTKLSVKCY	0.98/0.011790597
130	50	CANQSTSVY	0.83/0.1571116
131	883	TSGLGTVDV	0.65/0.33360293
132	482	HTDINFTAT	0.98/0.00059218188
133	523	SIRYIYNRV	0.35/0.62967485
134	552	FSFSKLNNF	0.45/0.52867297
135	36	SSTIVTGLL	0.93/0.040560127
136	1112	GTHIFSIVN	0.86/0.11035307
137	749	NSSDNGISA	0.25/0.71807843
138	1035	NFQAISNSI	0.68/0.28221563
139	944	SAAAIPFSL	0.87/0.091521307
140	81	YYDVNQYYI	0.01/0.95071361
141	716	YVSNGGNNC	0.54/0.41685514
142	499	VCKPHQVNI	0.67/0.28679491
143	1108	FCGNGTHIF	0.97/-0.014700882
144	229	GFPFNNWFL	0.03/0.92362977
145	959	NYVALQTDV	0.66/0.29095699
146	812	KTIEDALRL	0.44/0.50292724
147	410	TDFWTVAFa	0.74/0.20196062
148	141	GITISGETV	0.93/0.011474211
149	1172	RVTSRVMFQ	0.25/0.68868661
150	160	FYVPAAYKL	0.01/0.92154374
151	1100	KSQSNRYGF	0.11/0.81717377
152	770	TTSVQVEYL	0.49/0.43589624
153	243	TLVDGVSRL	0.13/0.79565075
154	1045	AIYDRLDSI	0.00/0.92458091
155	1194	NCNVTFVNI	0.10/0.82143975
156	64	FYIDVGNHR	0.00/0.91957366
157	1028	LTSQLRHNF	0.99/-0.070453131
158	475	ALPIYYQHT	0.88/0.036254557
159	236	FLLTNGSTL	0.54/0.37110512



160	700	ESRYGLQNL	0.00/0.90894505
161	224	GRIPNGFPF	0.46/0.44783211
162	230	FPFNNWFL	0.24/0.66438825
163	609	VPYPVSGIR	0.14/0.75986932
164	960	YVALQTDVL	0.58/0.3188048
165	1017	VVNQQGSAL	0.81/0.088394453
166	595	ALYVTWSEG	1.00/-0.11064179
167	120	SSSSFDCIV	0.39/0.49728833
168	109	GINTTDFL	0.00/0.88301886
169	76	ALHTGYYDV	0.11/0.7710844
170	68	VGNHRSAFA	0.85/0.030503775
171	791	TYVCNGNPR	0.41/0.47005482
172	1199	FVNISRVEL	0.01/0.86752135
173	703	YGLQNLLQL	0.73/0.14329516
174	267	KSSTGFVYF	0.22/0.65221262
175	715	YYVSNGGNN	0.97/-0.099784989
176	57	VYSANGFFY	0.04/0.82896505
177	456	DGFYSANFL	0.94/-0.07281743
178	172	SVKCYFNYS	0.92/-0.058566303
179	1031	QLRHNFQAI	0.12/0.73975815
180	395	GFIEAVNFN	0.93/-0.077976637
181	310	LKSGVIVFK	0.95/-0.10067565
182	1273	IDQINSTYV	0.52/0.32710996
183	177	FNYS CVFSV	0.08/0.76687706
184	105	ICKFGINTT	0.96/-0.11791621
185	287	YQHNSVADV	0.22/0.62057052
186	1064	TGRLAALNA	0.88/-0.040067279
187	982	AINNIVASF	0.33/0.50952376
188	389	FKYFDLGFI	0.89/-0.050648976
189	185	VVNATVTVN	0.98/-0.14390927
190	1197	VTFVNISRV	0.17/0.66507703
191	165	AYKLT KLSV	0.25/0.57898383
192	148	TVRLHLYNV	0.05/0.77594486
193	136	LGAPLGITI	0.80/0.024552374
194	1291	NYIKWPWWV	/0.81982151
195	316	VFKTLQYDV	0.95/-0.133608
196	97	LNASVTLKI	0.70/0.11555113
197	865	SRIAGRSAL	0.05/0.7630997
198	996	AITQTAEAI	0.91/-0.099155362

199	416	AFATFVDVL	0.11/0.70016861
200	740	GSLIPVRPR	0.00/0.80968904
201	369	LPPTVREIV	0.59/0.21811733
202	1298	WVWLHISVV	/0.80704834
203	373	VREIVVART	0.93/-0.1252681
204	827	NDVSSMLTF	0.98/-0.17792716
205	74	AFALHTGYY	0.68/0.12115824
206	832	MLTFDSNAF	0.74/0.060872202
207	412	FWTVAFATF	0.90/-0.099847397
208	967	VLQENQKIL	0.76/0.036650192
209	921	ADAERMAMY	0.24/0.55454404
210	1302	IISVVFVVL	/0.79428752
211	497	CYVCKPHQV	0.04/0.75423386
212	294	DVMRYNLNF	0.40/0.38656581
213	901	SIADLACAQ	0.95/-0.16644765
214	357	INTTHVSTF	0.89/-0.1069322
215	1147	ICVDGIYGY	0.96/-0.17819768
216	352	FINSTINTT	0.74/0.041253138
217	247	GVSRLYQPL	0.57/0.20914439
218	1131	TVLLPTDYK	0.16/0.6128775
219	1170	VFRVTSRVM	0.81/-0.03960289
220	209	DCNGYTDNI	0.26/0.50820824
221	1285	LLNRFENYI	/0.76809085
222	391	YFDLGFIEA	0.19/0.57516701
223	158	RTFYVPAAY	0.53/0.23405711
224	232	FNNWFLLTN	0.83/-0.071336717
225	465	DDNVLPETY	0.95/-0.19146001
226	1342	STKLPYYEF	/0.75702633
227	184	SVVNATVTV	0.35/0.40585993
228	558	NNFQKFKTI	0.04/0.71521064
229	940	GGLTSAAAI	0.82/-0.065385197
230	226	IPNGFPFNN	0.00/0.75326527
231	653	TYVSNSGNL	0.13/0.62016707
232	22	NANLSMLQL	0.07/0.67580278
233	1225	AQNLPKYVK	0.02/0.72072651
234	349	YYCFINSTI	0.02/0.71576597
235	365	FVGVLPTTV	0.69/0.045548071
236	808	TSACKTIED	0.89/-0.16008049
237	384	FYINGFKYF	0.11/0.61735784

238	159	TFYVPAAYK	0.15/0.57711365
239	1196	NVTFVNISR	0.14/0.58609672
240	614	SGIREFSNL	0.01/0.7142468
241	274	YFNATGSDV	0.31/0.4094302
242	854	SSVLPQRNI	0.22/0.49273507
243	1086	SSRRLAQQK	0.05/0.66261708
244	934	IGGMVLGGL	0.92/-0.21023028
245	627	CTKYNIYDY	0.04/0.6686317
246	354	NSTINTTHV	0.89/-0.18209401
247	1118	IVNSAPDGL	0.63/0.067213964
248	332	SSGVLDTTI	0.95/-0.25758336
249	750	SSDNGISAI	0.00/0.68918248
250	999	QTAEAIHTV	0.04/0.64864443
251	950	FSLALQARL	0.00/0.68863499
<b>252</b>	<b>1292</b>	<b>YIKWPWWVW</b>	<b>/0.68663607</b>
253	1048	DRLDSIQAD	0.92/-0.23370948
254	297	RYNLNFSAN	0.13/0.55394579
255	279	GSDVNCNGY	0.70/-0.021040362
256	508	SLNGNTSVC	0.31/0.36716509
257	205	TVCDDCNGY	0.24/0.43709719
258	985	NIVASFSSV	0.00/0.67709251
259	1235	NFDLTPFNL	0.01/0.66392842
260	548	GTCPFSFSK	0.01/0.66124711
261	908	AQYYNGIMV	0.00/0.67011179
262	313	GVIVFKTLQ	0.90/-0.23226724
263	535	SPGDSSWHI	0.00/0.66516522
264	978	SFNKAINNI	0.17/0.49166828
265	466	DNVLPETYV	0.18/0.48133486
266	1190	VQIYNCNVT	0.93/-0.26963089
267	442	FEKLQCEHL	0.11/0.54846185
268	556	KLNNFQKFK	0.00/0.65676664
269	608	GVPYPVSGI	0.05/0.60612364
270	629	KYNIYDYVG	0.95/-0.29640713
271	730	TYSNFGICA	0.12/0.53332877
272	422	DVLVNVSAT	0.65/0.0032103136
273	937	MVLGGLTSA	0.95/-0.29802898
274	526	YIYNRVKSG	0.97/-0.3220123
275	153	LYNVTRTFY	0.53/0.11368219
276	975	LAASFNKAI	1.00/-0.3567849

277	1033	RHNFQAISN	0.99/-0.34988227
278	1094	KINECVKSQ	0.73/-0.090891295
279	258	TCLWPVPGL	0.00/0.63689405
280	516	CVRTSHFSI	0.98/-0.34396749
281	929	YTGSLIGGM	0.99/-0.35574078
282	326	FYCSNSSSG	0.95/-0.32770028
283	726	TAVMTYSNF	0.00/0.62184941
284	1084	VRSSRRLAQ	0.99/-0.36831842
285	544	YLKSGTCPF	0.11/0.50779975
286	407	ASATDFWTV	0.00/0.61754287
287	1072	AFVSQVLNK	0.01/0.60477023
288	295	VMRYNLNFS	0.96/-0.3457583
289	905	LACAQYYNG	0.66/-0.046858515
290	915	MVLPGVADA	0.04/0.57104125
291	836	DSNAFSLAN	0.98/-0.37150207
292	898	KGLSIADLA	0.57/0.038495119
293	562	KFKTICFST	0.88/-0.27337283
294	741	SLIPVRPRN	0.10/0.50452706
295	9	VLPLASCFF	0.92/-0.31727456
296	175	CYFNYSCVF	0.95/-0.34774304
297	1077	VLNKYTEVR	0.66/-0.059900419
298	755	ISAITANL	0.02/0.5761487
299	1202	ISRVELHTV	0.53/0.062218
300	549	TCPFSFSKL	0.18/0.41131222
301	641	IRSSNQSLA	0.00/0.58966843
302	920	VADAERMAM	0.59/-0.0010617828
303	810	ACKTIEDAL	0.15/0.42832587
304	852	NLSSVLPQR	0.05/0.52769584
305	374	REIVVARTG	0.64/-0.062689593
306	555	SKLNNFQKF	0.64/-0.065544177
307	521	HFSIRYIYN	0.03/0.54244334
308	1195	CNVTFVNIS	0.84/-0.26922709
309	1189	FVQIYNCNV	0.00/0.56962022
310	1219	KTLQEFAQN	0.78/-0.21340889
311	503	HQVNISLNG	0.96/-0.39355616
312	455	QDGFYSANF	0.94/-0.37839705
313	406	TASATDFWT	0.88/-0.32314226
314	334	GVLDTTIPF	0.02/0.53645384

315	619	FSNLVLNNC	0.01/0.54575229
316	82	YDVNQYYIY	0.81/-0.25705896
317	571	VAVPGSCNF	0.11/0.44260339
318	1163	VLYS DNGVF	0.91/-0.35876137
319	479	YYQHTDINF	0.13/0.41587506
320	1253	KQLEAKTAS	0.95/-0.40503389
321	180	SCVFSVVNA	0.11/0.43204577
322	78	HTGYYDVNQ	0.94/-0.40009667
323	125	DCIVNLLFT	0.98/-0.44244182
324	1301	LIISVVFVV	/0.53657419
325	885	GLGTVDVDY	0.95/-0.4163575
326	1275	QINSTYVDL	0.01/0.51961701
327	649	AGGITYVSN	0.98/-0.45085953
328	1037	QAISNSIQA	0.01/0.51902963
329	838	NAFSLANVT	0.81/-0.283673
330	225	RIPNGFPFN	0.99/-0.46715179
331	621	NLVLNNCTK	0.00/0.52102561
332	536	PGDSSWHIY	0.82/-0.2989773
333	672	IFIVTPCNQ	0.96/-0.44288359
334	919	GVADAERMA	0.04/0.4766117
335	424	LVNVSATKI	0.76/-0.24394201
336	154	YNVTRTFYV	0.01/0.50194357
337	1101	SQSNRYGFC	0.85/-0.3474219
338	600	WSEGNSITG	0.75/-0.2532342
339	1134	LPTDYKNVK	0.69/-0.19471857
340	504	QVNISLNGN	0.89/-0.39688785
341	721	GNNCTTAVM	0.01/0.48010274
342	869	GRSALEDLL	0.06/0.4283327
343	157	TRTFYVPAA	0.01/0.47811724
344	1184	PVLSDFVQI	0.54/-0.058458381
345	5	LILLVLPLA	0.97/-0.48936344
346	1062	LITGRLAAL	0.00/0.47893931
347	490	TASFGGSCY	0.05/0.42709924
348	1295	WPWWVWLII	/0.4757909
349	1305	VVFVVL LSL	/0.47473929
350	1226	QNLPKYVKP	0.92/-0.45107071
351	1164	LYSDNGVFR	0.01/0.45767625
352	181	CVFSVVNAT	0.01/0.45508949
353	506	NISLNGNTS	0.87/-0.40562111

