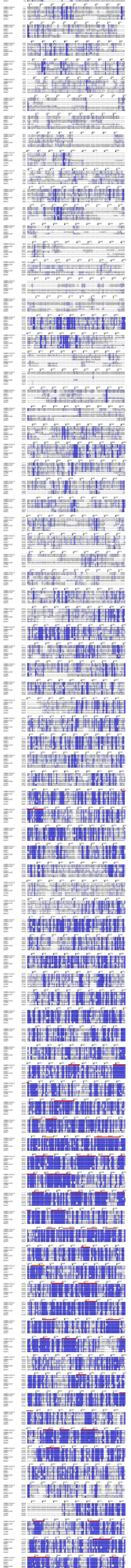
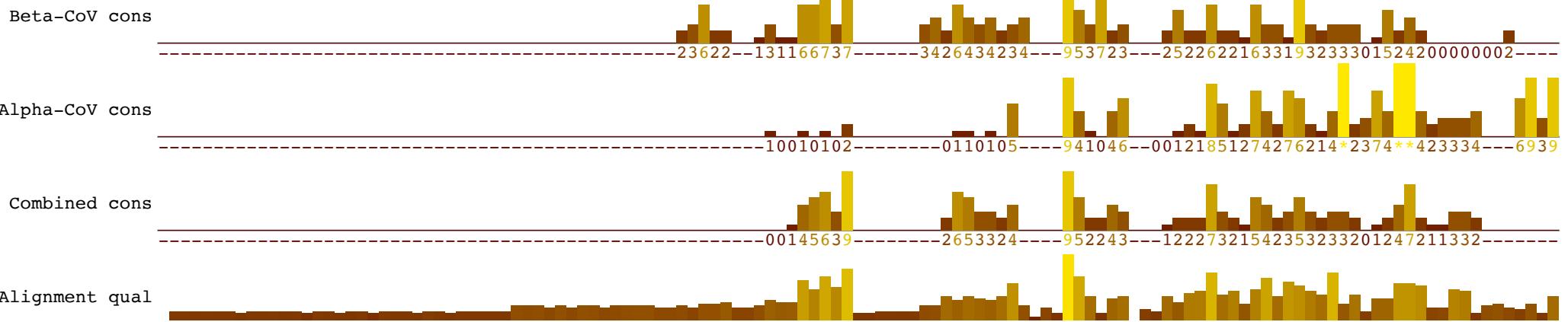
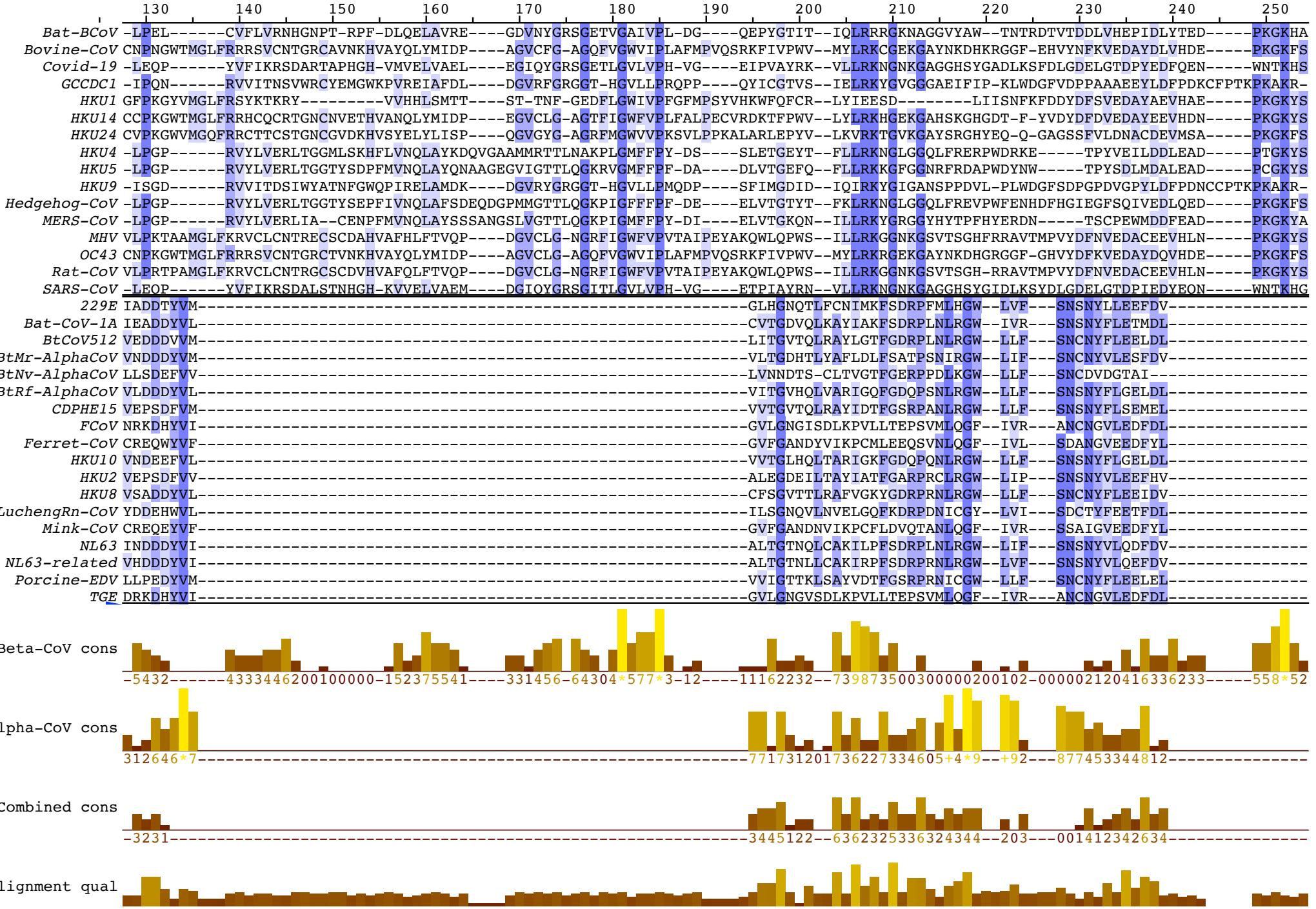
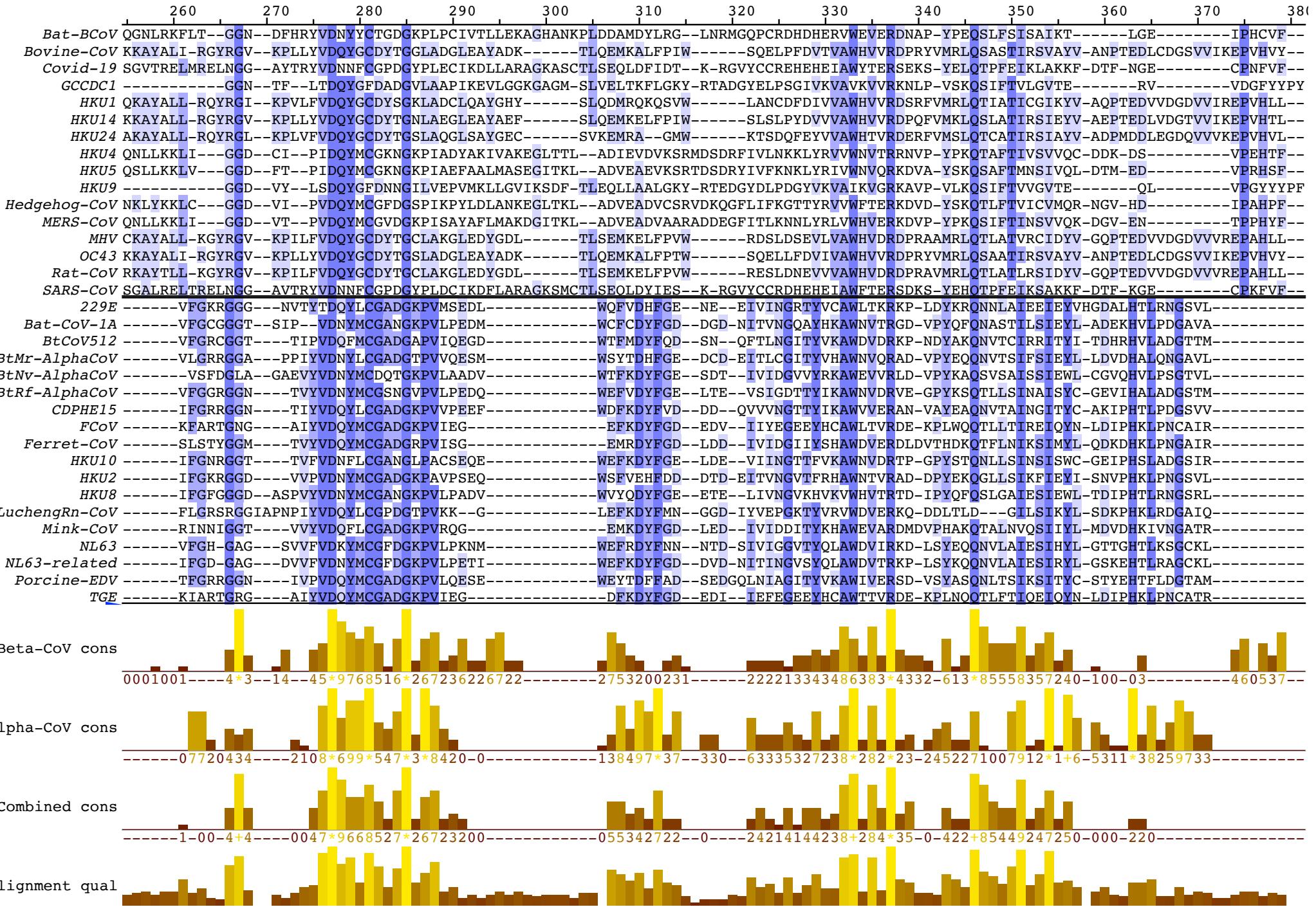