354	302	FSANSVDNL	0.04/0.42407014
355	495	GSCYVCKPH	0.97/-0.50770133
356	1055	ADQQVDRLI	0.68/-0.21796388
357	1132	VLLPTDYKN	0.90/-0.43882807
358	767	SNWTTSVQV	0.00/0.46070684
359	973	KILAASFNK	0.61/-0.15005263
360	781	TSTPIVDC	0.04/0.41970129
361	483	TDINFTATA	0.52/-0.06251456
362	472	TYVALPIYY	0.00/0.45712689
363	821	SAHLETNDV	0.53/-0.074047723
364	524	IRYIYNRVK	0.00/0.45478019
365	725	TTAVMTYSN	0.64/-0.18585488
366	239	TNGSTLVDG	0.89/-0.43667911
367	837	SNAFSLANV	0.00/0.45020023
368	32	VPDNSSTIV	0.03/0.41572927
369	206	VCDDCNGYT	0.77/-0.32461314
370	1139	KNVKAWSGI	0.68/-0.23558417
371	385	YINGFKYFD	0.81/-0.36698015
372	66	IDVGNHRSA	0.94/-0.49907895
373	291	SVADVMRYN	0.00/0.44077119
374	1251	ELKQLEAKT	0.97/-0.53238036
375	1152	IYGYVLRQP	0.81/-0.37764397
376	398	EAVNFNVTT	0.84/-0.40778528
377	1267	VELQGLIDQ	0.91/-0.47956997
378	538	DSSWHIYLK	0.02/0.40856322
379	1114	HIFSIVNSA	0.00/0.42137042
380	1148	CVDGIYGYV	0.00/0.42057261
381	92	TNEIGLNAS	0.59/-0.17816339
382	866	RIAGRSALE	0.77/-0.35818712
383	1209	TVIPDYVDV	0.03/0.38178166
384	843	ANVTSFGDY	0.72/-0.30949526
385	912	NGIMVLPGV	0.52/-0.11400895
386	786	VVDCATYVC	0.01/0.39261562
387	380	RTGQFYING	0.00/0.39763001
388	344	PSSQPYYCF	0.59/-0.19545169
389	845	VTSFGDYNL	0.00/0.39244636
390	116	FLSNSSSSF	0.79/-0.39952442
391	292	VADVMRYNL	0.02/0.36732007
392	1041	NSIQAIYDR	0.00/0.38661732

393	762	NLSIPSNWT	0.71/-0.32634225
394	635	YVGTGIIRS	0.61/-0.2314161
395	1344	KLPYYEFEK	/0.37793187
396	427	VSATKIQNL	0.01/0.36410775
397	803	LLKQYTSAC	0.89/-0.51625964
398	486	NFTATASFG	0.97/-0.59844676
399	580	PLEATWHYT	0.99/-0.6187496
400	1294	KWPWWVWLI	/0.36874339
401	695	MTAVNESRY	0.00/0.36783213
402	856	VLPQRNIHS	0.66/-0.29358821
403	1181	PRLPVLSDF	0.54/-0.17438213
404	250	RLYQPLRLT	0.88/-0.51563493
405	307	VDNLKSGVI	0.62/-0.25756919
406	1063	ITGRLAALN	0.89/-0.53545237
407	170	KLSVKCYFN	0.64/-0.28554086
408	494	GGSCYVCKP	0.90/-0.54914357
409	1201	NISRVELHT	0.65/-0.30017658
410	670	GNIFIVTPC	0.54/-0.19053574
411	207	CDDCNGYTD	0.83/-0.48688557
412	270	TGFVYFNAT	0.62/-0.28113561
413	1115	IFSIVNSAP	0.99/-0.65832253
414	1223	EFAQNLPKY	0.91/-0.58286731
415	359	TTHVSTFVG	0.67/-0.34596887
416	1158	RQPNLVLYS	0.91/-0.58726713
417	776	EYLQITSTP	0.97/-0.65306558
418	1254	QLEAKTASL	0.85/-0.53660417
419	509	LNGNTSVCV	0.80/-0.48935029
420	322	YDVLFYCSN	0.89/-0.58518661
421	28	LQLGVPDNS	0.79/-0.49172582
422	1203	SRVELHTVI	0.57/-0.27584068
423	19	CNSNANLSM	0.72/-0.43013285
424	747	PRNSSDNGI	0.79/-0.50811028
425	330	NSSSGVLDT	0.83/-0.56128745
426	760	TANLSIPSN	0.96/-0.69625857
427	742	LIPVRPRNS	0.90/-0.64085629
428	233	NNWFLLTNG	0.97/-0.71680874
429	597	YVTWSEGNS	0.95/-0.69910405
430	610	PYPVSGIRE	0.96/-0.70919913
431	835	FDSNAFSLA	0.64/-0.39265703

432	1103	SNRYGFCGN	0.67/-0.42632403
433	718	SNGGNNCTT	0.61/-0.38121932
434	293	ADVMRYNLN	0.75/-0.5249897
435	314	VIVFKTLQY	0.52/-0.29505227
436	874	EDLLFSKVV	0.99/-0.77758162
437	14	SCFFTCNSN	0.65/-0.44179279
438	1095	INECVKSQS	0.85/-0.64417014
439	1143	AWSGICVDG	0.94/-0.73505255
440	327	YCSNSSSGV	0.95/-0.74568568
441	463	FLDDNVLPE	0.81/-0.61323046
442	61	NGFFYIDVG	0.95/-0.75551754
443	337	DTTIPFGPS	0.97/-0.77914943
444	820	LSAHLETND	0.91/-0.73122578
445	48	WICANQSTS	0.86/-0.69190279
446	688	QQSIIGAMT	0.87/-0.71448635
447	850	DYNLSSVLP	0.63/-0.50259028
448	364	TFVGVLPPT	0.78/-0.65289224
449	200	RVVNYTVCD	0.86/-0.7376643
450	355	STINTTHVS	0.87/-0.74978859
451	542	HIYLKSGTC	0.69/-0.57703562
452	217	IFSVQQDGR	0.52/-0.40725701
453	301	NFSANSVDN	0.51/-0.39917181
454	1109	CGNGTHIFS	0.69/-0.58205052
455	1236	FDLTPFNLT	0.97/-0.86773462
456	1096	NECVKSQSN	0.99/-0.8964071
457	10	LPLASCFFT	0.80/-0.70812199
458	543	IYLKSGTCP	0.79/-0.70586974
459	418	ATFVDVLVN	0.63/-0.54928233
460	1043	IQAIYDRLD	0.76/-0.68606589
461	440	SPFEKLQCE	0.87/-0.80543131
462	179	YSCVFSVVN	0.64/-0.57767518
463	127	IVNLLFTEQ	0.61/-0.55888341
464	1018	VNQQGSALN	0.98/-0.92953148
465	39	IVTGLLPTH	0.86/-0.81432357
466	722	NNCTTAVMT	0.74/-0.70055333
467	692	IGAMTAVNE	0.60/-0.56386442
468	963	LQTDVLQEN	0.98/-0.94548141
469	604	NSITGVPYP	0.52/-0.48684409
470	452	FGLQDGFYS	0.86/-0.82733097



471	459	YSANFLDDN	0.78/-0.7535541
472	315	IVFKTLQYD	0.78/-0.75511057
473	939	LGGLTSAAA	0.82/-0.79697613
474	245	VDGVSRLYQ	0.91/-0.89247564
475	45	PTHWICANQ	0.69/-0.6774849
476	790	ATYVCNGNP	0.81/-0.80119268
477	739	DGSLIPVRP	0.94/-0.93604532
478	763	LSIPSNWTT	0.90/-0.89723455
479	350	YCFINSTIN	0.60/-0.59842528
480	319	TLQYDVLFY	0.61/-0.60901685
481	652	ITYVSNSGN	0.97/-0.96911796
482	450	LQFGLQDGF	0.92/-0.91916096
483	568	FSTVAVPGS	0.70/-0.70282629
484	1015	QDVVNQQGS	0.98/-0.98322824
485	462	NFLDDNVLP	0.80/-0.80737677
486	183	FSVVNATVT	0.66/-0.6792846
487	421	VDVLNVSA	0.65/-0.67314091
488	732	SNFGICADG	0.90/-0.93035591
489	644	SNQSLAGGI	0.70/-0.73116338
490	990	FSSVNDAIT	0.51/-0.54952064
491	12	LASCFFTCN	0.56/-0.60483011
492	626	NCTKYNIYD	0.82/-0.8731964
493	1241	FNLTYLNLS	0.97/-1.0258481
494	744	PVRPRNSSD	0.84/-0.9073214
495	403	NVTTASATD	0.72/-0.79478634
496	794	CNGNPRCKN	0.75/-0.83111047
497	766	PSNWTTSVQ	0.76/-0.85006615
498	235	WFLLTNGST	0.79/-0.88042498
499	37	STIVTGLLP	0.88/-0.97046583
500	895	SCTKGLSIA	0.85/-0.94052322
501	660	NLLGFKNVS	0.72/-0.81066823
502	918	PGVADAERM	0.76/-0.85941545
503	561	QKFKTICFS	0.87/-0.98208006
504	1257	AKTASLFQT	0.85/-0.97013702
505	775	VEYLQITST	0.59/-0.71757214
506	498	YVCKPHQVN	0.69/-0.82879719
507	1021	QGSALNHLT	0.90/-1.0565682
508	246	DGVSRLYQP	0.61/-0.77281509
509	62	GFFYIDVGN	0.94/-1.124061

510	822	AHLETNDVS	0.92/-1.1094466
511	557	LNNFQKFKT	0.67/-0.86273161
512	1269	LQGLIDQIN	0.87/-1.073234
513	1026	NHLTSQLRH	0.58/-0.78799615
514	103	LKICKFGIN	0.55/-0.76576516
515	620	SNLVLNNCT	0.78/-1.0111585
516	818	LRLSAHLET	0.51/-0.74520102
517	46	THWICANQS	0.58/-0.82444777
518	734	FGICADGSL	0.55/-0.80259024
519	21	SNANLSMLQ	0.62/-0.87332856
520	651	GITYVSNSG	0.71/-0.96740514
521	1117	SIVNSAPDG	0.80/-1.0609345
522	375	EIVVARTGQ	0.95/-1.2191328
523	257	LTCLWPVPG	0.69/-0.96791555
524	704	GLQNLLQLP	0.71/-0.99244215
525	26	SMLQLGVPD	0.72/-1.0130671
526	593	VGALYVTWS	0.85/-1.1463538
527	161	YVPAAYKLT	0.95/-1.2536919
528	851	YNLSSVLPQ	0.63/-0.94170498
529	114	FDFLSNSSS	0.81/-1.1220574
530	691	IIGAMTAVN	0.74/-1.0588107
531	529	NRVKSGSPG	0.57/-0.89537498
532	496	SCYVCKPHQ	0.72/-1.0461098
533	540	SWHIYLKSG	0.69/-1.0174474
534	961	VALQTDVLQ	0.59/-0.9450207
535	675	VTPCNQPDQ	0.51/-0.86667655
536	219	SVQQDGRIP	0.63/-1.0192834
537	882	VTSGLGTVD	0.54/-0.93667796
538	578	NFPLEATWH	0.51/-0.9163556
539	487	FTATASFGG	0.57/-0.99628377
540	1056	DQQVDRLIT	0.95/-1.3798289
541	457	GFYSANFLD	0.52/-0.95852554
542	1123	PDGLLFLHT	0.87/-1.3505087
543	3	LFLILLVLP	0.95/-1.4529949
544	958	LNVALQTD	0.73/-1.2332524
545	596	LYVTWSEGN	0.92/-1.4279859
546	329	SNSSSGVLD	0.74/-1.2559305
547	280	SDVNCNGYQ	0.96/-1.5016393
548	1252	LKQLEAKTA	0.81/-1.3727952

	549	1183	LPVLSDFVQ	0.57/-1.1331272
	550	893	YKSCTKGLS	0.89/-1.4584225
	551	241	GSTLVDGVS	0.63/-1.2090417
	552	341	PFGPSSQPY	0.80/-1.3816066
	553	551	PFSFSKLNN	0.56/-1.167181
	554	325	LFYCSNSSS	0.52/-1.1365713
	555	25	LSMLQLGVP	0.64/-1.2976153
	556	139	PLGITISGE	0.87/-1.5836941
	557	1262	LFQTTVELQ	0.55/-1.3558081
	558	435	LLYCDSPFE	0.66/-1.4793635
	559	935	GGMVLGGLT	0.58/-1.4473077
	560	336	LDTTIPFGP	0.68/-1.572947
	561	853	LSSVLPQRN	0.79/-1.7629417
	562	1160	PNLVLYSDN	0.62/-1.7062135
	563	1129	LHTVLLPTD	0.96/-2.1223174
<b>HCoV-229E</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	1	235	SYADVLVNV	0.79/1.8674984
	2	1056	QYNQTILNL	0.82/1.3531035
	3	483	VTNGTIYSI	0.90/1.0448798
	4	1101	WLN RVET YI	0.98/0.95296113
	5	295	YHKHTFIVL	0.81/1.1011519
	6	320	RPAVVNITL	0.71/1.1982227
	7	892	HTLTKYTEV	0.81/1.0859292
	8	46	GYIPSNFAF	0.99/0.87869412
	9	769	AIQSRLNYV	0.90/0.95884354
	10	607	TYVCNGNVR	0.99/0.803791
	11	1010	NCNVTFVNI	0.95/0.82143975
	12	427	KPVEGVSSF	0.82/0.95057829
	13	1077	ELNYTVQKL	0.96/0.78826668
	14	1036	TLQELSYKL	0.93/0.77813212
	15	216	AVNFNVTNA	0.24/1.4507931
	16	277	TSPIQSVEL	0.87/0.81769459
	17	896	KYTEVRASR	0.84/0.83155092
	18	406	SHNIGSLYV	0.92/0.73114542
	19	201	FYINGYRYF	0.42/1.1868815
	20	558	IIAVQPRNV	1.00/0.60525918
	21	49	PSNFAFNW	0.78/0.80918015

22	368	FSFGKVNNF	0.86/0.7209464
23	516	FTSYGFSNV	0.99/0.57238523
24	52	FAFNNWFL	0.93/0.62976246
25	289	IVSLPVYHK	0.62/0.93506074
26	985	NYRITSRI	0.59/0.95204458
27	525	VEMPKFFYA	0.88/0.64569606
28	685	GRSAIEDIL	0.99/0.52356122
29	937	AAPEGLVFL	0.80/0.70872321
30	323	VVNITLANF	0.71/0.78885227
31	731	MVLPGVADA	0.92/0.57104125
32	89	RFTTGfVYF	0.48/1.0076679
33	916	KSQSKRYGF	0.72/0.76151462
34	162	VLGNFYCFV	0.29/1.1825707
35	1025	TIVPEYIDV	0.70/0.76114642
36	1029	EYIDVNKTL	0.61/0.84239534
37	232	ALASYADVL	0.85/0.59711401
38	417	SDGDVITGV	0.98/0.4623578
39	206	YRYFSLGNV	0.82/0.61256483
40	249	ANIIYCNSV	0.99/0.43179498
41	583	SNWTTSVQV	0.93/0.46070684
42	451	VSGVGVIRI	0.50/0.8906717
43	294	VYHKHTFIV	0.67/0.71297662
44	613	NVRCVELLK	0.92/0.45735117
45	111	NASSDVIRY	0.54/0.81078762
46	1118	CISVVLIFV	/1.340019
47	567	SYDSVSAIV	0.03/1.2935833
48	785	QENQKILAA	0.96/0.35162883
49	882	RLAALNVFV	0.02/1.2836399
50	346	TTQFVG VKF	0.74/0.54546205
51	991	SRIMFEPRI	0.69/0.59218388
52	394	AMPIMANLV	0.77/0.50575436
53	760	SAASIPFSL	0.99/0.28566631
54	115	DVIRYNINF	0.67/0.60410097
55	661	VSSFGDYNL	0.62/0.6461
56	135	KTSYGAVVF	1.00/0.25763561
57	694	FSKLVTSGL	0.91/0.34420326
58	369	SFGKVNNFV	0.98/0.26941735
59	178	TTSafVGAL	0.56/0.6823268
60	307	FELRRGPGR	0.88/0.3609089