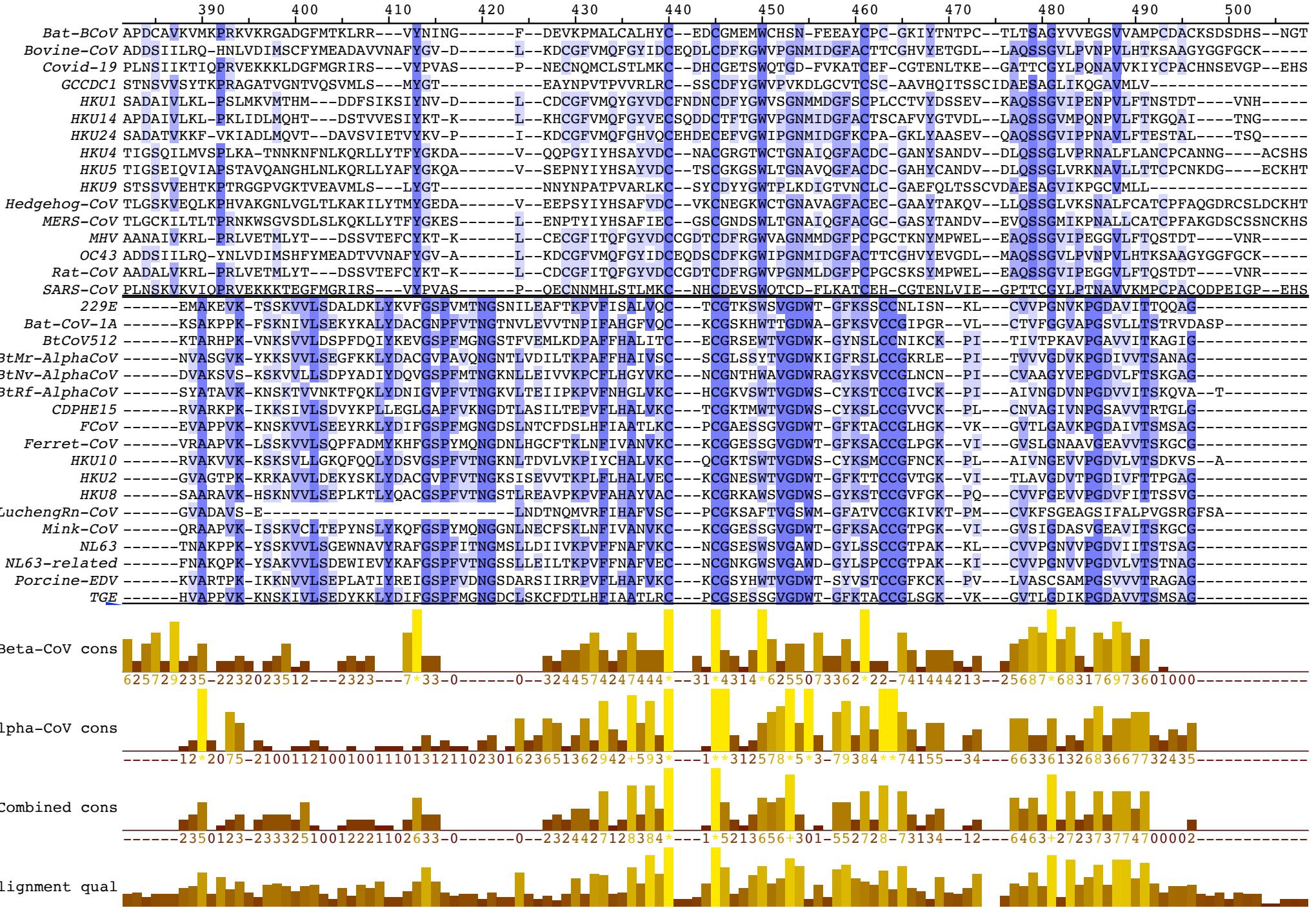
# ORF1ab polyprotein

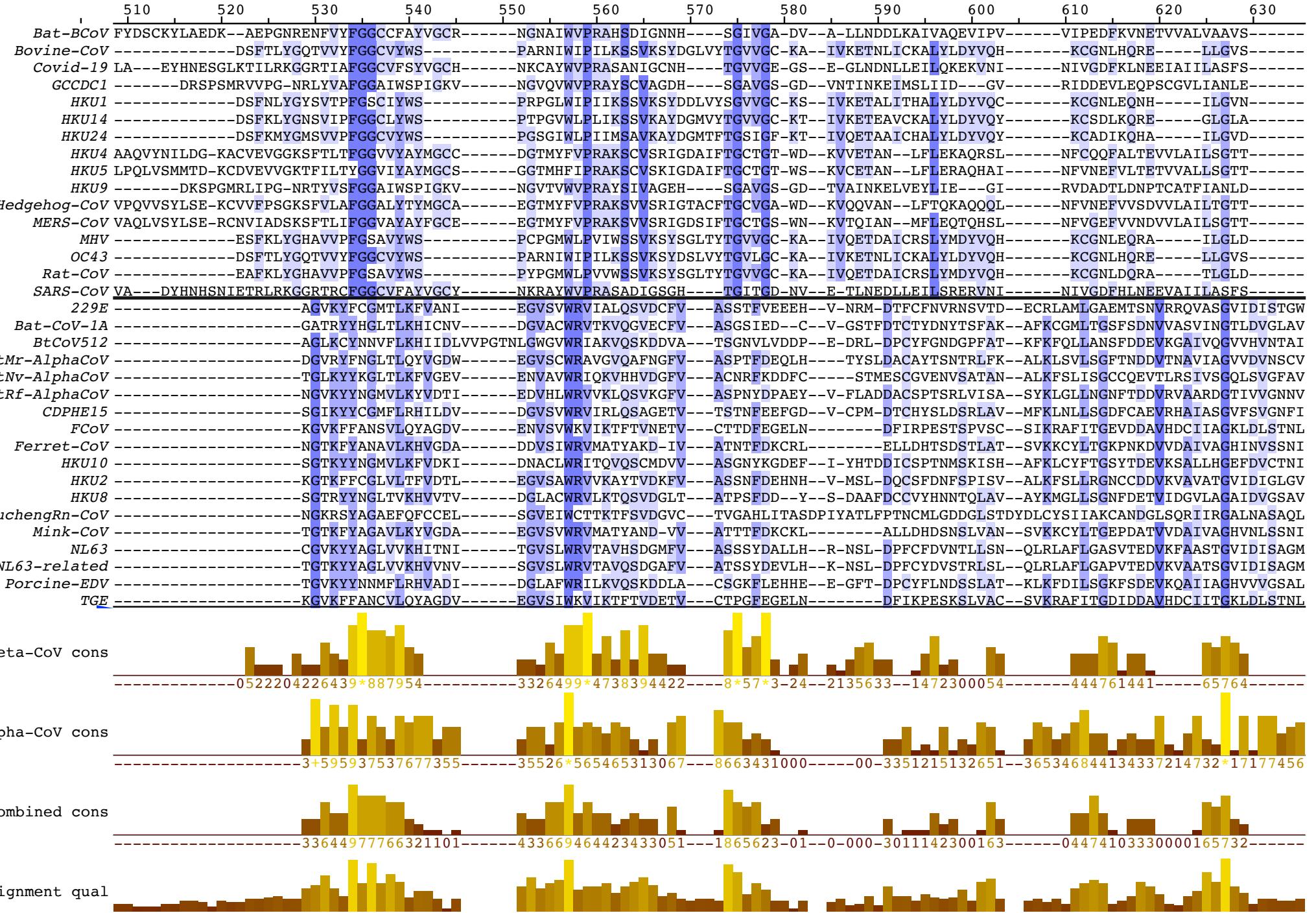


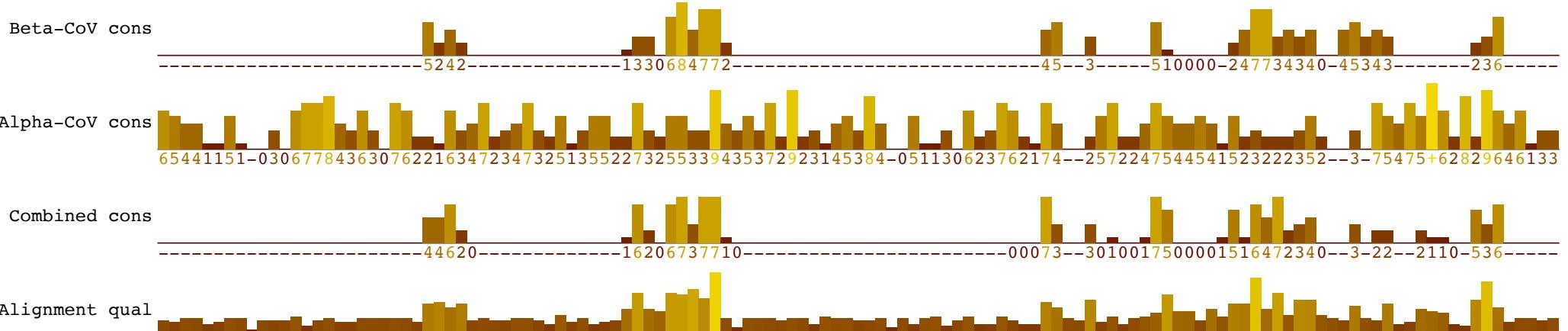
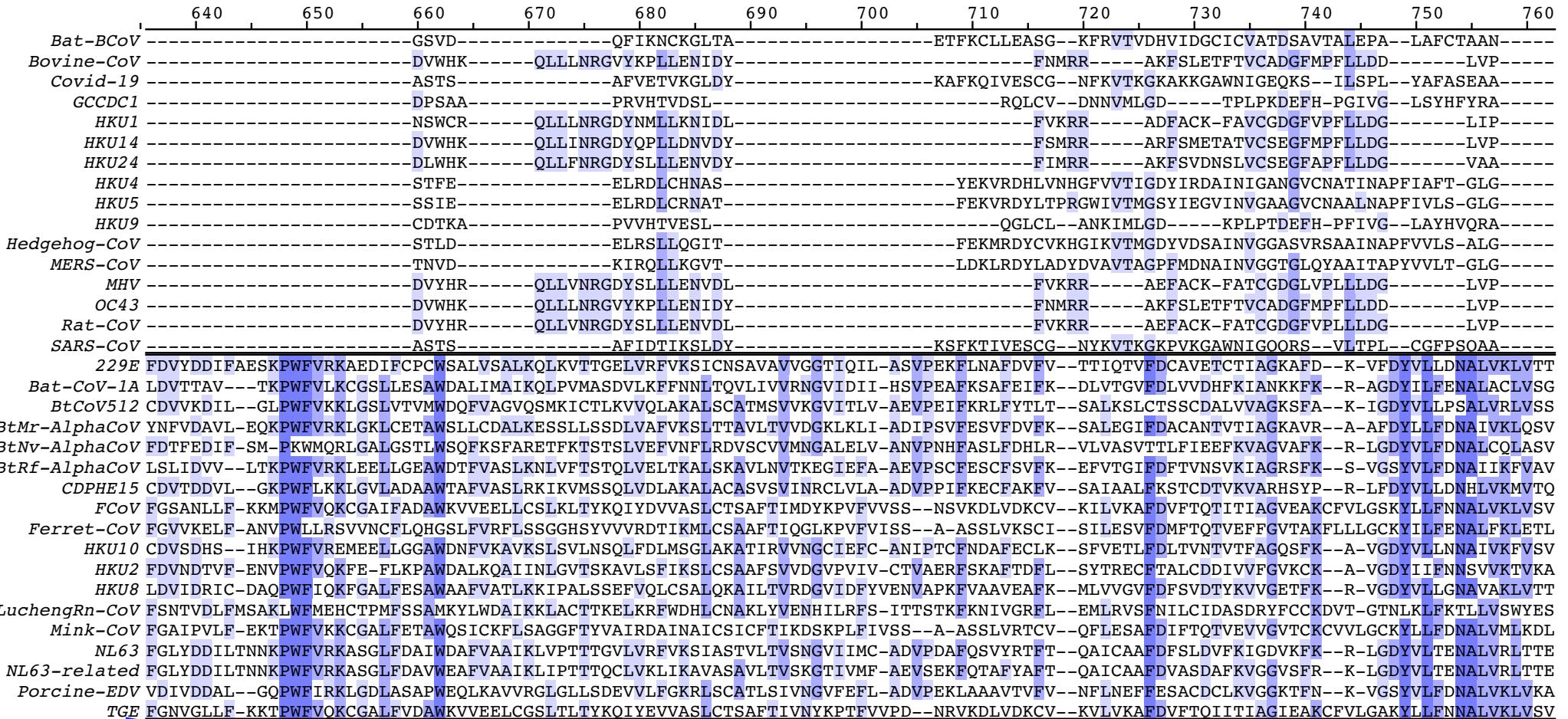