61	847	QLRQNFQAI	0.67/0.56653056
62	689	IEDILFSKL	0.59/0.64639245
63	393	CAMPIMANL	0.39/0.84452801
64	678	RSGSRVAGR	0.28/0.9520858
65	302	VLVYNFELR	0.85/0.38186099
66	903	SRQLAQQKV	0.88/0.34964962
67	771	QSRLNYVAL	0.09/1.1311114
68	189	TVREFVISR	0.15/1.0555086
69	252	IYCNSVINR	0.48/0.71575058
70	1103	NRVETYIKW	0.30/0.89489893
71	374	NNFVKFGSV	0.86/0.33249972
72	127	LRRGTILFK	0.64/0.54941191
73	376	FVKFGSVCF	0.58/0.60784418
74	735	GVADAERMA	0.71/0.4766117
75	316	CYNCRPAVV	0.28/0.90276199
76	766	FSLAIQSRL	0.95/0.21696195
77	601	IVVDCSTYV	0.25/0.91605442
78	166	FYCFVNTTI	0.72/0.44509881
79	1087	TLIDNINST	0.80/0.36106377
80	3	VLLVAYALL	0.96/0.19537306
81	747	GSLIGGIAL	0.09/1.0482509
82	718	IADLACAQY	0.92/0.21199025
83	66	VVDGVVRSF	0.58/0.54851216
84	342	TSHFTTQFV	0.47/0.65637686
85	930	HIFSLVNAA	0.75/0.37491738
86	124	EENLRRGTI	0.06/1.0626125
87	401	LVNHKSHNI	0.61/0.50902711
88	914	CVKSQSKRY	0.86/0.25632977
89	885	ALNVFVSHT	0.94/0.17364435
90	397	IMANLVN HK	0.85/0.25302582
91	990	TSRIMFEPR	0.98/0.11959916
92	513	SENFTSYGF	0.75/0.34772899
93	443	CTKYNIYDV	0.51/0.58622773
94	725	QYYNGIMVL	0.27/0.82550391
95	537	TYNCTDAVL	0.00/1.0931999
96	73	SFQPLLLNC	0.98/0.10832622
97	301	IVLYVNFEL	0.84/0.24266543
98	972	VLRQPNLAL	0.64/0.43922754
99	297	KHTFIVLYV	0.71/0.36403013

100	112	ASSDVIRYN	0.70/0.3726522
101	877	RLITGRLAA	0.90/0.1717323
102	1041	SYKLPNYTV	0.61/0.44278541
103	1094	STLVDLKWL	0.85/0.19147491
104	1	MFVLLVAYA	0.53/0.50741409
105	209	FSLGNVEAV	0.01/1.0208665
106	1084	KLQTLIDNI	0.73/0.2981202
107	1121	VVLIFVVSM	/1.0266072
108	450	DVSGVGVIR	0.14/0.88629946
109	318	NCRPAVVNI	0.15/0.86753865
110	203	INGRYFSL	0.83/0.1868726
111	970	GYVLRQPNL	0.07/0.94512873
112	492	TPCNPPDQL	0.83/0.18329478
113	640	LESADVSEM	0.92/0.091135185
114	94	FVYFNGTGR	0.58/0.43057183
115	70	VVRSFQPLL	0.94/0.065829315
116	1096	LVDLKWLNR	0.02/0.98490518
117	282	SVELPVSIV	0.02/0.98416244
118	299	TFIVLYVNF	0.18/0.82148897
119	462	DTFLNGITY	0.05/0.95137086
120	553	CADGSIIAV	0.49/0.50823926
121	1008	IENCNVTFV	0.14/0.84796279
122	1032	DVNKTLQEL	0.22/0.76726182
123	345	FTTQFVGVK	0.49/0.49421911
124	675	SLPRSGSRV	0.00/0.97343014
125	840	SLNHLTSQL	0.02/0.95158788
126	870	QADQQVDRL	0.81/0.16036435
127	947	TVLLPTQYK	0.10/0.86981699
128	996	EPRIPTIAD	0.96/0.0095154556
129	864	DRLDIIQAD	0.99/-0.028714089
130	1005	FVQIENCNV	0.56/0.38784067
131	1102	LNRVETYIK	0.98/-0.039698991
132	994	MFEPRIPTI	0.10/0.82822264
133	782	DVLQENQKI	0.94/-0.018259514
134	861	AIYDRLDII	0.00/0.91534075
135	5	LVAYALLHI	0.74/0.17394567
136	643	ADVSEMLTF	0.99/-0.077213855
137	119	YNINFEENL	0.05/0.85968561
138	364	GNCPFSGFK	0.42/0.48731197

139	703	GTVDADYKK	0.99/-0.10022226
140	185	ALPKTVREF	0.17/0.71973496
141	874	QVDRLITGR	0.66/0.22945282
142	646	SEMLTFDKK	0.94/-0.053857407
143	36	SENVFAVES	0.98/-0.10024764
144	472	STSGNLLGF	0.80/0.078257942
145	802	IVDAFTGVN	0.42/0.45761822
146	104	DCKGFYSNA	0.15/0.72682753
147	681	SRVAGRSAI	0.26/0.61071974
148	405	KSHNIGSLY	0.27/0.60015863
149	722	ACAQYYNGI	0.74/0.12818655
150	1001	TIADFVQIE	0.29/0.57621536
151	75	QPLLLNCLW	0.98/-0.11717826
152	26	HSVCNGCVG	1.00/-0.14273432
153	608	YVCNGNVRC	0.65/0.20553062
154	200	HFYINGYRY	0.10/0.75470995
155	974	RQPNLALYK	0.41/0.44467609
156	1080	YTVQKLQTL	0.74/0.11050933
157	532	YASNGTYNC	0.59/0.25669996
158	223	NAATTVCTV	0.43/0.41024426
159	876	DRLITGRLA	0.79/0.045018785
160	812	AITQTSQAL	0.42/0.41468226
161	281	QSVELPVS	0.96/-0.12663657
162	348	QFVGVKFDR	0.93/-0.1090081
163	453	GVGVIRISN	0.83/-0.025506206
164	697	LVTSGLGTV	0.97/-0.16600661
165	311	RGPGRCYNC	0.02/0.7799757
166	611	NGNVRCVEL	0.01/0.78743139
167	858	SIQAIYDRL	0.69/0.1074006
168	434	SFMNVTLNK	0.88/-0.083120717
169	854	AISSSIQAI	0.56/0.23386998
170	1035	KTLQELSYK	0.06/0.72787522
171	34	GHSENVFAV	0.14/0.64571548
172	762	ASIPFSLAI	0.26/0.52224994
173	801	NIVDAFTGV	0.09/0.68411889
174	1114	WWWLCISVV	/0.77228223
175	319	CRPAVVNIT	0.61/0.16148364
176	1107	TYIKWPWWV	0.00/0.77137517
177	789	KILAASFNK	0.92/-0.15005263

178	378	KFGSVCFSL	0.06/0.70669931
179	980	LYKEGNYYR	0.24/0.52449807
180	566	VSYDSVSAI	0.10/0.66293112
181	149	TLVSGDAHI	0.14/0.60636099
182	573	AIVTANLSI	0.88/-0.13653229
183	940	EGLVFLHTV	0.89/-0.14936414
184	2	FVLLVAYAL	0.00/0.73925828
185	169	FVNTTIGNE	0.95/-0.21603387
186	446	YNIYDVSGV	0.90/-0.16686651
187	724	AQYYNGIMV	0.06/0.67011179
188	234	ASYADVLVN	0.96/-0.23019959
189	656	FTLANVSSF	0.66/0.064119553
190	986	YYRITSRIM	0.01/0.70525022
191	1100	KWLNRVETY	0.12/0.59135612
192	910	KVNECVKSQ	0.54/0.16973449
193	353	KFDRWSASI	0.08/0.62899224
194	448	IYDVSGVGV	0.24/0.46726532
195	196	SRTGHFYIN	0.99/-0.28548587
196	143	FYCTNNTLV	0.02/0.67709199
197	907	AQQKVNECV	0.92/-0.22454834
198	137	SYGAVVFYC	0.00/0.69264725
199	981	YKEGNYYRI	0.00/0.69189128
200	341	DTSHFTTQF	0.29/0.39790601
<b>201</b>	<b>1108</b>	<b>YIKWPWWWW</b>	<b>/0.68663607</b>
202	668	NLSSVIPSL	0.04/0.6464634
203	1086	QTLIDNINS	0.99/-0.30454623
204	688	AIEDILFSK	0.29/0.38699398
205	765	PFS LAIQSR	0.89/-0.21756993
206	929	THIFSLVNA	0.69/-0.018795332
207	524	VVEMPKFFY	0.98/-0.30957933
208	1019	SRSELQTIV	0.16/0.50968915
209	387	KDIPGGCAM	0.11/0.55922917
210	851	NFQAISSSI	0.67/-0.0030036502
211	218	NFNVTNAAT	0.76/-0.094476459
212	356	RWSASINTG	0.90/-0.23586429
213	154	DAHIPSGTV	0.20/0.46166747
214	198	TGHFYINGY	0.76/-0.099687321
215	241	VNVSQTAIA	0.84/-0.17998223



216	521	FSNVVEMPK	0.81/-0.15167492
217	253	YCNSVINRL	0.00/0.6486805
218	560	AVQPRNVSY	0.01/0.63705648
219	536	GTYNCTDAV	0.62/0.020826311
220	982	KEGNYRIT	0.99/-0.35082543
221	1092	INSTLVDLK	0.97/-0.33162479
<b>222</b>	<b>936</b>	<b>NAAPEGLVF</b>	<b>0.04/0.59776222</b>
223	571	VSAIVTANL	0.03/0.59471685
224	717	SIADLACAQ	0.79/-0.16644765
225	136	TSYGAVVFY	0.17/0.4473033
226	542	DAVLTYSSF	0.01/0.60335384
227	113	SSDVIRYNI	0.01/0.60002135
228	1012	NVTFVNISR	0.02/0.58609672
229	1089	IDNINSTLV	0.51/0.094785561
230	312	GPGRCYNCR	0.61/-0.0062245289
231	1070	TLENKSAEL	0.08/0.51648165
232	818	QALQTVATA	0.00/0.59629874
233	165	NFYCFVNTT	0.53/0.065864476
234	792	AASFNKAMT	0.65/-0.057975537
235	415	SWSDGDVIT	0.99/-0.39962906
236	609	VCNGNVRCV	0.14/0.45011757
237	902	ASRQLAQK	0.04/0.54710832
238	246	TAIANIYC	0.00/0.58374967
239	578	NLSIPSNWT	0.91/-0.32634225
240	732	VLPGVADAE	0.98/-0.39679437
241	1049	VPDLVVEQY	0.02/0.55750201
242	1015	FVNISRSEL	0.04/0.53516845
243	737	ADAERMAMY	0.02/0.55454404
244	886	LNVFVSHTL	0.96/-0.38847497
245	24	TSHSVCNGC	0.13/0.44104516
246	671	SVIPSLPRS	0.60/-0.029356677
247	74	FQPLLNCL	0.01/0.55990136
248	1045	PNYTVPDV	0.91/-0.3424814
249	1022	ELQTIVPEY	0.81/-0.24544621
250	829	KIQDVVNQQ	0.97/-0.40984551
251	464	FLNGITYTS	0.66/-0.10644623
252	835	NQQGNSLNH	0.98/-0.43019055

253	511	MLSENFTSY	0.55/-0.0020421399
254	197	RTGHFYING	0.00/0.54784924
255	546	TYSSFGVCA	0.00/0.54592906
256	519	YGFSNVVEM	0.07/0.47486415
257	1091	NINSTLVDL	0.00/0.5390921
258	1156	STKLPHYDV	/0.53783653
259	1116	WLCISVCLI	/0.53187977
260	439	TLNKCTKYN	0.90/-0.37017565
261	225	ATTVCTVAL	0.14/0.38865657
262	963	LCVDGINGY	0.86/-0.33411288
263	867	DIIQADQQV	0.57/-0.046201068
264	351	GVKFDRWSA	0.00/0.52357451
265	905	QLAQQKVNE	0.97/-0.44737215
266	641	ESADVSEML	0.00/0.51824056
267	825	TALNKIQDV	0.79/-0.2730957
268	523	NVVEPKFF	0.85/-0.33492762
269	445	KYNIYDVSG	0.86/-0.35250506
270	865	RLDIIQADQ	0.81/-0.30272767
271	917	SQSKRYGFC	0.88/-0.37717579
272	77	LLNCLWSV	0.14/0.36169184
273	973	LRQPNLALY	0.01/0.49040965
274	161	TVLGNFYCF	0.01/0.48814844
275	310	RRGPGRCYN	0.00/0.49270729
276	51	NFAFNNWFL	0.00/0.48776454
277	744	MYTGSLIGG	0.85/-0.36553556
278	69	GVVRSFQPL	0.06/0.42212265
279	878	LITGRLAAL	0.00/0.47893931
280	701	GLGTVDADY	0.95/-0.4719181
281	612	GNVRCVELL	0.00/0.47415948
282	952	TQYKDVEAW	0.67/-0.19763242
283	262	RCDQLSFDV	0.55/-0.080345705
284	898	TEVRASRQL	0.00/0.46636497
285	908	QQKVNECVK	0.58/-0.11443267
286	708	DYKKCTKGL	0.10/0.36496436
287	37	ENVFAVESG	0.90/-0.4378157
288	602	VVDCSTYVC	0.00/0.46013971
289	327	TLANFNETK	0.00/0.45979244
290	597	TSTPIVDC	0.04/0.41970129