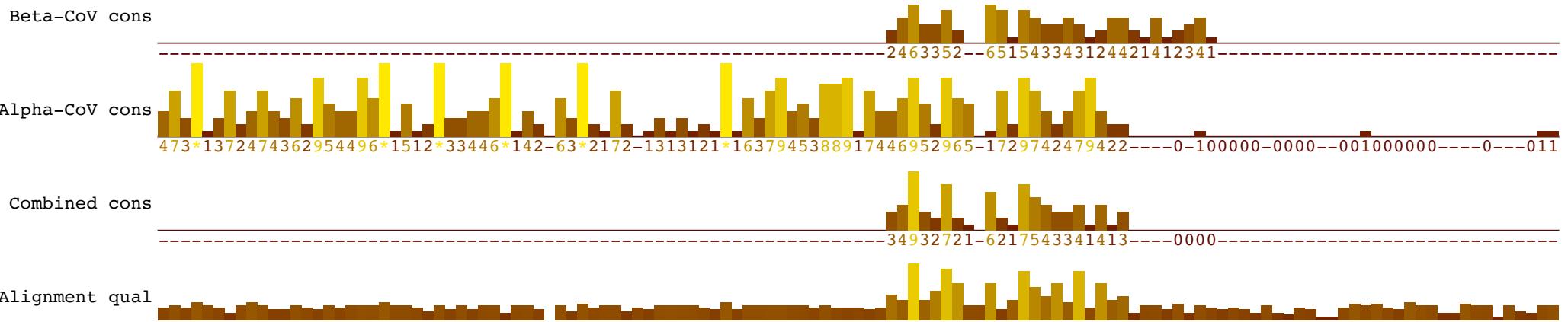


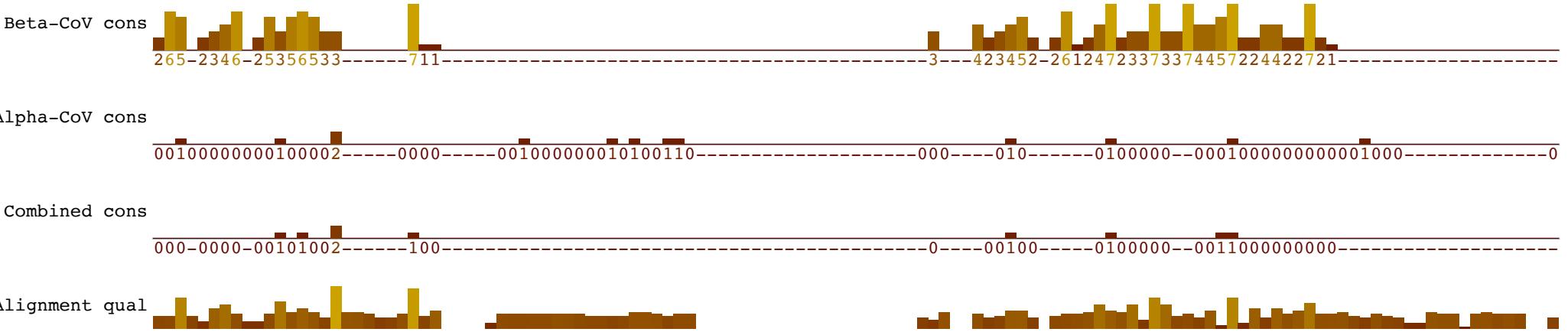
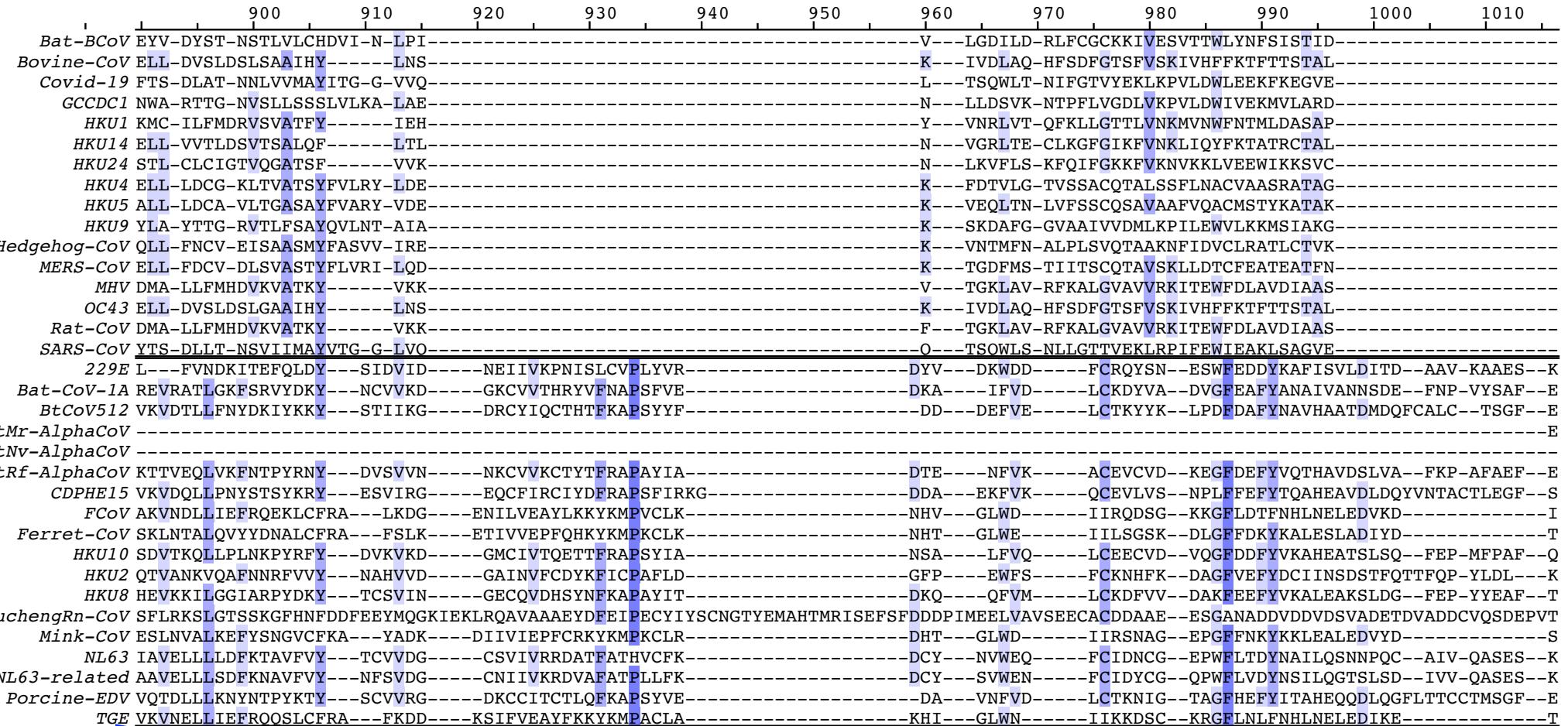