291	88	SRFTTGfVY	0.52/-0.061584829
292	586	TTSVQVEYL	0.02/0.43589624
293	255	NSVINRLRC	0.03/0.42406909
294	372	KVNNFVKFG	0.00/0.45339313
295	626	ACKTIEDAL	0.02/0.42832587
296	1039	ELSYKLPNY	0.00/0.44555067
297	819	ALQTVATAL	0.00/0.44195641
298	308	ELRRGPGRC	0.58/-0.14132533
299	155	AHIPSGTVL	0.00/0.43257788
300	322	AVVNITLAN	0.80/-0.37110654
301	650	TFDKKAFTL	0.00/0.42624886
302	245	QTAIANIIY	0.01/0.41188351
303	846	SQLRQNFQA	0.66/-0.24359989
304	628	KTIEDALRN	0.77/-0.35564078
305	38	NVFAVESGG	0.73/-0.32178622
306	315	RCYNCRPAV	0.01/0.39397177
307	889	FVSHTLTky	0.63/-0.22663645
308	633	ALRNSAMLE	0.53/-0.12694683
309	358	SASINTGNC	0.01/0.39131931
310	396	PIMANLVNH	0.84/-0.4402206
311	606	STYVCNGNV	0.03/0.36592377
312	1142	FFSCFASSI	/0.39565224
313	887	NVFVSHTLT	0.96/-0.56489893
314	942	LVFLHTVLL	0.73/-0.33703446
315	857	SSIQAiyDR	0.02/0.37213588
316	517	TSYGFSNVV	0.00/0.38948488
317	508	VGAMLSENF	0.74/-0.35252647
318	62	NTSSVVDGV	0.01/0.37533965
319	632	DALRNSAML	0.01/0.36842498
320	452	SGVGVIRIS	0.85/-0.47314963
321	1113	WWVWLCISV	/0.3685549
322	1138	GCCGFFSCF	/0.36600621
323	1117	LCISVVLIF	/0.36038831
324	590	QVEYLQITS	0.99/-0.6333917
325	156	HIPSGTVLG	0.77/-0.41714308
326	1088	LIDNINSTL	0.97/-0.62490563
327	179	TSAFVGALP	0.76/-0.41598162
328	470	YTSTSGNLL	0.96/-0.62167019
329	912	NECVKSQSK	0.51/-0.17224333

330	367	PFSFGKVNN	0.96/-0.63819861
331	236	YADVLVNVS	0.62/-0.30480027
332	605	CSTYVCNGN	0.84/-0.52834667
333	716	LSIADLACA	0.96/-0.65097904
334	803	VDAFTGVND	0.55/-0.25543738
335	1081	TVQKLQTLI	0.53/-0.24006154
336	745	YTGSLIGGI	0.54/-0.25176861
337	1026	IVPEYIDVN	0.90/-0.6121174
338	592	EYLQITSTP	0.94/-0.65306558
339	790	ILAASFNKA	0.86/-0.57923679
340	738	DAERMAMYT	0.70/-0.42524277
341	843	HLTSQLRQN	0.90/-0.62566165
342	1011	CNVTFVNIS	0.54/-0.26922709
343	78	LLNCLWSVS	0.69/-0.42631526
344	402	VNHKSHNIG	0.88/-0.62056885
345	772	SRLNYVALQ	0.60/-0.34824453
346	273	GFYSTSPIQ	0.99/-0.73964479
347	993	IMFEPRIPT	0.55/-0.30814447
348	625	SACKTIEDA	0.97/-0.73039223
349	883	LAALNVFVS	0.98/-0.74296752
350	1034	NKTLQELSY	0.99/-0.75423339
351	1061	ILNLTSEIS	0.97/-0.73497931
352	1054	VEQYNQTIL	0.52/-0.28622372
353	687	SAIEDILFS	0.81/-0.57788787
354	557	SIIAVQPRN	0.81/-0.57822068
355	292	LPVYHKHTF	0.51/-0.30073016
356	1013	VTFVNISRS	0.57/-0.370745
357	192	EFVISRTGH	0.67/-0.47310592
358	568	YDSVSAIVT	0.75/-0.55609741
359	455	GVIRISNDT	0.72/-0.53619664
360	475	GNLLGFKDV	0.61/-0.42674079
361	97	FNGTGRGDC	0.74/-0.5576958
362	925	CGNGTHIFS	0.76/-0.58205052
363	272	DGFYSTSPI	0.75/-0.57516659
364	649	LTFDKKAFT	0.81/-0.6382898
365	509	GAMLSEFT	0.87/-0.69861697
366	237	ADVLVNVSQ	0.98/-0.83313
367	181	AFVGALPKT	0.81/-0.67288943
368	96	YFNGTGRGD	0.98/-0.84910663

369	19	TNGTNTSHS	0.95/-0.81915444
370	361	INTGNCPFS	0.57/-0.44375531
371	168	CFVNTTIGN	0.92/-0.8073517
372	204	NGYRYFSLG	0.61/-0.49905751
373	152	SGDAHIPSG	0.70/-0.59317155
374	850	QNFQAISSS	0.91/-0.8067722
375	1021	SELQTIVPE	0.65/-0.5495362
376	13	IAGCQTTNG	0.72/-0.62907467
377	1073	NKSAELNYT	0.75/-0.6682695
378	274	FYSTSPIQS	0.66/-0.58502157
379	43	ESGGYIPSN	0.86/-0.78829773
380	333	ETKGPLCVD	0.60/-0.53306782
381	12	HIAGCQTTN	0.88/-0.82711406
382	207	RYFSLGNVE	0.57/-0.51871077
383	400	NLVNHKSHN	0.91/-0.85980582
384	106	KGFYSNASS	0.91/-0.86585437
385	712	CTKGLSIAD	0.76/-0.72347179
386	331	FNETKGPLC	0.89/-0.85467211
387	411	SLYVSWSDG	0.65/-0.61546482
388	989	ITSRIMFEP	0.57/-0.54334517
389	879	ITGRLAALN	0.56/-0.53545237
390	268	FDVPDGFYS	0.77/-0.76135237
391	663	SFGDYNLSS	0.90/-0.92363067
392	831	QDVVNQQGN	0.96/-0.98436847
393	698	VTSGLGTVD	0.91/-0.93667796
394	939	PEGLVFLHT	0.81/-0.84117946
395	17	QTTNGTNTS	0.97/-1.0132252
396	564	RNVSYDSVS	1.00/-1.0474162
397	305	VNFELRRGP	0.97/-1.0341897
398	468	ITYTSTSGN	0.71/-0.77757767
399	968	INGYVLRQP	0.56/-0.6406024
400	335	KGPLCVDTS	0.72/-0.80694304
401	157	IPSGTVLGN	0.81/-0.89919577
402	80	NCLWSVSGS	0.66/-0.75737334
403	399	ANLVNHKSH	0.93/-1.0292441
404	418	DGDVITGVP	0.78/-0.88356208
405	1082	VQKLQTLID	0.78/-0.89669535
406	285	LPVSIVSLP	0.52/-0.65196154
407	644	DVSEMLTFD	0.70/-0.84566535

408	555	DGSIIAVQP	0.90/-1.0553293
409	734	PGVADAERM	0.70/-0.85941545
410	535	NGTYNCTDA	0.55/-0.72438715
411	813	ITQTSQALQ	0.79/-0.98447689
412	328	LANFNETKG	0.55/-0.74500319
413	693	LFSKLVTS	0.98/-1.1754057
414	593	YLQITSTPI	0.79/-0.99429385
415	875	VDRLITGRL	0.68/-0.88516301
416	76	PLLLNCLWS	0.82/-1.0499466
417	589	VQVEYLQIT	0.89/-1.1386482
418	654	KAFTLANVS	0.97/-1.2275132
419	172	TTIGNETTS	0.91/-1.1715195
420	962	GLCVDGING	0.73/-0.99284111
421	128	RRGTILFKT	0.70/-0.96898117
422	265	QLSFDVPDG	0.89/-1.1631618
423	528	PKFFYASNG	0.75/-1.0233617
424	505	QAVVGAMLS	0.56/-0.85011124
425	29	CNGCVGHSE	0.61/-0.90378002
426	515	NFTSYGFSN	0.77/-1.0820661
427	705	VDADYKKCT	0.93/-1.2732464
428	1031	IDVNKTLQE	0.84/-1.1900464
429	866	LDIIQADQQ	0.99/-1.3408012
430	56	NWFLLTNTS	0.80/-1.1595931
431	105	CKGFYSNAS	0.51/-0.87994452
432	496	PPDQLVVYQ	0.66/-1.033688
433	366	CPFSFGKVN	0.64/-1.0220149
434	1030	YIDVNKTLQ	0.54/-0.94559946
435	1085	LQTLIDNIN	0.53/-0.94715365
436	67	VDGVVRSFQ	0.63/-1.0473691
437	976	PNLALYKEG	0.99/-1.4108035
438	788	QKILAASFN	0.83/-1.3005132
439	757	GLTSAASIP	0.56/-1.0697704
440	9	ALLHIAGCQ	0.55/-1.0601386
441	436	MNVTLNKCT	0.88/-1.4132269
442	79	LNCLWSVSG	0.93/-1.4709827
443	827	LNKIQDVVN	0.82/-1.3779429
444	329	ANFNETKGP	0.90/-1.4704893
445	68	DGVVRSFQP	0.78/-1.3561402
446	868	IIQADQQVD	0.84/-1.4393487

	447	1067	EISTLENKS	0.78/-1.3808414
	448	1068	ISTLENKSA	0.76/-1.3634631
	449	22	TNTSHSVCN	0.59/-1.2011995
	450	30	NGCVGHSEN	0.65/-1.2884225
	451	467	GITYTSTSG	0.54/-1.3242179
	452	695	SKLVTSGLG	0.60/-1.4684949
	453	749	LIGGIALGG	0.52/-1.765249
<b>HCoV-OC43</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	1	1047	ALNNLLQQL	0.95/1.4112217
	2	698	FRNIKCNYV	0.95/1.1709293
	3	987	FYLVNQYRI	0.98/1.090522
	4	1273	NRLQEAIKV	0.94/1.1290627
	5	881	GVNFNVDDI	0.62/1.3894502
	6	1085	LINGRLTAL	0.91/1.0044934
	7	1089	RLTALNAYV	0.68/1.0179382
	8	1227	MLNTSIPNL	0.93/0.76203536
	9	456	FNPSTWNKR	0.97/0.72124296
	10	1303	WYVWLLICL	/1.6803307
	11	1	MFLILLISL	0.95/0.72279412
	12	620	IELGVCVNY	0.92/0.74972224
	13	732	AYNSTAISV	0.21/1.4529222
	14	209	TYDVNATYL	0.19/1.46477
	15	646	YYNSWQNLL	0.99/0.6301557
	16	1068	EILSRDAL	0.99/0.61409618
	17	438	TSCQLYYNL	0.23/1.3659761
	18	240	KFLFNVYLG	0.88/0.68603567
	19	841	SFCDNINAI	0.31/1.2510239
	20	445	NLPAANVSV	0.86/0.69940644
	21	455	RFPSTWNK	0.68/0.85397591
	22	1050	NLLQQLSNR	0.93/0.59233873
	23	724	SYLGCVVNA	0.99/0.53149928
	24	1274	RLQEAIKVL	0.84/0.67054773
	25	1123	KSQSSRINF	0.96/0.5384127
	26	1012	IANAFNNAL	0.99/0.49337339
	27	707	FNNSLTRQL	0.86/0.60523812
	28	49	TNGLGTYVYV	0.53/0.91235578
	29	1276	QEAIVLNH	0.99/0.45116678

30	61	VYLNTTLFL	0.83/0.59556931
31	78	STYRNMALK	0.18/1.2296772
32	73	YPTSGSTYR	0.71/0.68884631
33	1281	VLNHSYINL	0.35/1.0482089
34	59	DRVYLNTTL	0.91/0.48591595
35	425	YLQSSNYRI	0.96/0.41883581
36	1033	ALVKIQAVV	1.00/0.37157227
37	1065	SLQEILSRL	0.31/1.0593795
38	1234	NLPDFKEEL	0.04/1.3193041
39	645	TYYNSWQNL	0.19/1.1651561
40	44	DTVDTVNL	0.93/0.41931319
41	180	ICHPNLGNH	0.92/0.4269388
42	900	SECSKASSR	0.94/0.39299948
43	55	YYVLDRVYL	0.71/0.62291031
44	1193	MYTGSGYYY	0.97/0.36147644
45	848	AILTEVNEL	0.14/1.1749989
46	774	FEPFTVNSV	0.01/1.3022989
47	53	GTYYVLDRV	0.69/0.60199689
48	886	VDDINFSPV	0.83/0.45986232
49	883	NFNVDDINF	0.78/0.50752128
50	573	CQPQAFLGW	0.99/0.28651332
51	823	AFVCGDYAA	0.99/0.28228767
52	801	FTIGNMEEF	0.98/0.29156448
53	238	VTKFLFNVY	0.62/0.63532879
54	934	TGGAEIRDL	0.93/0.3175666
55	103	DFINGIFAK	0.96/0.28254372
56	372	NMSSLMSFI	0.68/0.54977987
57	221	FYQEGGTFY	0.81/0.41757152
58	232	FTDTGFVTK	0.82/0.40702178
59	314	IAPPTGVYE	0.95/0.27336411
60	86	KGTDLLSTL	0.90/0.31778329
61	249	MALSHYYVM	0.56/0.65690464
62	300	MSDFMSEIK	0.98/0.23444278
63	75	TSGSTYRNM	0.01/1.2010616
64	553	GEHCSGLAV	0.90/0.30660072
65	172	NMCEYPHTI	0.90/0.30410877
66	104	FINGIFAKV	0.55/0.65009989
67	695	ALLFRNIKC	1.00/0.190667
68	102	SDFINGIFA	1.00/0.19053218



69	1185	FVNVNNTWM	0.44/0.74167686
70	93	TLWFKPPFL	0.48/0.69472086
71	1121	CVKSQSSRI	0.64/0.5260796
72	1029	ATNSALVKI	0.98/0.17734067
73	353	KSVPSPLNW	0.90/0.25572466
74	705	YVFNNLSTR	0.92/0.23398838
75	1255	DLSLDYINV	0.30/0.84797395
76	257	MPLTCISRR	0.95/0.19667284
77	315	APPTGVYEL	0.02/1.1244147
78	288	NQDGIIFNA	0.53/0.61375342
79	7	ISLPTAFAV	0.04/1.1025963
80	443	YYNLPAANV	0.11/1.0288998
81	1149	LYFIHFNYV	0.03/1.1084429
82	1043	ANAEALNNL	0.05/1.0871314
83	312	QSIAPPTGV	0.83/0.30513685
84	415	KVDLQLGNL	0.01/1.1246018
85	286	AFNQDGIIF	0.37/0.76161999
86	759	RRSRRAITT	0.58/0.54861985
87	1184	YFVNVNNTW	1.00/0.11331374
88	237	FVTKFLFNV	0.50/0.61279072
89	983	AGVPFYLVN	0.24/0.86902301
90	89	DLLSTLWFK	0.49/0.60690967
91	838	EYGSFCDNI	0.03/1.0662257
92	693	EPALLFRNI	0.20/0.89291034
93	165	EVSVCQYNM	0.43/0.6575204
94	1205	ITENNVVVM	0.61/0.47194715
95	1297	YYVKWPWYV	/1.0811943
96	587	LQGDKCNIF	0.79/0.29042741
97	593	NIFANFILH	0.84/0.23956208
98	840	GSFCDNINA	0.77/0.30676157
99	1062	ISASLQEIL	0.54/0.53070597
100	299	CMSDFMSEI	0.00/1.0705229
101	1218	VNYTKAPYV	0.75/0.31957114
102	755	YSKNRRSRR	0.09/0.97754
103	481	FTNHSVVYA	0.03/1.0358614
104	663	GFRDYITNR	0.73/0.33455557
105	72	YYPTSGSTY	0.90/0.16318297
106	1267	DLQVEMNRL	0.01/1.0440607
107	486	VVYAQHCFK	0.81/0.23959963

108	1264	TFLDLQVEM	0.03/1.0192909
109	190	KELWHLDTG	0.99/0.054476873
110	408	FAIPNRRKV	0.31/0.73333435
111	100	FLSDFINGI	0.03/1.0039439
112	889	INFSPVLGC	0.97/0.05905104
113	214	ATYLYFHFY	0.50/0.52759043
114	869	MNGVTLSTK	0.92/0.1047512
115	387	CNNIDAANKI	0.97/0.051703862
116	1315	AMLVLLFFI	/1.0168347
117	1284	HSYINLKDI	0.75/0.26674026
118	1008	NQKLIANAF	0.25/0.76483172
119	410	IPNRRKVDL	0.00/1.0144714
120	981	AAAGVPFYL	0.06/0.95135741
121	175	EYPHTICHP	0.98/0.029419481
122	26	RLKGSFNRR	0.08/0.92219217
123	68	FLNGYYPTS	0.99/0.0075039235
124	279	TPRQYLLAF	0.65/0.34707301
<b>125</b>	<b>1298</b>	<b>YVKWPWYVW</b>	<b>/0.9882908</b>
126	1220	YTKAPYVML	0.10/0.88798981
127	152	STQDGVNKL	0.58/0.40302599
128	831	ACKLQLVEY	0.41/0.57250738
129	452	SVSRFNPST	0.88/0.097310705
130	916	FDKVKLSDV	0.99/-0.013587218
131	395	IYGMCFSSI	0.00/0.96874448
132	660	NLYGFRDYI	0.31/0.65430472
133	1114	AMEKVNECV	0.98/-0.027050296
134	664	FRDYITNRT	0.91/0.041126691
135	248	GMALSHYYV	0.41/0.53802265
136	223	QEGGTFYAY	0.51/0.43651074
137	336	RRKPDLPNC	0.58/0.35549882
138	598	FILHDVNNG	0.89/0.041984006
139	762	RRAITTGYR	0.01/0.91189376
140	1208	NNVVVMSTC	0.97/-0.062852773
141	971	TSASLFPPW	0.58/0.31571442
142	1278	AIKVLNHSY	0.88/0.009819656
143	984	GVPFYLVNQ	0.97/-0.081793283
144	1126	SSRINFCGN	0.80/0.087054246
145	50	NGLGTYYYVL	0.49/0.39410372
146	47	DVTNGLGTY	0.64/0.244054

147	625	CVNYDLYGI	0.71/0.17380978
148	355	VPSPLNWER	0.84/0.043685485
149	1213	MSTCAVNYT	0.76/0.12337384
150	182	HPNLGNHFK	0.23/0.64654865
151	703	CNYVFNNL	0.83/0.044655497
152	1088	GRLTALNAY	0.82/0.053010075
153	326	YTVQPIADV	0.65/0.22291911
154	757	KNRRSRRAI	0.12/0.7488378
155	142	SVVVQPRTI	0.99/-0.12375199
156	519	TNYLTCDNL	0.00/0.86476576
157	13	FAVIGDLNC	0.56/0.29687663
158	239	TKFLFNVYL	0.24/0.61649385
159	435	TTATSCQLY	0.98/-0.12369374
160	795	IQIPSEFTI	0.40/0.44760283
161	166	V SVCQYNMC	0.81/0.031736858
162	690	NSSEPALLF	0.20/0.64170804
163	1144	NAPYGLYFI	0.08/0.76023932
164	460	TWNKRFGFI	0.46/0.37223574
165	854	NELDDTTQL	0.95/-0.11790109
166	737	AISVQTCDL	0.83/0.0016002471
167	211	DVNATYLYF	0.04/0.79043859
168	1194	YTGSGYYYP	0.70/0.12167664
169	478	TGVFTNHSV	0.61/0.20905354
170	338	KPDLPCNI	0.04/0.77749754
171	222	YQEGGTFYA	0.10/0.71632531
172	491	HCFKAPKNF	0.94/-0.12924902
173	1260	YINVTFLDL	0.07/0.73527251
174	494	KAPKNFCPC	0.61/0.18369692
175	64	NTTLFLNGY	0.29/0.50264266
176	1250	TSVAPDLSL	0.71/0.08076331
177	253	HYYVMPLTC	0.99/-0.20029993
178	960	ENQISGYTL	0.42/0.36562494
179	974	SLFPPWTAA	0.09/0.69513002
180	208	FTYDVNATY	0.55/0.22922013
181	1061	AISASLQEI	0.33/0.44906176
182	406	DKFAIPNRR	0.85/-0.07174264
183	276	TPLTPRQYL	0.00/0.77599705
184	997	GLGVTMDVL	0.00/0.77527918
185	1313	GVAMLVLLF	/0.76736083

186	1282	LNHSYINLK	0.54/0.22670868
187	1092	ALNAYVSQQ	0.85/-0.096656997
188	962	QISGYTLAA	0.77/-0.020037817
189	56	YVLDRVYLN	0.34/0.40926692
190	1165	KVSPGLCIA	0.35/0.39918266
191	1310	CLAGVAMLV	/0.74744067
192	448	AANVSVSRF	0.04/0.70554959
193	725	YLGCVVNAY	0.17/0.57212179
194	678	CYSGRVSAA	0.03/0.7107723
195	382	ADSFTCNNI	0.08/0.66009748
196	1080	AQIDRLING	0.01/0.72729886
197	973	ASLFPPWTA	0.00/0.73282348
198	878	LKDGVNFNV	0.99/-0.25906731
199	388	NNIDAAKIY	0.54/0.18869519
200	787	EPVGGLYEI	0.17/0.55786069
201	941	DLICVQSYK	0.14/0.58405131
202	1240	EELDQWFKN	0.97/-0.25948147
203	699	RNIKCNYVF	0.82/-0.11039526
204	754	DYSKNRRSR	0.29/0.4176939
205	306	EIKCKTQSI	0.52/0.18669072
206	522	LTCDNLCTL	0.57/0.13631168
207	1230	TSIPNLPDF	0.32/0.3855904
208	996	NGLGVTMDV	0.62/0.079774929
209	289	QDGIIFNAV	0.00/0.69829607
210	778	TVNSVNDSL	0.00/0.69697261
211	714	QLQPINYSF	0.59/0.10689114
212	826	CGDYAACKL	0.82/-0.12324492
213	66	TLFLNGYYP	0.70/-0.0035975393
214	890	NFSPVLGCL	0.02/0.67492761
215	397	GMCFSITI	0.03/0.66154645
216	1022	AIQQGFDAT	0.72/-0.029113002
217	347	EAWLNDKSV	0.83/-0.14267198
218	129	AITIGSTFV	0.12/0.55958619
219	784	DSLEPVGGL	0.74/-0.060698413
220	944	CVQSYKGIK	0.99/-0.31229481
221	1098	SQQLSDSTL	0.80/-0.13068828
222	623	GVCVNYDLY	0.05/0.61745976
223	1158	PTKYVTAKV	0.85/-0.18294795

224	365	TFSNCNFNM	0.19/0.47604251
225	763	RAITGYRF	0.00/0.66576004
226	281	RQYLLAFNQ	0.88/-0.2160557
227	972	SASLFPPWT	0.95/-0.28683702
228	1293	GTYEYYVKW	/0.66268852
229	828	DYAACKLQL	0.06/0.59971396
230	1003	DVLSQNQKL	0.69/-0.0322844
231	1104	STLVKFSAA	0.70/-0.044472735
232	860	TQLQVANSL	0.01/0.6449722
233	18	DLNCPLDPR	0.53/0.1165145
234	158	NKLQGLLEV	0.14/0.50596177
235	340	DLPNCNIEA	1.00/-0.35617448
236	528	CTLDPITFK	0.07/0.57312374
237	599	ILHDVNNGL	0.00/0.64300865
238	668	ITNRTFMIH	0.01/0.63270915
239	592	CNIFANFIL	0.53/0.10874037
240	923	DVGFEAYN	0.96/-0.32327137
241	201	SCLYKRNFT	0.98/-0.34380037
242	363	RKTFSNCNF	0.83/-0.19796853
243	918	KVKLSDVGF	0.00/0.63123157
244	771	FTNFEPFTV	0.16/0.4672843
245	392	AAKIYGMCF	0.16/0.46608809
246	817	VTIDCAAFV	0.14/0.48426328
247	259	LTCISRRDI	0.90/-0.27793737
248	766	TTGYRFTNF	0.07/0.54384314
249	689	ANSSEPALL	0.62/-0.0077838697
250	702	KCNYVFNNS	0.62/-0.0081875223
251	380	IQADSFTCN	0.91/-0.29818823
252	295	NAVDCMSDF	0.19/0.42143147
253	291	GIIFNAVDC	0.93/-0.32119415
254	1084	RLINGRLTA	0.14/0.46520304
255	224	EGGTFYAYF	0.04/0.56507794
256	802	TIGNMEEFI	0.60/0.0014550198
257	911	IEDLLFDKV	0.70/-0.10481566
258	1130	NFCGNGNHI	0.11/0.47477456
259	843	CDNINAILT	0.93/-0.34822341
260	932	NCTGGAEIR	0.88/-0.30516226

261	87	GTDLLSTLW	0.99/-0.4160785
<b>262</b>	<b>1143</b>	<b>QNAPYGLYF</b>	<b>0.00/0.57365265</b>
263	1006	SQNQKLIAN	0.54/0.023150896
264	935	GGAEIRDLI	0.19/0.37275014
265	11	TAFAVIGDL	0.01/0.55076349
266	1239	KEELDQWFK	0.96/-0.40040811
267	92	STLWFKPPF	0.68/-0.12043133
268	849	ILTEVNELL	0.03/0.52947398
269	586	CLQGDKCNI	0.72/-0.16110256
270	994	RINGLGVTM	0.08/0.47722697
271	342	PNCNIEAWL	0.82/-0.26332217
272	965	GYTLAATSA	1.00/-0.44381501
273	866	NSLMNGVTL	0.09/0.465982
274	260	TCISRRDIG	0.67/-0.11730873
275	1072	RLDALEAEA	0.02/0.5316099
276	437	ATSCQLYYN	0.87/-0.3211714
277	424	GYLQSSNYR	0.01/0.53880786
278	60	RVYLNNTTLF	0.52/0.026021448
279	432	RIDTTATSC	0.90/-0.35685537
280	1272	MNRLQEAIK	0.00/0.541886
281	612	DLQKANTEI	0.72/-0.18053657
282	8	SLPTAFAVI	0.01/0.52927823
283	1137	HIISLVQNA	0.10/0.43765557
284	781	SVNDSLEPV	0.02/0.51138399
285	671	RTFMIHSCY	0.56/-0.029309502
286	206	RNFTYDVNA	0.72/-0.19035185
287	864	VANSLMNGV	0.60/-0.072806232
288	1113	QAMEKVNEC	0.63/-0.10346227
289	261	CISRRDIGF	0.10/0.42390412
290	20	NCPLDPRLK	1.00/-0.48350687
291	591	KCNIFANFI	0.04/0.4709378
292	619	EIELGVCVN	0.51/0.00065446274
293	1058	RFGAISASL	0.77/-0.25970955
294	1300	KWPWYVWLL	/0.50615282
295	358	PLNWERKTF	0.89/-0.38479811
296	979	WTAAAGVPF	0.02/0.483316
297	329	QPIADVYRR	0.01/0.49214361
298	1221	TKAPYVMLN	0.64/-0.13991298

299	1004	VLSQNQKLI	0.52/-0.022422484
300	156	GVNKLQGLL	0.01/0.48647109
301	1271	EMNRLQEAI	0.00/0.49444084
302	1055	LSNRFGAIS	0.84/-0.34586669
303	412	NRRKVDLQL	0.04/0.4505505
304	1056	SNRFGAISA	0.00/0.48756828
305	514	TCPAGTNYL	0.00/0.48576885
306	263	SRRDIGFTL	0.09/0.39504491
307	351	NDKSVPSPL	0.88/-0.39511623
308	476	QPTGVFTNH	0.04/0.44456138
309	1151	FIHFNYVPT	0.76/-0.27616536
310	855	ELDDTTQLQ	0.96/-0.47711623
311	1096	YVSQQLSDS	0.98/-0.49812739
312	931	NNCTGGAEI	0.06/0.42177788
313	846	INAILTEVN	0.91/-0.42993354
314	641	EVNATYYNS	0.63/-0.15279713
315	681	GRVSAAYHA	0.00/0.46978696
316	217	LYFHFYQEG	0.72/-0.25059039
317	666	DYITNRTFM	0.00/0.46560219
318	77	GSTYRNMAL	0.02/0.44499329
319	1291	DIGTYEYYV	/0.46381712
320	207	NFTYDVNAT	0.75/-0.28705922
321	124	YSEFPAITI	0.03/0.43068298
322	534	TFKAPDTYK	0.08/0.37780319
323	1262	NVTFLDLQV	0.69/-0.23278799
324	644	ATYYNSWQN	0.80/-0.34594176
325	194	HLDTGVVSC	0.00/0.45387067
326	588	QGDKCNIFA	1.00/-0.54648073
327	674	MIHSCYSGR	0.00/0.44851207
328	176	YPHTICHPN	0.69/-0.25099717
329	368	NCNFMSSL	0.00/0.43511016
330	1017	NNALHAIQQ	0.65/-0.21524797
331	616	ANTEIELGV	0.93/-0.50233594
332	654	LYDSNGNLY	0.01/0.41692321
333	202	CLYKRNFY	0.02/0.40617812
334	457	NPSTWKNRF	0.05/0.37604116
335	1202	PEPITENNV	0.03/0.39578709
336	758	NRRSRRAIT	0.03/0.39316024
337	275	VTPLTPRQY	0.53/-0.1079905