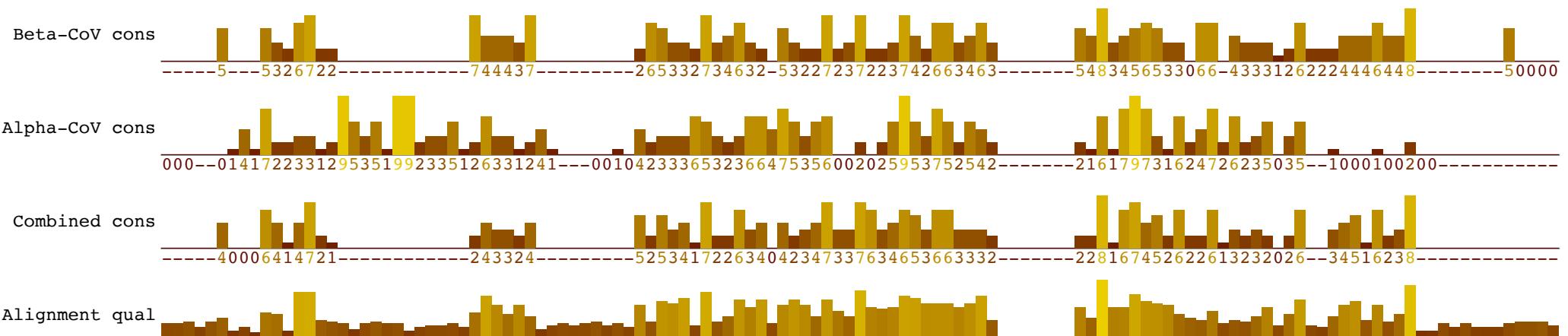
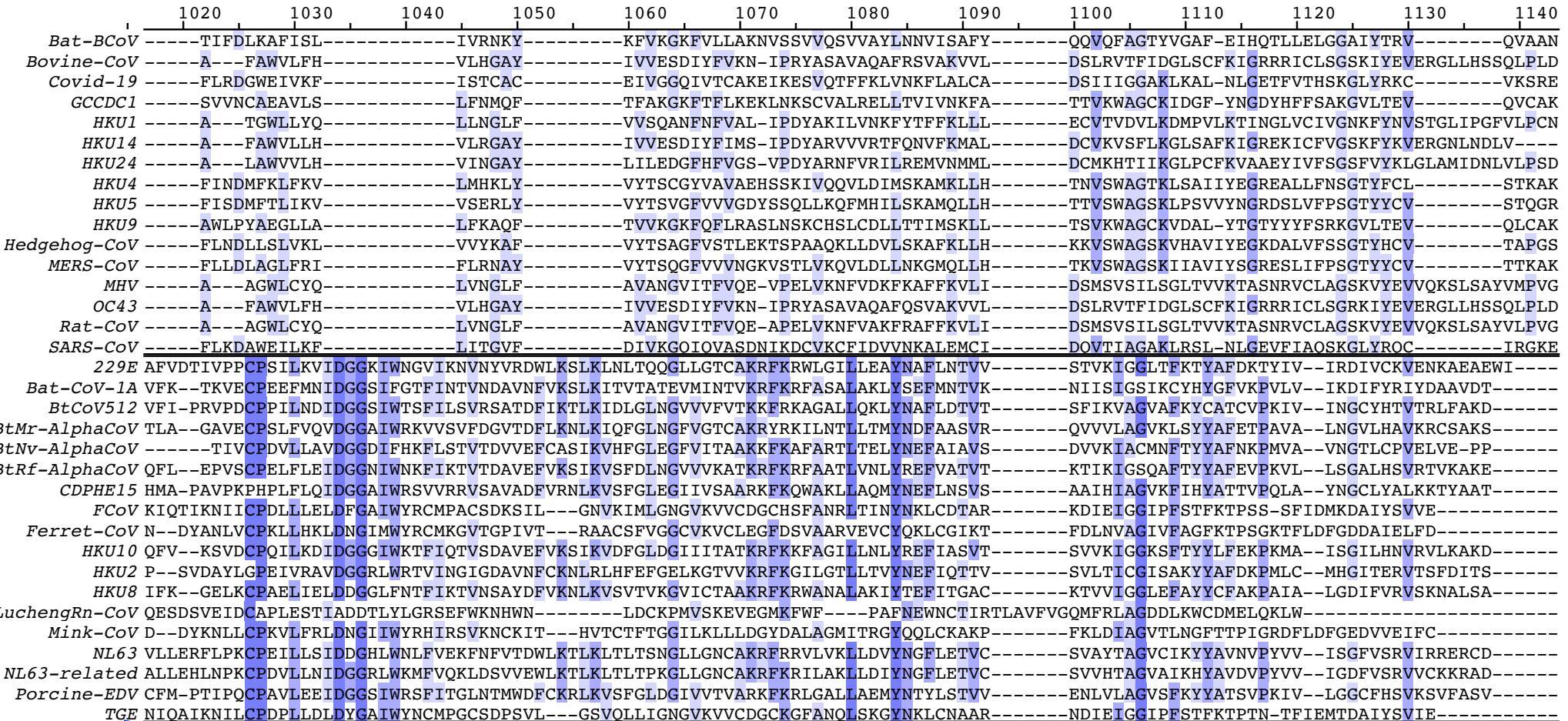


Bat-BCoV 770 - 880  
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 Covid-19  
 GCCDC1  
 HKU1  
 HKU14  
 HKU24  
 HKU4  
 HKU5  
 HKU9  
 Hedgehog-CoV  
 MERS-CoV  
 MHV  
 OC43  
 Rat-CoV  
 SARS-CoV  
 229E  
 Bat-CoV-1A  
 BtCoV512  
 BtMr-AlphaCoV  
 BtNv-AlphaCoV  
 BtRf-AlphaCoV  
 CDPHE15  
 FCoV  
 Ferret-CoV  
 HKU10  
 HKU2  
 HKU8  
 LuchengRn-CoV  
 Mink-CoV  
 NL63  
 NL63-related  
 Porcine-EDV  
 TGE

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 RAYYLAV--SGQAFCDYADKICHAVVSKSK--  
 RVVRSIFSRTLETAQNSVRVLQKAATILDGISQYSLRL--I--DAMM  
 CWYGVLTAKSFGAFKELLQSEEVRLSHFCANI--RRC--L--DRAL  
 RSYYLIQ--SGIFFTSLMSQFSQEVSDMCL--  
 RTYYLVK--SGQAFCDMLCEFGQEVADLSK--  
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 CWYGALASRTFEAFRDFVRTEERFAQFFGKV--CAP--I--NGCV  
 ESFKKIAAVPFKVGSSFLKTWEYLSDCIVYRVLPLYELE--D--VSDFV  
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 RSYYLIK--SGQAYTSMMVNFSSHEVIDMCM--  
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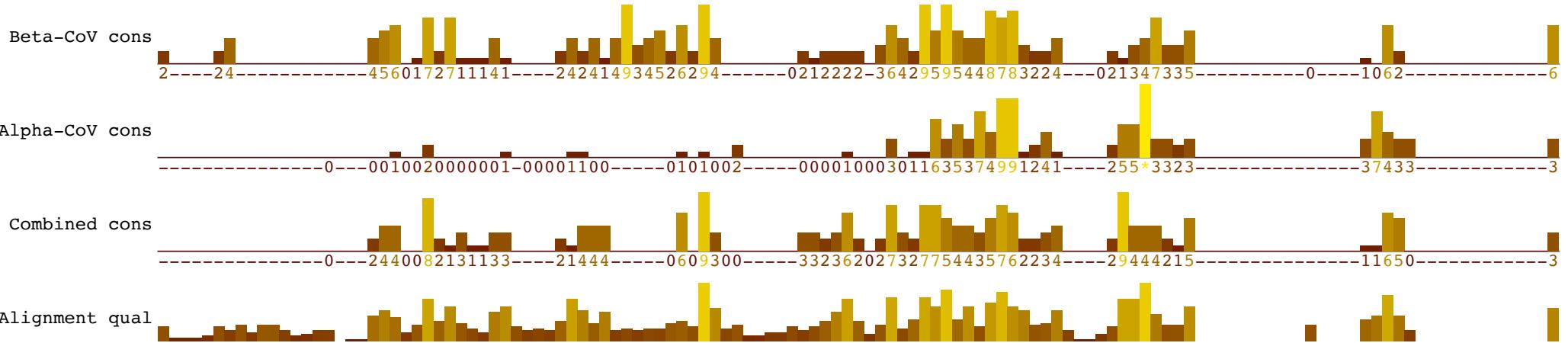