338	988	YLVNQYRIN	0.00/0.41799404
339	216	YLYFHFYQE	0.56/-0.14373867
340	244	NVYLGMAIS	0.01/0.40391588
341	1317	LVLLFFICC	/0.41201937
342	1192	WMYTGSGYY	0.74/-0.33025172
343	1082	IDRLINGRL	0.93/-0.52091464
344	691	SSEPALLFR	0.02/0.38692398
345	607	LTCSTDLOK	0.52/-0.11330045
346	1112	AQAMEKVNE	0.93/-0.52470152
347	453	VSRFNPSTW	0.01/0.39453873
348	1316	MLVLLFFIC	/0.40448677
349	712	TRQLQPINY	0.04/0.36290021
350	196	DTGVVSCLY	0.98/-0.57715749
351	1040	VVNANAEAL	0.01/0.39005797
352	403	ITIDKFAIP	0.60/-0.20006665
353	418	LQLGNLGYL	0.03/0.36814366
354	269	FTLEYWVTP	0.84/-0.44348394
355	605	NGLTCSTDL	0.93/-0.53401282
356	1309	ICLAGVAML	/0.39466956
357	48	VTNGLGTYT	0.03/0.36353205
358	458	PSTWNRFG	0.89/-0.497645
359	111	KVKNTKVFK	0.01/0.38051629
360	513	GTCPAGTNY	0.01/0.3764456
361	992	QYRINGLGV	0.00/0.38442158
362	1071	SRLDALEAE	0.59/-0.21367453
363	686	AYHANSSEP	0.59/-0.21487044
364	1101	LSDSTLVKF	0.69/-0.31845964
365	1204	PITENNVVV	0.70/-0.33970453
366	873	TLSTKLKDG	0.95/-0.5958323
367	943	ICVQSYKGI	0.81/-0.46412512
368	344	CNIEAWLND	0.58/-0.23931148
369	1241	ELDQWFKNQ	0.98/-0.64667895
370	1277	EAIKVLNHS	0.88/-0.55048873
371	1134	NGNHISLV	0.54/-0.21083952
372	955	PLLSENQI	0.65/-0.32494524
373	680	SGRVSAAYH	0.55/-0.22501637
374	852	EVNELDIT	0.86/-0.54075608
375	673	FMIHSCYSG	0.58/-0.26337873
376	302	DFMSEIKCK	0.63/-0.33209923



377	1135	GNHIISLVQ	0.80/-0.50584485
378	413	RRKVDLQLG	0.71/-0.41620099
379	516	PAGTNYLTC	0.92/-0.62977525
380	304	MSEIKCKTQ	0.69/-0.40222402
381	647	YNSWQNLLY	0.77/-0.49874867
382	35	DTGPPSIST	0.84/-0.56895408
383	858	DTTQLQVAN	0.96/-0.69962055
384	825	VCGDYAACK	0.86/-0.59972232
385	393	AKIYGMCFS	1.00/-0.75505759
386	218	YFHFYQEGG	0.87/-0.63256838
387	682	RVSAAYHAN	0.65/-0.41568367
388	110	AKVKNTKVF	0.86/-0.6297661
389	337	RKPDLPNCN	0.87/-0.64290963
390	163	LLEVSVCQY	0.74/-0.51421165
391	1211	VVMSTCAVN	0.57/-0.34500854
392	597	NFILHDVNN	0.70/-0.48115938
393	807	EEFIQTSSP	0.79/-0.57134192
394	488	YAQHCFKAP	0.98/-0.76170797
395	1167	SPGLCIAGN	0.89/-0.67335075
396	909	SAIEDLLFD	0.53/-0.31535
397	604	NNGLTCSTD	0.68/-0.47649818
398	706	VFNNSLTRQ	0.99/-0.79064358
399	1025	QGFDATNSA	0.97/-0.78151497
400	1150	YFIHFNYVP	0.84/-0.65757591
401	547	KSLVGIGEH	0.86/-0.67854085
402	132	IGSTFVNTS	0.69/-0.51562853
403	745	LTVGSGYCV	0.58/-0.41749306
404	894	VLGCLGSEC	0.99/-0.82944512
405	865	ANSLMNGVT	0.97/-0.82130289
406	1060	GAISASLQE	0.97/-0.82240708
407	743	CDLTVGSGY	0.64/-0.49756575
408	19	LNCPLDPRL	0.60/-0.46049867
409	5	LLISLPTAF	0.78/-0.64332399
410	362	ERKTFSNCN	0.51/-0.37516632
411	29	GSFNNRDTG	0.56/-0.43610297
412	1176	RGIAPKSGY	0.83/-0.70812954
413	274	WVTPLTPRQ	0.56/-0.44276648
414	179	TICHPNLGN	0.94/-0.83615439
415	183	PNLGNHFKE	0.82/-0.72560525

416	1160	KYVTAKVSP	0.88/-0.78985969
417	715	LQPINYSFD	0.70/-0.61148709
418	768	GYRFTNFEP	0.81/-0.72203329
419	305	SEIKCKTQS	0.81/-0.73675677
420	343	NCNIEAWLN	0.78/-0.70741334
421	834	LQLVEYGSF	0.59/-0.52017472
422	603	VNNGLTCS	0.81/-0.74144409
423	736	TAISVQTC	0.90/-0.83421642
424	602	DVNNGLTCS	0.65/-0.58624337
425	1030	TNSALVKIQ	0.70/-0.6403108
426	1232	IPNLPDFKE	0.71/-0.65284227
427	185	LGNHFKELW	0.53/-0.49048956
428	540	TYKCPQTKS	0.66/-0.634436
429	999	GVTMDVLSQ	0.54/-0.51486845
430	440	CQLYYNLPA	0.93/-0.90855461
431	734	NSTAISVQT	0.75/-0.72984075
432	160	LQGLLEVS	0.54/-0.5299666
433	370	NFMSSLMS	0.52/-0.51276669
434	733	YNSTAISVQ	0.83/-0.82465938
435	303	FMSEIKCKT	0.64/-0.64391806
436	792	LYEIQIPSE	0.92/-0.92620775
437	936	GAEIRDLIC	0.52/-0.53052956
438	1252	VAPDLSLDY	0.72/-0.73208717
439	170	QYNMCEYPH	0.52/-0.53476248
440	290	DGIIFNAVD	0.88/-0.89752012
441	661	LYGFRDYIT	0.60/-0.63273785
442	1102	SDSTLVKFS	0.96/-0.99682303
443	1249	QTSVAPDLS	0.93/-0.9701338
444	731	NAYNSTAIS	0.72/-0.76168058
445	915	LFDKVKLSD	0.91/-0.95737381
446	76	SGSTYRNMA	0.52/-0.56749591
447	800	EFTIGNMEE	0.74/-0.78988985
448	84	ALKGTDLLS	0.62/-0.67112019
449	1237	DFKEELDQW	0.99/-1.0653404
450	515	CPAGTNYLT	0.61/-0.68654272
451	143	VVVQPRTIN	0.92/-1.0031782
452	830	AACKLQLVE	0.99/-1.1013187
453	1118	VNECVKSQS	0.74/-0.85144344
454	1129	INFCGNGNH	0.66/-0.77528715

455	271	LEYWVTPLT	0.95/-1.0843846
456	626	VNYDLYGIS	0.88/-1.0144346
457	439	SCQLYYNLP	0.96/-1.1044019
458	126	EFPAITIGS	0.98/-1.1310392
459	164	LEVSVQCYN	0.52/-0.68212951
460	423	LGYLQSSNY	0.94/-1.1135489
461	379	FIQADSFTC	0.85/-1.0255097
462	738	ISVQTCDLT	0.69/-0.8835284
463	655	YDSNGNLYG	0.57/-0.76773168
464	624	VCVNYDLYG	0.90/-1.0994467
465	90	LLSTLWFKP	0.97/-1.1744841
466	294	FNAVDCMSD	0.63/-0.83843424
467	1023	IQQGFDATN	0.58/-0.79554506
468	1138	IISLVQNAP	0.64/-0.85762903
469	899	GSECSKASS	0.70/-0.94197746
470	414	RKVDLQLGN	0.64/-0.8833033
471	297	VDCMSDFMS	0.95/-1.199791
472	63	LNTTLFLNG	0.58/-0.83857498
473	582	SADSCLQGD	0.74/-1.0009609
474	433	IDTTATSCQ	0.97/-1.2312266
475	629	DLYGISGQG	0.95/-1.2499934
476	585	SCLQGDKCN	0.93/-1.2617111
477	94	LWFKPPFLS	0.73/-1.0640988
478	1275	LQEAIKVLN	0.85/-1.1890652
479	687	YHANSSEPA	0.99/-1.3494791
480	23	LDPRLKGSF	0.65/-1.0288432
481	192	LWHLDTGVV	0.63/-1.0241095
482	272	EYWVTPLTP	0.61/-1.0087404
483	815	PKVTIDCAA	0.96/-1.3703482
484	173	MCEYPHTIC	0.54/-0.97772655
485	1283	NHSYINLKD	0.79/-1.2538952
486	898	LGSECSKAS	0.97/-1.459627
487	893	PVLGCLGSE	0.79/-1.2837736
488	1016	FNNALHAIQ	0.62/-1.1205932
489	1247	KNQTSVAPD	0.86/-1.4031885
490	966	YTLAATSAS	0.74/-1.3078469
491	376	LMSFIQADS	0.68/-1.2753651
492	649	SWQNLLYDS	0.79/-1.3856862
493	317	PTGVYELNG	0.64/-1.2363928

	494	853	VNELDDTTQ	0.91/-1.5430266
	495	794	EIQIPSEFT	0.60/-1.2380885
	496	147	PRTINSTQD	0.66/-1.3513955
	497	17	GDLNCPLDP	0.98/-1.6959279
	498	874	LSTKLKDGV	0.76/-1.4797166
	499	652	NLLYDSNGN	0.65/-1.3889788
	500	1052	LQQLSNRFG	0.57/-1.3478396
	501	1226	VMLNTSIPN	0.56/-1.3608976
	502	85	LKGTDLLST	0.98/-1.8358649
	503	46	VDVTNGLGT	0.68/-1.6244871
<b>HCoV-HKU1</b>				
<b>Viral protein name</b>	<b>Rank</b>	<b>Start position</b>	<b>Sequence</b>	<b>Score ANN/SVM</b>
<b>Spike</b>	1	700	SYVLNNISL	0.82/1.7620527
	2	1230	KLSDFESEL	0.73/1.5428952
	3	43	SYGLGTYI	0.96/1.2092847
	4	1043	ALNSLLQQL	0.62/1.4388165
	5	1085	RLTALNAYV	1.00/1.0179382
	6	272	NRGVITNAV	0.99/1.0100408
	7	493	ASCPIGTNY	0.98/0.98864166
	8	752	SSSSRRKRR	0.99/0.93621214
	9	69	KSGANFRDL	0.29/1.6294666
	10	421	SSCQLYYSL	0.88/1.0333743
	11	1279	SLNNSYINL	0.65/1.2588282
	12	105	RVKNTKLYV	0.88/1.0142424
	13	1046	SLLQQLFNK	0.97/0.91980637
	14	1216	FTKAPLVYL	0.85/1.0336339
	15	426	YYSLPAINV	0.67/1.1909905
	16	710	TQPYFDSYL	0.97/0.88413644
	17	759	RRSISASYR	0.94/0.8998011
	18	316	TVHRRIPDL	0.97/0.82574329
	19	586	NRCNIFSNF	0.99/0.7827496
	20	1140	NAPYGLLFM	0.98/0.7818804
	21	828	ACHDLLSEY	0.78/0.968353
	22	1175	APKQGYFIK	0.93/0.81055523
	23	1119	KSQSPRINF	0.98/0.74189283
	24	993	GLGVTMDVL	0.88/0.77527918
	25	963	TTAATVAAM	0.99/0.63779509
	26	687	ASSLALLYR	0.91/0.71152114

27	190	TYNVSTDWL	0.51/1.0546429
28	583	VSNRCNIF	0.90/0.65970885
29	760	RSISASYRF	0.92/0.63541909
30	719	GCVFNADNL	0.88/0.63192236
31	298	LPNTGVYDL	0.56/0.92549997
32	1258	TINATFLDL	0.35/1.1252987
33	608	DLLQPNTTEV	0.26/1.2138412
34	1025	ATNSALAKI	0.91/0.54701037
35	116	TLYSEFSTI	0.96/0.49510647
36	756	RRKRRSISA	0.65/0.79842193
37	659	GFKDFVTNK	0.03/1.414362
38	41	DVSYGLGTY	0.86/0.5842947
39	102	IFSRVKNTK	0.64/0.78915923
40	20	CTNFAINDK	0.75/0.67871498
41	494	SCPIGTNYR	0.99/0.43613207
42	689	SLALLYRNL	0.10/1.3084838
43	66	YFPKSGANF	0.90/0.48607673
44	1248	IAPNLTLNL	0.87/0.49571062
45	770	TFEPFNVSF	0.71/0.64479112
46	1282	NSYINLKDI	0.96/0.36903748
47	557	GVLGDGSYNV	0.86/0.46236517
48	903	SRSFFEDLL	0.44/0.86461849
49	616	VFTDVCVDY	1.00/0.30138995
50	984	SLNVQYRIN	0.96/0.33455348
51	902	SSRSFFEDL	0.58/0.71380652
52	668	TYNIFPCYA	0.15/1.1424939
53	1076	VQIDRLING	0.98/0.30891106
54	1277	IKSLNNSYI	0.99/0.29657575
55	203	YQERGTFYA	0.92/0.35979012
56	754	SSRRKRRSI	0.29/0.97791422
57	1276	SIKSLNNSY	1.00/0.25899736
58	716	SYLGCVFNA	0.46/0.79580845
59	670	NIFPCYAGR	0.86/0.39556392
60	224	FSLYLGTL	0.90/0.34920887
61	380	GSCFKSIVL	0.52/0.72519269
62	698	KCSYVLNNI	0.09/1.1537617
63	1262	TFLDLYYEM	0.08/1.1465407
64	1029	ALAKIQSVV	0.99/0.2335248
65	824	SNYAACHDL	0.67/0.55271113