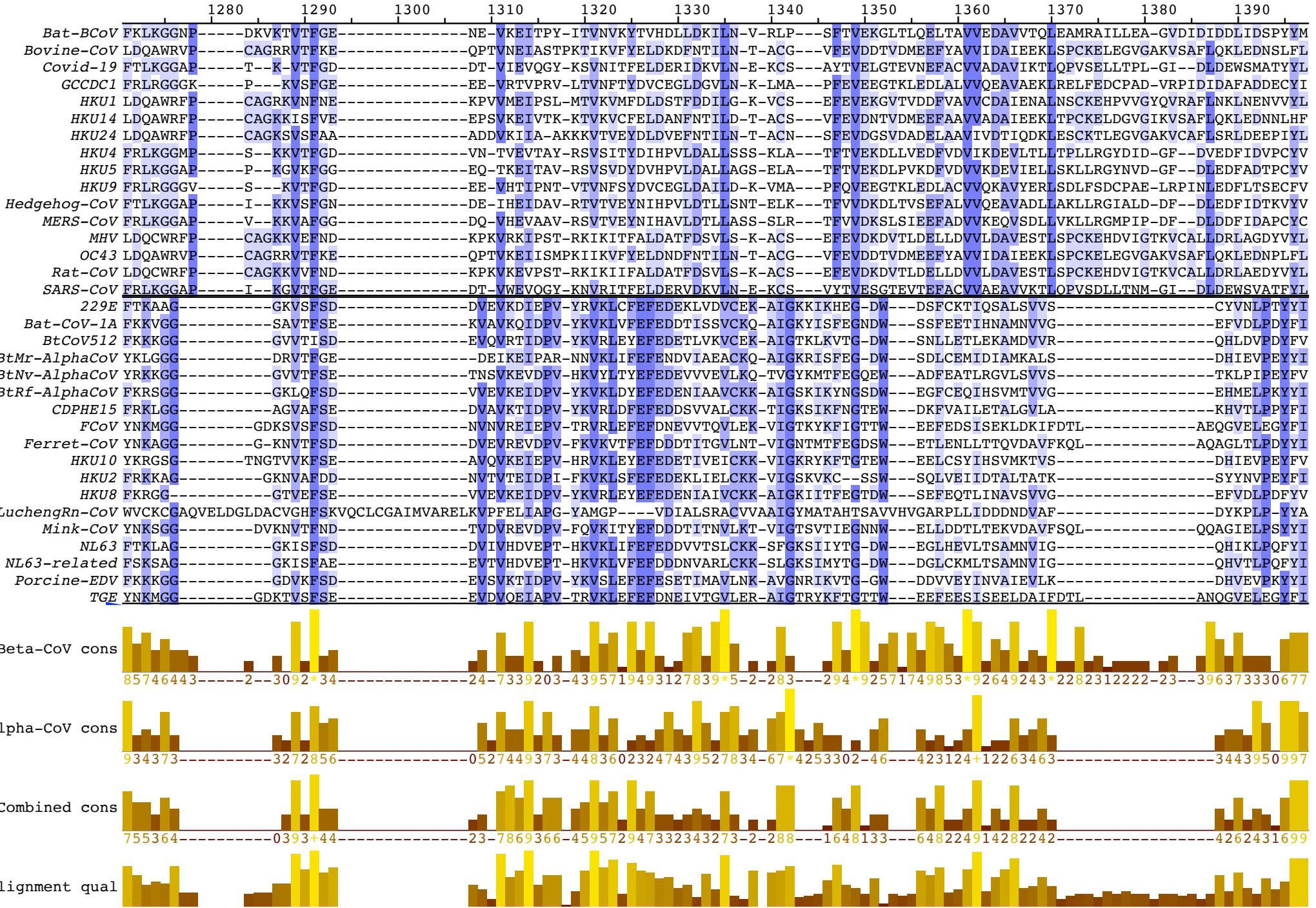


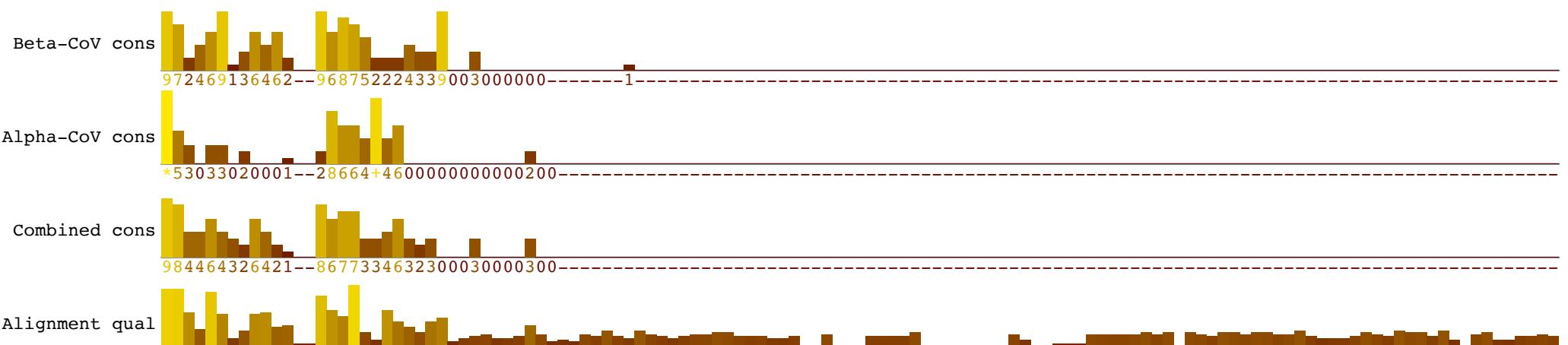
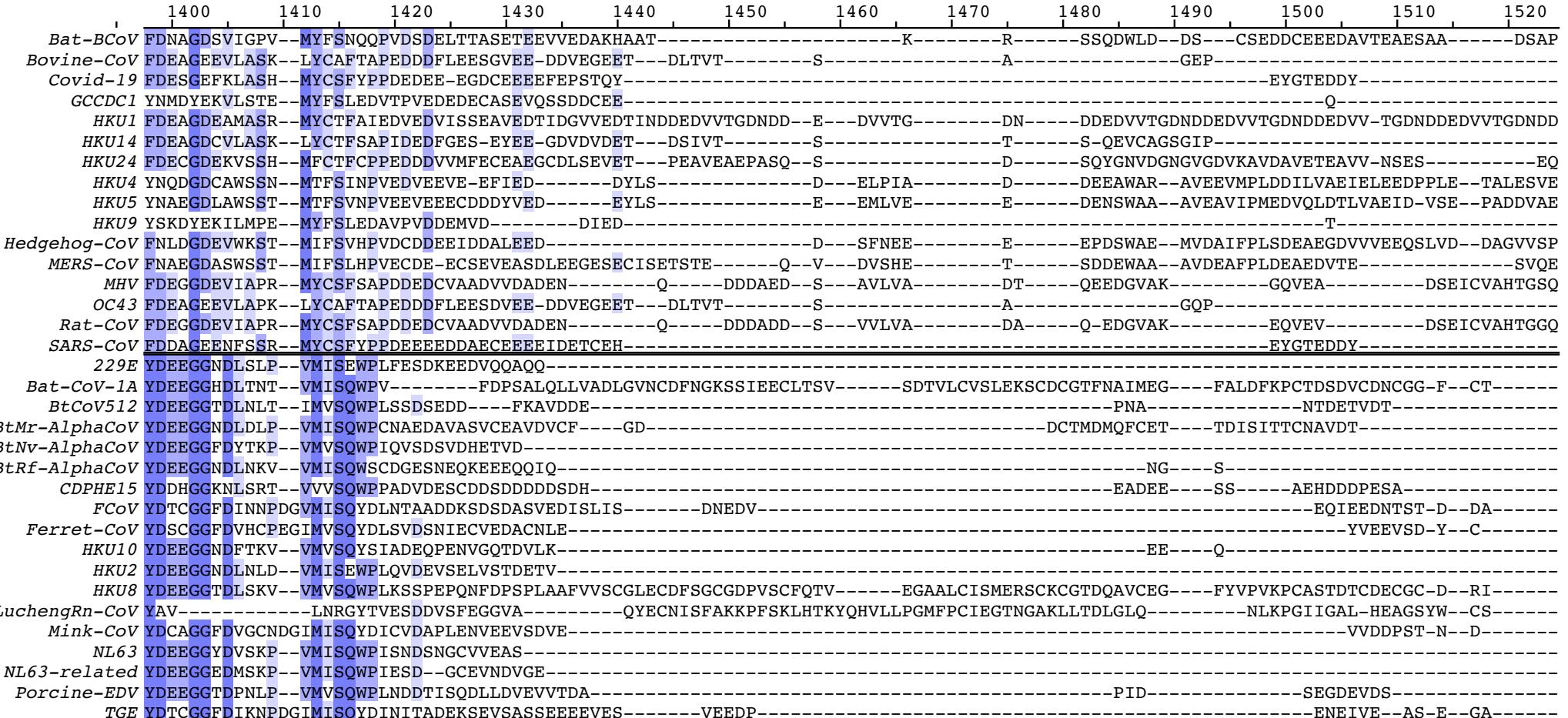


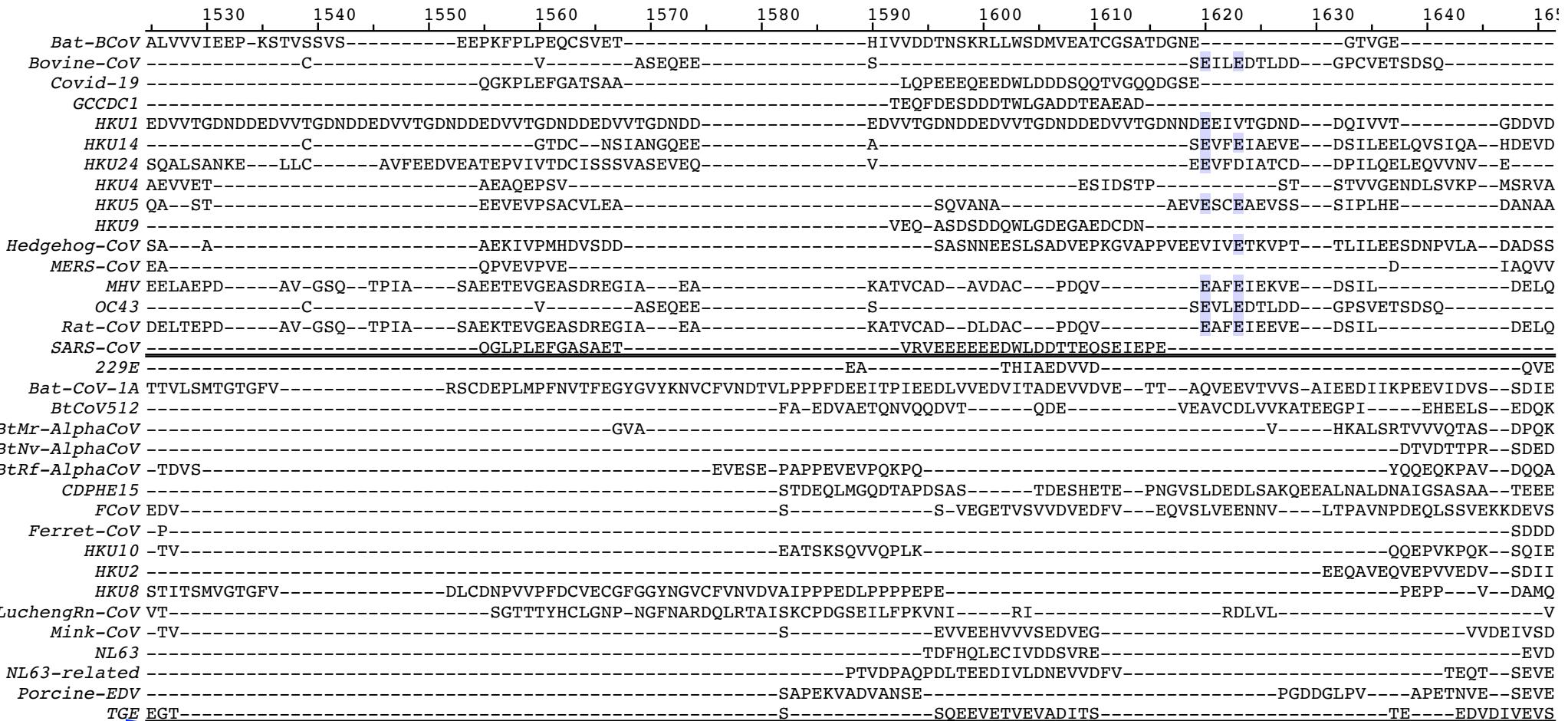
1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260