66	1039	SNAQALNSL	0.53/0.69256061
67	663	FVTNKTYNI	0.92/0.29439927
68	1213	SVNFTKAPL	0.65/0.56402523
69	34	RISEYVVDV	0.60/0.61107435
70	31	TVPRISEYV	0.84/0.36983368
71	123	TIVIGSVFI	0.75/0.45831722
72	121	FSTIVIGSV	0.89/0.31395181
73	50	YILDRVYLN	0.88/0.32378538
74	520	CLPDPITAY	0.94/0.24984492
75	1068	RLDALEAQV	0.16/1.0275264
76	1106	GAALAMEKV	0.97/0.2122842
77	694	YRNLCESYV	0.03/1.1489538
78	1036	VVNSNAQAL	0.55/0.62669954
79	44	YGLGTYIIL	0.51/0.66332393
80	587	RCNIFSNFI	0.57/0.59998654
81	168	SSRNESWHF	0.94/0.22365044
82	940	CVQSFNGIK	0.97/0.18017776
83	1310	SFIIFLVLL	/1.1462725
84	799	TIVGQEEFI	0.99/0.14878547
85	336	FNVPSPLNW	0.87/0.26876333
86	684	HQNASSLAL	0.60/0.52976512
87	1280	LNNSYINLK	0.98/0.14938808
88	1173	GIAPKQGYF	0.89/0.23258714
89	768	FVTFEPFNV	0.74/0.38140978
90	667	KTYNIFPCY	0.96/0.16056305
91	167	GSSRNESWH	0.67/0.44543268
92	664	VTNKTYNIF	0.65/0.46142895
93	154	TMCEYPHTI	0.88/0.22894775
94	259	TPLSKRQYL	0.69/0.41730156
95	969	AAMFPPWSA	0.96/0.14246828
96	566	SCLCSTDAF	0.78/0.32080457
97	98	FNNGIFSRV	0.50/0.59960068
98	763	SASYRFVTF	0.42/0.66799322
99	1081	LINGRLTAL	0.08/1.0044934
100	216	SGMPTTFLF	0.87/0.21339683
101	30	TTVPRISEY	0.49/0.59254183
102	638	VSAVYYNSW	0.76/0.31132417
103	749	SPSSSSSRR	0.98/0.086673944
104	1113	KVNECVKSQ	0.89/0.16973449

105	1061	SLQEILSRL	0.00/1.0593795
106	933	SEIRDLLCV	0.97/0.088650187
107	55	VYLNTTILF	0.94/0.11184547
108	1008	IATAFNAL	0.86/0.18873703
109	1120	SQSPRINFC	0.71/0.33327033
110	512	HTDWCRSC	0.76/0.27946536
111	713	YFDSYLGCV	0.99/0.047453498
112	623	DYDLYGITG	0.93/0.10714629
113	271	DNRGVITNA	0.38/0.6541886
114	1309	FSFIIFLVL	/1.02016
115	1181	FIKHNDHWM	0.34/0.67452595
116	456	SSHSVVYSR	0.16/0.8521888
117	438	NYNPSSWNR	0.01/1.0021574
118	269	KFDNRGVIT	0.92/0.091904369
119	825	NYAACHDLL	0.92/0.090832085
120	766	YRFVTFEPF	0.65/0.35859663
121	838	TFCDNINSI	0.01/0.99608652
122	857	TQLHVADTL	0.14/0.85599145
123	706	ISLATQPYF	0.51/0.48360961
124	835	EYGTFCdni	0.00/0.98854271
<b>125</b>	<b>1296</b>	<b>YVKWPWYVW</b>	<b>/0.9882908</b>
126	67	FPKSGANFR	0.00/0.98805195
127	229	GTLLSHYYV	0.38/0.60497684
128	282	CSSSFFSEI	0.37/0.61421477
129	83	TYLSTLWYQ	0.92/0.057406196
130	445	NRRYGFNNF	0.00/0.97131068
131	758	KRRSISASY	0.46/0.50867717
132	941	VQSFNGIKV	0.59/0.3746817
133	99	NNGIFSRVK	0.46/0.5046264
134	6	FILPTTLAV	0.12/0.8440887
135	771	FEPFNVSFV	0.00/0.96360683
136	1295	MYVKWPWYV	/0.96271032
137	982	PFSLNVQYR	0.70/0.25872781
138	351	NCNFNLSTL	0.08/0.87856975
139	1220	PLVYLNHSV	0.99/-0.032345436
140	793	KIPTNFTIV	0.23/0.72663501
141	942	QSFNGIKVL	0.83/0.12580053
142	927	NNCTGGSEI	0.86/0.090929945
143	192	NVSTDWLYF	0.56/0.39015094

144	195	TDWLYFHFY	0.98/-0.032061155
145	790	YEIKIPTNF	0.77/0.17685198
146	1101	SLVKFGAAL	0.04/0.90519989
147	1084	GRLTALNAY	0.89/0.053010075
148	218	MPTTFLFSL	0.30/0.64085994
149	1217	TKAPLVYLN	0.90/0.039178237
150	53	DRVYLNTTI	0.99/-0.052404722
151	784	ESVGGLYEI	0.61/0.32512498
152	582	CVSNNRCNI	0.01/0.92421002
153	775	NVSFVNDSI	0.27/0.66251473
154	896	GPHCGSSSR	0.75/0.18243433
155	641	VYYNSWQNL	0.01/0.91599745
156	15	IGDFNCTNF	0.57/0.35013327
157	344	WERKIFSNC	0.81/0.10647642
158	179	SEPLCLFKK	0.93/-0.013901221
159	649	LLYDSNGNI	0.73/0.18122959
160	447	RYGFNNFNL	0.02/0.89119812
161	755	SRRKRRSIS	0.89/0.017578754
162	76	DLSLKGTTY	0.67/0.23461964
163	880	HFDVDNINF	0.69/0.21297986
164	1102	LVKFGAALA	0.63/0.26450716
165	1246	TSIAPNLT	0.66/0.22695478
166	839	FCDNINSIL	0.87/0.014076705
167	1254	LNLHTINAT	0.95/-0.070832522
168	821	FVCSNYAAC	0.00/0.87835355
169	1223	YLNHSVPKL	0.00/0.87416547
170	261	LSKRQYLLK	0.40/0.47387658
171	731	SVSSCALRM	0.06/0.81062214
172	262	SKRQYLLKF	0.97/-0.1017003
173	311	VKPVATVHR	0.99/-0.13332111
174	379	YGSCFKSIV	0.12/0.73650034
175	478	KPSFASSCK	0.85/0.0063641689
176	858	QLHVADTLM	0.67/0.18622172
177	1023	FSATNSALA	0.92/-0.066135006
178	470	NNTFCPCAK	0.67/0.18375089
179	1133	HILSLVQNA	0.38/0.46644373
180	979	AGIPFSLNV	0.59/0.25530636
181	674	CYAGRVSAA	0.28/0.5643572
182	792	IKIPTNFTI	0.79/0.052412036



183	204	QERGTFYAY	0.31/0.52384156
184	90	YQKPFLSDF	0.00/0.83050707
185	921	GFVEAYNNC	0.72/0.11041821
186	452	NFNLSSHVS	0.65/0.17807557
187	82	TTYLSTLWY	0.96/-0.13333662
188	810	NSPKVTIDC	0.92/-0.097998412
189	32	VPRISEYVV	0.91/-0.089489515
190	878	NLHFDVDNI	0.00/0.81835696
191	205	ERGTFYAYY	0.19/0.62816863
192	213	YADSGMPTT	0.79/0.026366717
193	126	IGSVFINNS	0.99/-0.17778451
194	1301	WYVWLLISF	/0.81108318
195	682	AFHQNASL	0.23/0.5762915
196	323	DLPDCDIDK	0.55/0.24927616
197	375	ESKIYGSCF	0.23/0.56911828
198	442	SSWNRRYGF	0.13/0.66621604
199	468	SVNNTFCPC	0.69/0.10515734
200	1018	SIQNGFSAT	0.94/-0.14581676
201	424	QLYYSLPAI	0.91/-0.11824045
202	715	DSYLGCVFN	0.89/-0.1002855
203	1307	ISFSFIIFL	/0.78575864
204	870	TLSSNLNTN	0.53/0.25374782
205	975	WSAAAGIPF	0.42/0.36285401
206	802	GQEEFIQTN	0.95/-0.16816374
207	1117	CVKSQSPRI	0.24/0.53208529
208	1049	QQLFNKFGA	1.00/-0.22821043
209	983	FSLNVQYRI	0.00/0.77109989
210	553	EEKCGVLDG	0.94/-0.16928054
211	842	NINSILDEV	0.76/0.0093573244
212	355	NLSTLLRLV	0.21/0.55837403
213	296	SLLPNTGVY	0.79/-0.022870965
214	774	FNVSFVND	0.99/-0.22480162
215	268	LKFDNRGVI	0.96/-0.19895001
216	319	RRIPDLPC	0.95/-0.19023402
217	1199	EPISDKNVV	0.98/-0.22196621
218	621	CVDYDLYGI	0.98/-0.22576904
219	1255	NLHTINATF	0.02/0.7275797
220	990	RINGLGVTM	0.27/0.47722697
221	609	LLQPNTVEF	0.92/-0.17327496

222	431	AINVTINNY	0.01/0.73662746
223	446	RRYGFNNFN	0.95/-0.20542774
224	764	ASYRFVTFE	0.71/0.034522759
225	1344	GHHDFVIKT	/0.73421119
226	914	KVKLSDVGF	0.10/0.63123157
227	84	YLSTLWYQK	0.10/0.62805235
228	283	SSSFFSEIQ	0.97/-0.24306024
229	250	DNETLQYWV	0.99/-0.2639652
230	569	CSTDAFLGW	0.36/0.36553576
231	148	ITACQYTCM	0.66/0.062803268
232	230	TLLSHYYVL	0.03/0.69149378
233	367	SFSCNNFDE	0.95/-0.23018078
234	24	AINDKNTTV	0.00/0.71801456
235	1136	SLVQNAPYG	0.51/0.20785211
236	968	VAAMFPPWS	0.91/-0.19743027
237	584	SNNRCNIFS	0.70/0.010838591
238	1153	KPISFKTVL	0.14/0.56974466
239	220	TTFLFSLYL	0.12/0.58921455
240	1235	ESELSHWFK	0.87/-0.16129879
241	591	FSNFILNGI	0.03/0.67748901
242	831	DLLSEYGTF	0.53/0.17644984
243	1126	NFCGNGNHI	0.23/0.47477456
244	1094	SQQLSDISL	0.34/0.36361471
245	476	CAKPSFASS	0.93/-0.2329537
246	912	FDKVKLSDV	0.71/-0.013587218
247	696	NLKCSYVLN	0.78/-0.084990188
248	378	IYGSCFKSI	0.26/0.43265229
249	778	FVNDIESV	0.02/0.67162651
250	1211	TCSVNFTKA	0.95/-0.26021539
251	303	VYDLSGFTV	0.05/0.63793452
252	71	GANFRDLSL	0.10/0.58244558
253	661	KDFVTNKTY	0.96/-0.27860918
254	861	VADTLMQGV	0.89/-0.21024097
255	1313	IFLVLLFFI	/0.67886396
256	845	SILDEVNGL	0.01/0.66860581
257	279	AVDCSSSFF	0.01/0.66734623
258	393	IPNSRRSDL	0.07/0.60429302
259	94	FLSDFNNGI	0.00/0.67069999
260	181	PLCLFKKNF	0.85/-0.17951239

261	109	TKLYVNKTL	0.76/-0.089771842
262	371	NNFDESKIY	0.54/0.12834157
263	309	FTVKPVATV	0.05/0.61695832
264	617	FTDVCVDYD	0.95/-0.28306966
265	736	ALRMGSGFC	0.84/-0.17342683
266	97	DFNNGIFSR	0.23/0.42996947
267	574	FLGWSYDTC	0.67/-0.010117306
268	1145	LLFMHFSYK	0.14/0.51861318
269	901	SSSRSFFED	0.84/-0.18454701
270	436	INNYPSSW	0.67/-0.019908136
271	383	FKSIVLDKF	0.90/-0.25461354
272	310	TVKPVATVH	0.62/0.025081417
273	386	IVLDKFAIP	0.96/-0.31530854
274	818	CSLFVCSNY	0.09/0.55363702
275	1251	NLTLNLHTI	0.01/0.63317409
276	265	QYLLKFDNR	0.00/0.64215101
277	637	EVSAVYYNS	0.73/-0.090024514
278	95	LSDFNNGIF	0.99/-0.35154806
279	461	VYSRYCFSV	0.00/0.63844054
280	613	NTEVFTDVC	0.93/-0.29226743
281	345	ERKIFSNCN	0.88/-0.24336764
282	737	LRMGSGFCV	0.85/-0.21364811
283	1096	QLSDISLVK	0.07/0.56491084
284	1278	KSLNNSYIN	0.93/-0.29574406
285	1009	ATAFNNALL	0.02/0.61373012
286	642	YYNSWQNLL	0.00/0.6301557
287	1201	ISDKNVVFM	0.00/0.62784627
288	244	AISSNTDNE	0.98/-0.35224845
289	1064	EILSRLDAL	0.01/0.61409618
290	680	SAAFHQNAS	0.92/-0.29897792
291	1195	YYYPEPISD	1.00/-0.37990603
292	49	YYILDRVYL	0.00/0.61815747
293	761	SISASYRFV	0.03/0.58708429
294	628	GITGQGIFK	0.64/-0.025004151
295	249	TDNETLQYW	0.68/-0.06721555
296	705	NISLATQPY	0.55/0.062014224
297	407	GFLQSSNYK	0.03/0.57213049
298	312	KPVATVHRR	0.02/0.57630943
299	513	TDWCRCSCSCL	0.17/0.42197058