<i>Bat-BCoV</i>	S---IQDLVLPSTVSK---KELTFVDGHAI DT---ELTSEEVTLEGIV EQLDTDTN KIAGDSIKGDLTVNNCVFVRFGDG---NL-SPATTD	C---QATKG---V
<i>Bovine-CoV</i>	V---YDLTMPSQVQK-AKQKPIYLKGSGSDF---SLADSVVVTTSLTPC---GYSEPPK-VADKICIVDNVYMAKAG---DKYYPV VVD	GHVG---L
<i>Covid-19</i>	E---TGLLM-PLKAP---KEIIFLEGETLPT---EVLTTEEVV LKTGDLQPL QOPTSEAPEALVGT P CINGLM LLEIKDT---EKYCALAPN	MVTNN---T
<i>GCCDC1</i>	T---LGAMLTPRQR---MEVEVLDGKYDAP---VAITVPELEELNGTLEEVFG---FDDLT LVRGS LVALASKVFVRTDDG---LLFRYVKSG	G---V-FLK---A
<i>HKU1</i>	A---QE---QQIYFFEGVAESV---IVEDDVIENVKSSLSSY---EYCQPPK-SVEKICII DNM YM GKG C---DKFFPIVMN	D---KNIC---L
<i>HKU14</i>	R---RDILVVPSVTQRAKNQQPVYLTGNACPV---NVDDD VVVEV TNPV TSC---GYQKPPQ-KCDKICIVDNVYMAKCG---EKFFPV VVN	E---DYIG---L
<i>HKU24</i>	M---IQ---SELLAHGDVIPV---TLESEVVEVV KQPLKPL---GYRKPPM-KAARLCVVDNL YMAQNG---DCLYPVIFD	G---ORIG---L
<i>HKU4</i>	TLQGQMLVLPGDYNK---KTLGILD PVPNA D T D V N A N S T V V D V V H G Q L E ---PTNEHGP SMIV G N Y V L V S D K L F V R T E D E ---EFYPLCTN	G---KVVST---L
<i>HKU5</i>	SLQDQFDL V I P G D L S K ---KQIGI LEPTPN S T T V D K K I N T N V V E V V V G Q L E ---PTKEHSPELVVGDYVIISNK I F V R S V E D S E T V F Y P L C T D	G---KIVPT---L
<i>HKU9</i>	R---LGLLLTPKQ QK---MEVEVLDGDFDAP---VTLTDLE ECTGV LEEVFG---ASDV KLVKG TL VSL ASK L F V R T E D G ---FLYRYVKSG	G---V-LGK---A
<i>Hedgehog-CoV</i>	V V G A H L E A T I P G E V V K ---KQLSM I T A T N Y S T T V D V R P K T R N V E L V Y G Q L E ---TTNMHS PD V V V G D Y V I V S D K V F V R S E E D G R L A F Y P M C T N	G---KAVPC---M
<i>MERS-CoV</i>	SVQQDLDVILPGEFSK---KQLGLLQPTDNSTT V S V T V S S N M V E T V V G Q L E ---QTNMHS PD V I V G D Y V I I S E K L F V R S K E E D G F A F Y P A C T N	G---HAVPT---L
<i>MHV</i>	C---SE---ATC---LVGEIEPA---VFEDDV V DV KAPL TYQ---GCCKPPT-SFEKICIVDKLYMAKCG---DQFYPV VVD	N---DTVG---V
<i>OC43</i>	V---YDLTMPSQVQK-AKQKPIYLKGSGSDF---SLADSVVVTTSLTPC---GYSEPPK-VAAKICIVDNVYMAKAG---DKYYPV VVD	DHVG---L
<i>Rat-CoV</i>	C---SE---ATC---LVGESEPA---VFEDDV V DV KAPL TYQ---GCCKPPT-SFEKICIVDKLYMAKCG---DQFYPV VVD	N---DTVG---V
<i>SARS-CoV</i>	O---LOLLM-PLKAP---KEVTFLEGDSHDT---VLTSEEV L K N G E L E A L E T P V D S F T N G A I V G T P V C V N G L M L L E I K D K ---EOYCALSPG	L---LATNN---V
	229E---ELFPHNDR I---KSFS T F E S A Y M P I A D P T H F D I E ---EVELLDA---EFV EP G C G G I L A V I D E H V F Y K K D G ---VYYPSNGT	NILPV---A
<i>Bat-CoV-1A</i>	FNVAVEAGL---NTVKTFSGGENP I-TFSRVEVA---SVELEHA---EYV K P E A N G H V S V I N G H T F Y T C G D ---Y Y Y P C D Q N	NCFSQ---C
<i>BtCoV512</i>	LQIPTEDGV---ADFNTFHCVFPV-NPTRIETD---SLEEEV---DFV EP G V D G K L V I L D D Y S F Y S D G T ---N Y Y P S D G K	G V V A S ---C
<i>BtMr-AlphaCoV</i>	LDVPCCEGA---AEFN V F E G V H L L -KPTA VELA---SDMELLET---D Y V E P C A G G V L R I C D D Y A W Y M D G D ---R F Y P Y G D N	G V I P L ---C
<i>BtNv-AlphaCoV</i>	SDVSKEALV---TDFMVFKSPDTHV-KPIRIETE---SCELHEV---DFV T P R D G G V V K I I D G Y A F Y E S N G ---N Y Y P S S V D	G V K N V ---T
<i>BtRf-AlphaCoV</i>	LNLSVQDV V---V E F N V F D G A T V E V -T P K R V E K E ---EVS LEEV---E F I P P V S N G S I A V V D G Y S F Y S C G D ---L Y Y P S N E N	AVVPL---C
<i>CDPHE15</i>	VGLQTEEGV---DGFYAFDH C Q L P V -T P R L L Q V E ---D V N L E E A ---D F K P P V G G G C V A V I N G Y S F Y V E G D ---S Y Y P T D C S	A I H S L ---R
<i>FCoV</i>	Y---GEALSF K T A S V P V T N S G I I T T D D W -S D P I L L E P A ---D Y V E P K D N G D V I V I A G Y T F Y K D E D ---D H F Y P Y G S G	M V V Q K ---M
<i>Ferret-CoV</i>	D---EPVRCFTGATV P V N V S D F E D D V R -G D V V -L E E C ---E Y Q P P R D E G I V V N L G G Y T F Y T T D N ---E K F Y P F G D C	K V V P K ---L
<i>HKU10</i>	CTIAAEDIL---VEMTVFDSF E I P I -QPSRVEVQ---Q V E F E E A ---E F V Q P Q A G G V L A V V N K H V F Y N K G D ---Y Y Y P A N E T	H A V P V ---C
<i>HKU2</i>	LGK P I L D G L ---S R C N V F A R G S N A V -V V D S V D H E ---L L D L E E C ---D F V E P A S H G V L V V S D D Y G F Y S C E T ---G I Y P Y A S D	G---K V L P L ---R
<i>HKU8</i>	FNR P T E A G A ---V D A L V F E G G D I E T -V C S R V Q F D ---T V A L E P V ---Q E V P V P S D G Y V A V V D G Y T F F T D G T ---H Y F P T G L S	R C Y T T ---C
<i>LuchengRn-CoV</i>	-----ETKSQ Y L W Y K L A V ---A A E P D K V E R G N E G S Y S G N G M D Y N T V F R V A L R K L K L S N A K A C	
<i>Mink-CoV</i>	-G---GKTRCFSGASIPV S V E K I V E D D V W -S T T I -L E E T ---E Y Q P P K N G G V V V V I D G Y T F Y T C N H ---E Q F Y P F G D S	K V V Q K ---L
<i>NL63</i>	MTFPCVSCV---T F F Y E F L D T C F G V S K P N A I D V E ---H L E L K E T ---V F V E P K D G G Q F F V S G D Y L W Y V V D D ---I Y Y P A S C N	G V L P V ---A
<i>NL63-related</i>	VPVPCDSRI---T S F H E F V D A V Y K V G K P T T I D V E ---H L E L K E T ---S F V E P K E G G Q F V V S D D Y F W Y I M D D ---I C Y P A S E D	G V L P V ---A
<i>Porcine-EDV</i>	FQIPVQAGI---EKF KVFLNCVHPV-VPRVIETS---F V E L E E T ---T F K P P A L N G G I A I V D G F A F Y D G T ---L Y Y P T D G N	S V P I ---C
<i>TGE</i>	O---GKALSFRDADPV V D N G T I S T A D W -S E P I L L E P A ---E Y V K P K N N G N V I V I A G Y T F Y K D E D ---E H F Y P Y G F G	K I V O R ---M





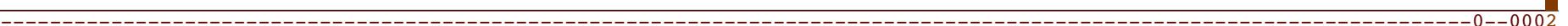




Beta-CoV cons



Alpha-CoV cons

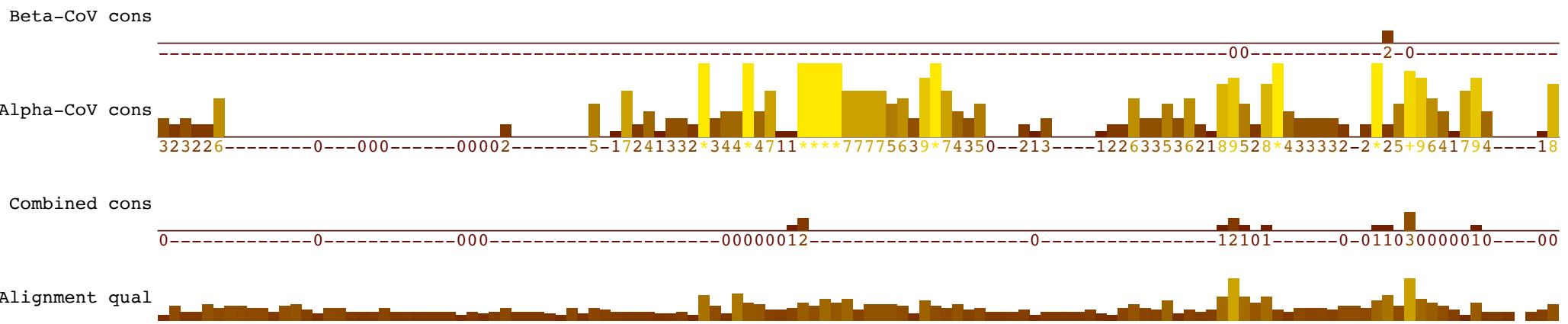


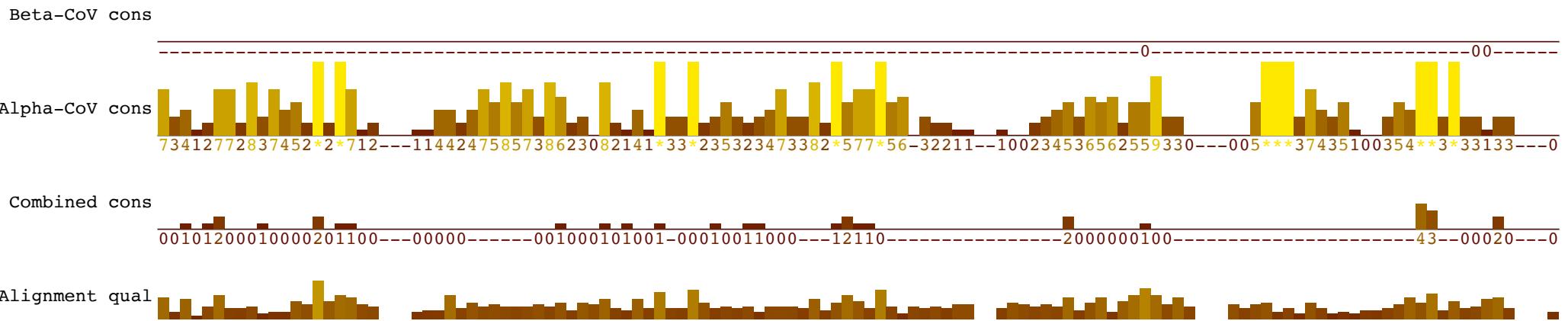
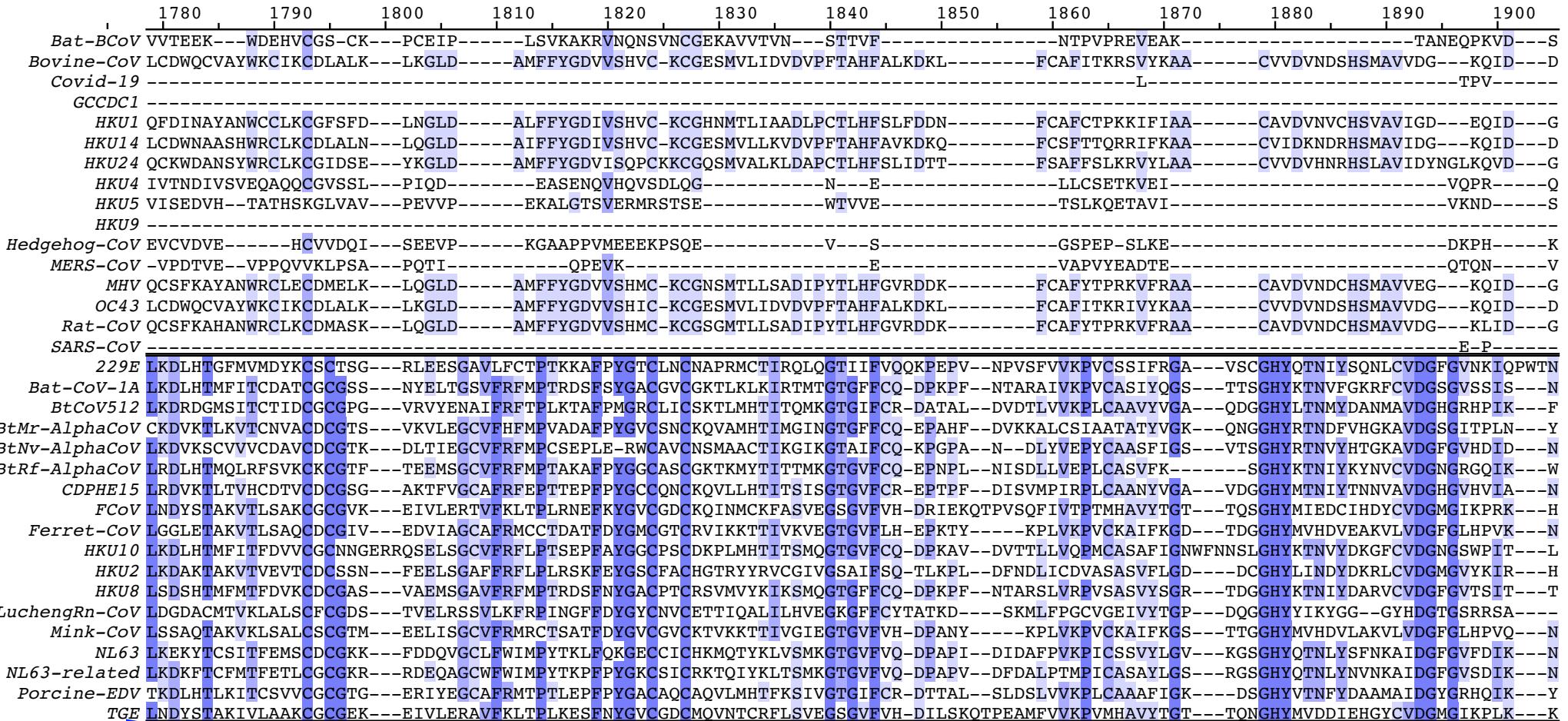
Combined cons