300	1289	DIGTYEMYV	/0.59184516
301	194	STDWLYFHF	0.01/0.58049051
302	707	SLATQPYFD	0.92/-0.33140434
303	612	PNTEVFTDV	0.85/-0.26779246
<b>304</b>	<b>1139</b>	<b>QNAPYGLLF</b>	<b>0.10/0.48203718</b>
305	22	NFAINDKNT	0.86/-0.29060754
306	676	AGRVSAAFH	0.99/-0.42362248
307	619	DVCVDYDLY	0.84/-0.27545188
308	354	FNLSTLLRL	0.01/0.55416564
309	420	SSSCQLYYS	0.91/-0.34786364
310	428	SLPAINVTI	0.11/0.4508478
311	846	ILDEVNGLL	0.13/0.43043316
312	1312	IIFLVLLFF	/0.55391827
313	237	VLPLTCNAI	0.17/0.38390309
314	1169	SGDVGIAPK	0.58/-0.026446645
315	212	YYADSGMPT	0.76/-0.20649614
316	823	CSNYAACHD	0.63/-0.076609213
317	956	ESQISGYTT	0.98/-0.43115406
318	104	SRVKNTKLY	0.94/-0.39415576
319	1249	APNLTLNLH	0.10/0.44537961
320	108	NTKLYVNKT	0.76/-0.21692109
321	92	KPFLSDFNN	0.60/-0.057320616
322	978	AAGIPFSLN	0.70/-0.15877418
323	1334	KCHNCCDEY	/0.53725745
324	170	RNESWHFDK	0.87/-0.33295051
325	1027	NSALAKIQS	0.92/-0.3902425
326	1304	WLLISFSFI	/0.52219583
327	1007	LIATAFNNA	0.51/0.0078548036
328	573	AFLGWSYDT	0.83/-0.31303054
329	285	SFFSEIQCK	0.87/-0.35358949
330	967	TVAAMFPPW	0.53/-0.017996401
331	656	NIIGFKDFV	0.72/-0.20812647
332	1035	SVVNSNAQA	0.64/-0.1282722
333	80	KGTTYLSTL	0.66/-0.14924444
334	1109	LAMEKVNEC	0.74/-0.22937302
335	1298	KWPWYVWLL	/0.50615282
336	686	NASSLALLY	0.52/-0.016611144
337	1013	NNALLSIQN	0.98/-0.47714226

338	110	KLYVNKTLY	0.51/-0.0086607937
339	63	FTGYFPKSG	0.87/-0.37039924
340	929	CTGGSEIRD	0.94/-0.44560625
341	567	CLCSTDAFL	0.00/0.48113686
342	765	SYRFVTFEP	0.90/-0.42077695
343	113	VNKTLYSEF	0.56/-0.084744278
344	1080	RLINGRLTA	0.01/0.46520304
345	1091	AYVSQQLSD	0.57/-0.096254408
346	1306	LISFSFIIF	/0.47340461
347	87	TLWYQKPFL	0.00/0.47070531
348	1058	ISSSLQEIL	0.06/0.40879305
349	208	TFYAYYADS	0.91/-0.44478649
350	1047	LLQQLFNKF	0.56/-0.096387161
351	729	DYSVSSCAL	0.01/0.45349998
352	643	YNSWQNLLY	0.96/-0.49874867
353	370	CNNFDESKI	0.06/0.39783651
354	278	NAVDCSSSF	0.00/0.45743779
355	1070	DALEAQVQI	0.00/0.45586622
356	56	YLNTTILFT	0.96/-0.51013712
357	815	TIDCSLFVC	0.84/-0.3902094
358	919	DVG FVEAYN	0.77/-0.32327137
359	881	FDVDNINFK	0.01/0.4343559
360	137	IVVQPHNGV	0.01/0.43382035
361	1160	VLVSPGLCI	0.60/-0.16010756
362	1244	NQTSIAPNL	0.00/0.42928267
363	369	SCNNFDESK	0.72/-0.29175091
364	1311	FIIFLVLLF	/0.4254223
365	234	HYYVLPLTC	0.95/-0.53184335
366	995	GVTMDVLNK	0.05/0.36752088
367	1315	LVLLFFICC	/0.41201937
368	301	TGVYDLSGF	0.99/-0.58161554
369	1011	AFNNALLSI	0.00/0.40789941
370	136	TIVVQPHNG	0.87/-0.47087208
371	14	VIGDFNCTN	0.86/-0.46146459
372	358	TLLRLVHTD	0.00/0.39603874
373	1340	DEYGGHHDF	/0.39453733
374	646	WQNLLYDSN	0.89/-0.49568818
375	140	QPHNGVLEI	0.03/0.36182254

376	191	YNVSTDWLY	0.67/-0.28154727
377	593	NFILNGINS	0.01/0.37841412
378	988	QYRINGLGV	0.00/0.38442158
379	1266	LYYEMNLIQ	0.99/-0.60719617
380	1274	QESIKSLNN	0.66/-0.27918132
381	1168	ISGDVGIAP	0.54/-0.16140511
382	225	SLYLGTTLS	0.74/-0.36583862
383	1291	GTYEMYVKW	/0.37300047
384	1284	YINLKDIGT	0.81/-0.43769982
385	117	LYSEFSTIV	0.00/0.37138757
386	246	SSNTDNETL	0.00/0.3675537
387	945	NGIKVLPPI	0.55/-0.1831929
388	808	QTNSPKVTI	0.00/0.36370318
389	1308	SFSFIIFLV	/0.36348684
390	1203	DKNVVFMT	0.68/-0.32249778
391	373	FDESKIYGS	0.82/-0.46546892
392	1104	KFGAALAME	0.98/-0.62602726
393	757	RKRRSISAS	0.76/-0.41062382
394	1034	QSVVNSNAQ	0.85/-0.50250075
395	28	KNTTVPRIS	0.84/-0.49389475
396	669	YNIFPCYAG	0.52/-0.17993155
397	466	CFSVNNTFC	0.59/-0.25503
398	973	PPWSAAAGI	0.86/-0.53075691
399	253	TLQYWVTPL	0.86/-0.54234401
400	703	LNNISLATQ	1.00/-0.69561064
401	1261	ATFLDLYYE	0.68/-0.37705261
402	867	QGVTLSSNL	0.52/-0.21802335
403	396	SRRSDLQLG	0.65/-0.35071912
404	961	GYTTAATVA	0.84/-0.54999436
405	856	TTQLHVADT	0.87/-0.58132348
406	500	NYRSCESTT	0.64/-0.374593
407	36	SEYVVDVSY	0.54/-0.27510673
408	644	NSWQNLLYD	0.61/-0.35769978
409	175	HFDKSEPLC	0.70/-0.45435513
410	5	IFILPTTLA	0.92/-0.67899069
411	1033	IQSVVNSNA	0.62/-0.38184421
412	805	EFIQTNSPK	0.90/-0.66261155
413	164	KSKGSSRNE	0.55/-0.32326481
414	89	WYQKPFLSD	0.66/-0.44014453

415	955	SESQISGYT	0.92/-0.70246871
416	96	SDFNNGIFS	0.77/-0.57073891
417	60	TILFTGYFP	0.86/-0.66235025
418	600	NSGTTCSND	0.54/-0.34397566
419	177	DKSEPLCLF	0.56/-0.36458908
420	1044	LNSLLQQLF	0.75/-0.56321456
421	26	NDKNTTVPR	0.56/-0.38498303
422	276	ITNAVDCSS	0.96/-0.79727375
423	741	SGFCVDYNS	0.96/-0.80289721
424	965	AATVAAMFP	0.96/-0.80290496
425	403	LGSSGFLQS	0.95/-0.79772719
426	480	SFASSCKSH	0.52/-0.37536045
427	267	LLKFDNRGV	0.84/-0.70074456
428	335	NFNVP SPLN	0.80/-0.66989397
429	1031	AKIQSVVNS	0.57/-0.44119322
430	620	VCVDYDLYG	0.92/-0.79279366
431	742	GFCVDYNSP	0.70/-0.58838526
432	657	IIGFKDFVT	0.86/-0.74926241
433	746	DYNSPSSSS	0.86/-0.75441479
434	827	AACHDLLSE	0.73/-0.62930206
435	1221	LVYLNHSVP	0.78/-0.67958717
436	934	EIRDLLCVQ	0.99/-0.89398253
437	221	TFLFSLYLG	0.69/-0.60269387
438	1193	SSYYYPEPI	0.59/-0.50616665
439	892	VGCLGPHCG	0.72/-0.64089218
440	397	RRSDLQLGS	0.60/-0.527383
441	280	VDCSSSFFS	0.85/-0.78407158
442	606	SNDLLQPNT	0.72/-0.65751758
443	333	LNNFNVPSP	0.94/-0.89129025
444	1132	NHILSLVQN	0.78/-0.74707211
445	882	DVDNINFKS	0.79/-0.7575608
446	776	VSFVNDSIE	0.85/-0.82012771
447	392	AIPNSRRSD	0.81/-0.78256021
448	734	SCALRMGSG	0.51/-0.49183573
449	672	FPCYAGRVS	0.51/-0.5027555
450	1191	TGSSYYYPE	0.72/-0.72422561
451	517	RCSCLPDPI	0.83/-0.83569933
452	326	DCDIDKWLN	0.74/-0.7476457
453	789	LYEIKIPTN	0.62/-0.63294201

454	811	SPKVTIDCS	0.88/-0.89853357
455	62	LFTGYFPKS	0.63/-0.65634241
456	172	ESWHFDKSE	0.83/-0.85967749
457	647	QNLLYDSNG	0.62/-0.6549701
458	401	LQLGSSGFL	0.99/-1.0250406
459	539	LVGVGEHCA	0.62/-0.66037379
460	256	YWVTPLSKR	0.54/-0.58719918
461	1075	QVQIDRLIN	0.71/-0.75773694
462	1090	NAYVSQQLS	0.67/-0.72905555
463	1163	SPGLCISGD	0.83/-0.89250164
464	1228	VPKLSDFES	0.61/-0.68824535
465	495	CPIGTNYRS	0.56/-0.65088614
466	1170	GDVGIAPKQ	0.90/-0.99249942
467	258	VTPLSKRQY	0.53/-0.6315164
468	423	CQLYYSLPA	0.85/-0.960454
469	565	VSCLCSTDA	0.58/-0.69245581
470	563	YNVSCLCST	0.68/-0.79259981
471	361	RLVHTDSFS	0.53/-0.65251529
472	381	SCFKSIVLD	0.62/-0.74942087
473	807	IQTNSPKVT	0.99/-1.1300118
474	683	FHQNASSLA	0.93/-1.0713918
475	1283	SYINLKDIG	0.72/-0.86827363
476	1086	LTALNAYVS	0.76/-0.90941691
477	551	VDEEKCGVL	0.64/-0.81204896
478	1260	NATFLDLYY	0.53/-0.72353102
479	1148	MHFSYKPIS	0.90/-1.094533
480	467	FSVNNTFCP	0.68/-0.88764661
481	1135	LSLVQNAPY	0.87/-1.0784129
482	434	VTINNYNPS	0.79/-1.0007724
483	1005	QKLIATAFN	0.92/-1.1477793
484	1257	HTINATFLD	0.70/-0.92788208
485	491	PSASCPIGT	0.54/-0.78325459
486	288	SEIQCKTKS	0.97/-1.2177181
487	1016	LLSIQNGFS	0.79/-1.0416696
488	708	LATQPYFDS	0.95/-1.2061844
489	559	LDGSYNVSC	0.63/-0.88731451
490	233	SHYYVLPLT	0.73/-0.98840856
491	599	INSGTTCSN	0.51/-0.77381362
492	1198	PEPISDKNV	0.53/-0.79441812



493	300	NTGVYDL SG	0.81/-1.0778341
494	1271	NLIQESIKS	0.55/-0.83062557
495	484	SCKSHKPPS	0.94/-1.2230892
496	155	MCEYPHTIC	0.68/-0.97772655
497	1243	KNQTSIAPN	0.95/-1.2651821
498	356	LSTLLRLVH	0.97/-1.2914599
499	745	VDYN SPSSS	0.81/-1.1406725
500	254	LQYWVTPLS	0.90/-1.2358166
501	888	FKSLVGCLG	0.67/-1.0074895
502	518	CSCLPDPIT	0.73/-1.0764705
503	416	IDTSSSCQ	0.97/-1.3411052
504	68	PKSGANFRD	0.78/-1.1525222
505	275	VITNAVDCS	0.92/-1.2972589
506	704	NNISLATQP	0.61/-0.99267762
507	1093	VSQQLSDIS	0.81/-1.1971889
508	506	STTVLDHTD	0.93/-1.324689
509	239	PLTCNAISS	0.52/-0.92619456
510	895	LGP HCGSSS	0.97/-1.3881777
511	299	PNTGVYDLS	0.84/-1.2595447
512	492	SASCPIGTN	0.60/-1.0339423
513	91	QKPFLSDFN	0.68/-1.124113
514	165	SKGSSRNES	0.61/-1.0572177
515	75	RDLSLKGTT	0.93/-1.3958825
516	772	EPFNVSFVN	0.51/-0.98344697
517	865	LMQGVTLSS	0.80/-1.3086614
518	173	SWHFDKSEP	0.52/-1.0291541
519	497	IGTNYRSCE	0.78/-1.304046
520	228	LGTL LSHYY	0.79/-1.3209309
521	106	VKNTKLYVN	0.66/-1.2002027
522	437	NNYNPSSWN	0.66/-1.2428289
523	2	LLIIFILPT	0.66/-1.2466662
524	556	CGVLDGSYN	0.66/-1.2651568
525	1042	QALNSLLQQ	0.60/-1.2210328
526	88	LWYQKPFLS	0.61/-1.2685105
527	1030	LAKIQSVVN	0.51/-1.1870013
528	568	LCSTDAFLG	0.99/-1.6797284
529	114	NKTLYSEFS	0.57/-1.3044412
530	1252	LTLNLHTIN	0.95/-1.7257947
531	952	PILSESQIS	0.67/-1.5090783

	532	1069	LDALDAQVQ	0.61/-1.5813453
	533	913	DKVKLSDBG	0.79/-1.8528379
	534	1250	PNLTLNLHT	0.52/-1.6937584
	535	847	LDEVNGLLD	0.57/-1.8290063