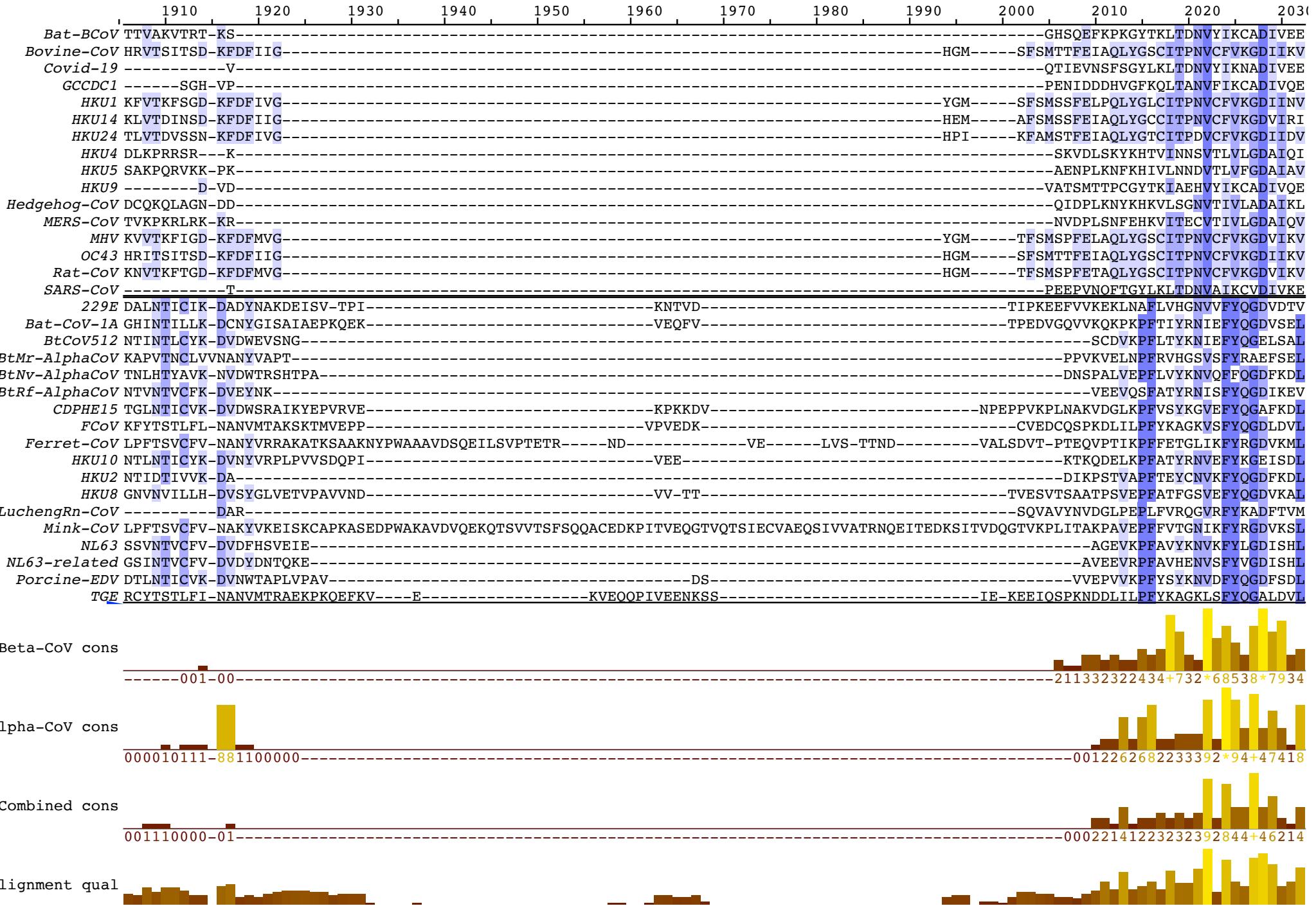


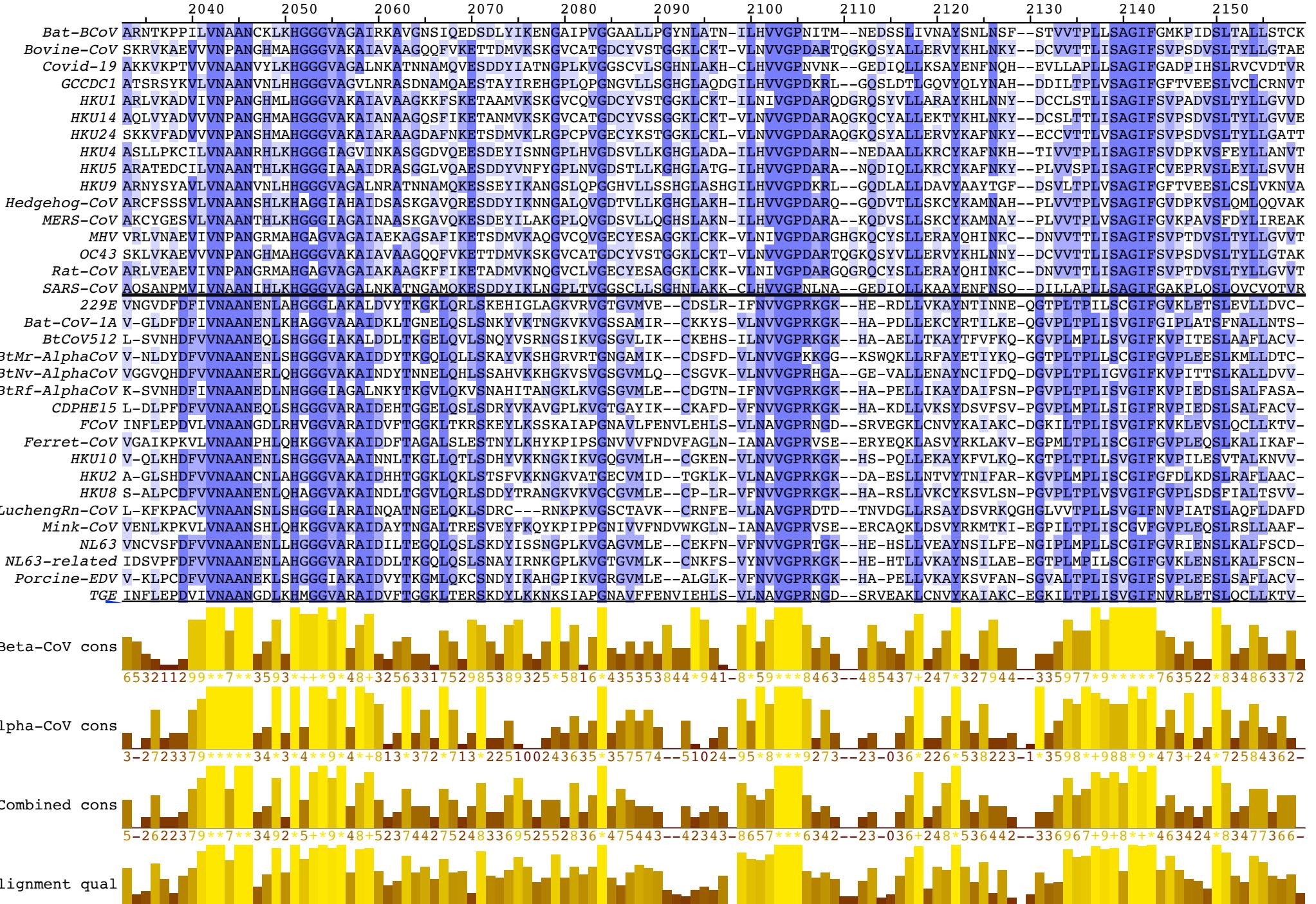
Alignment qual

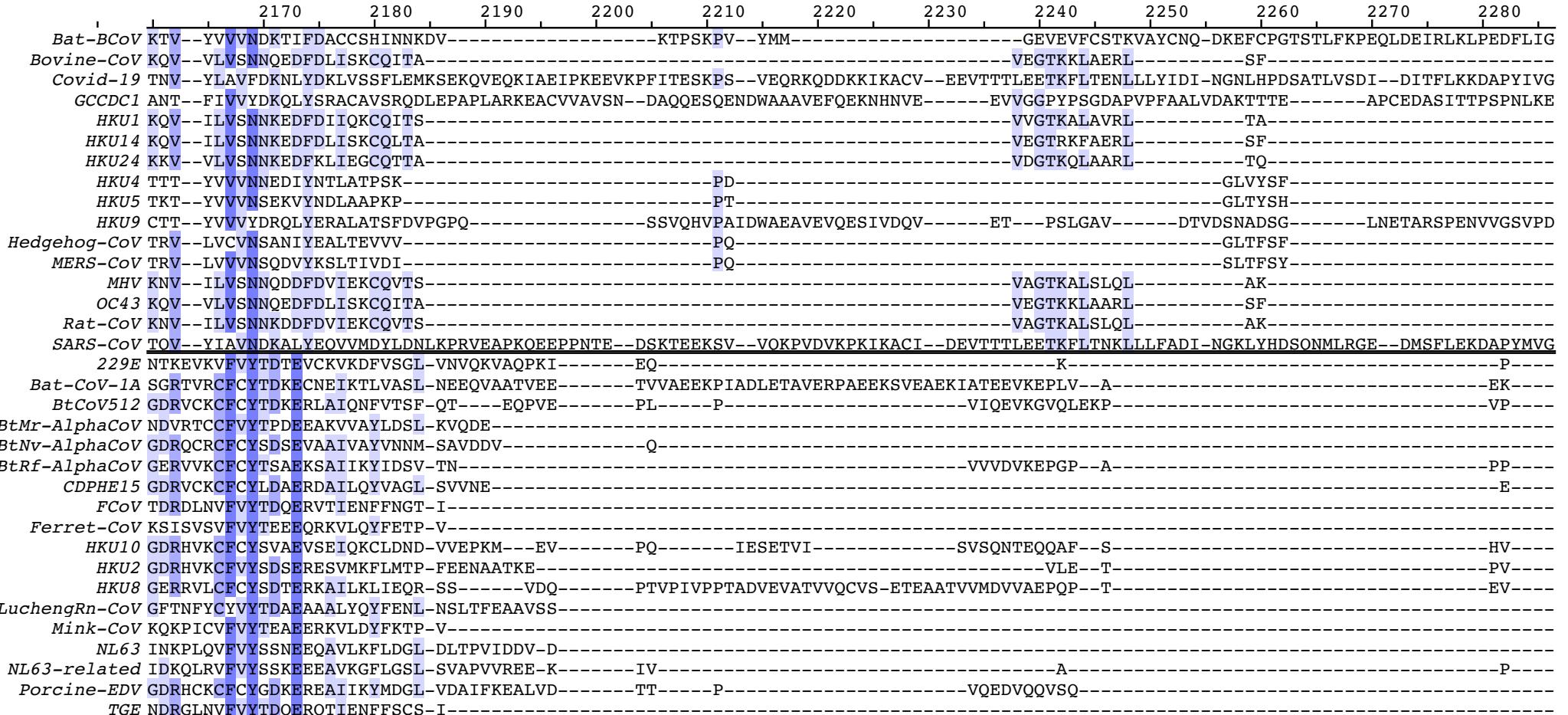


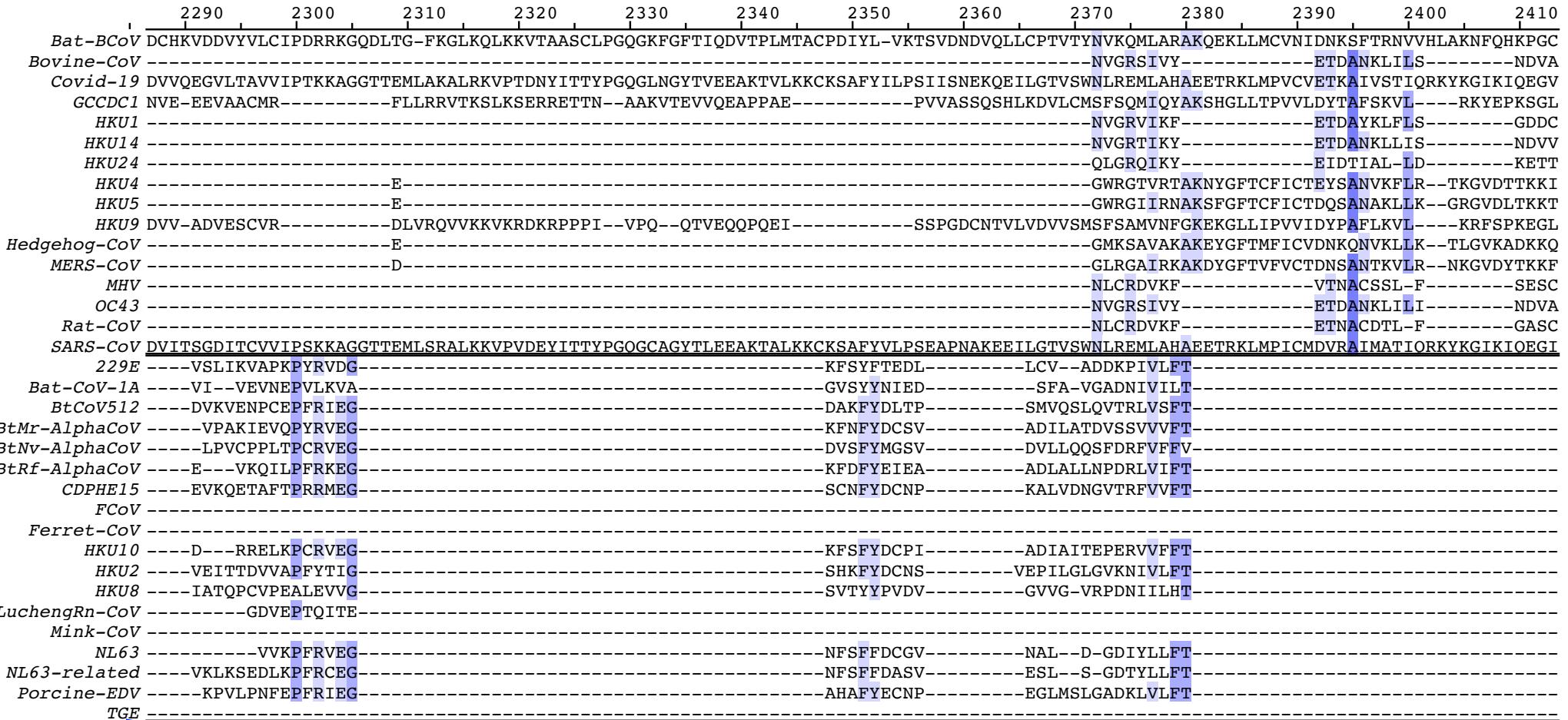


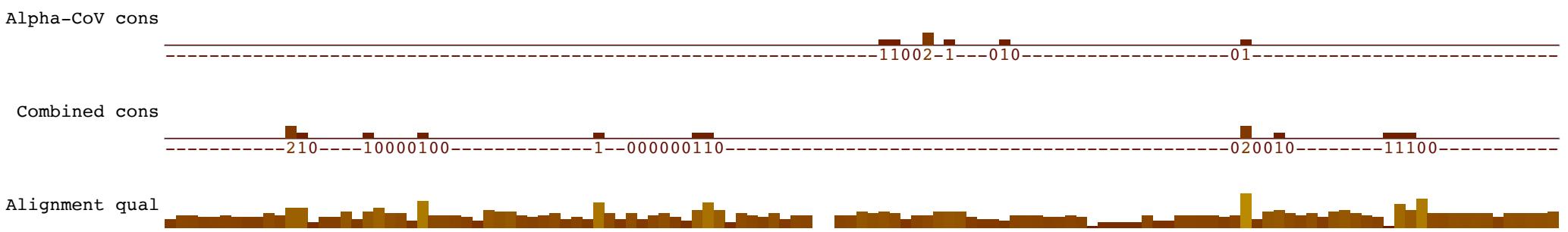
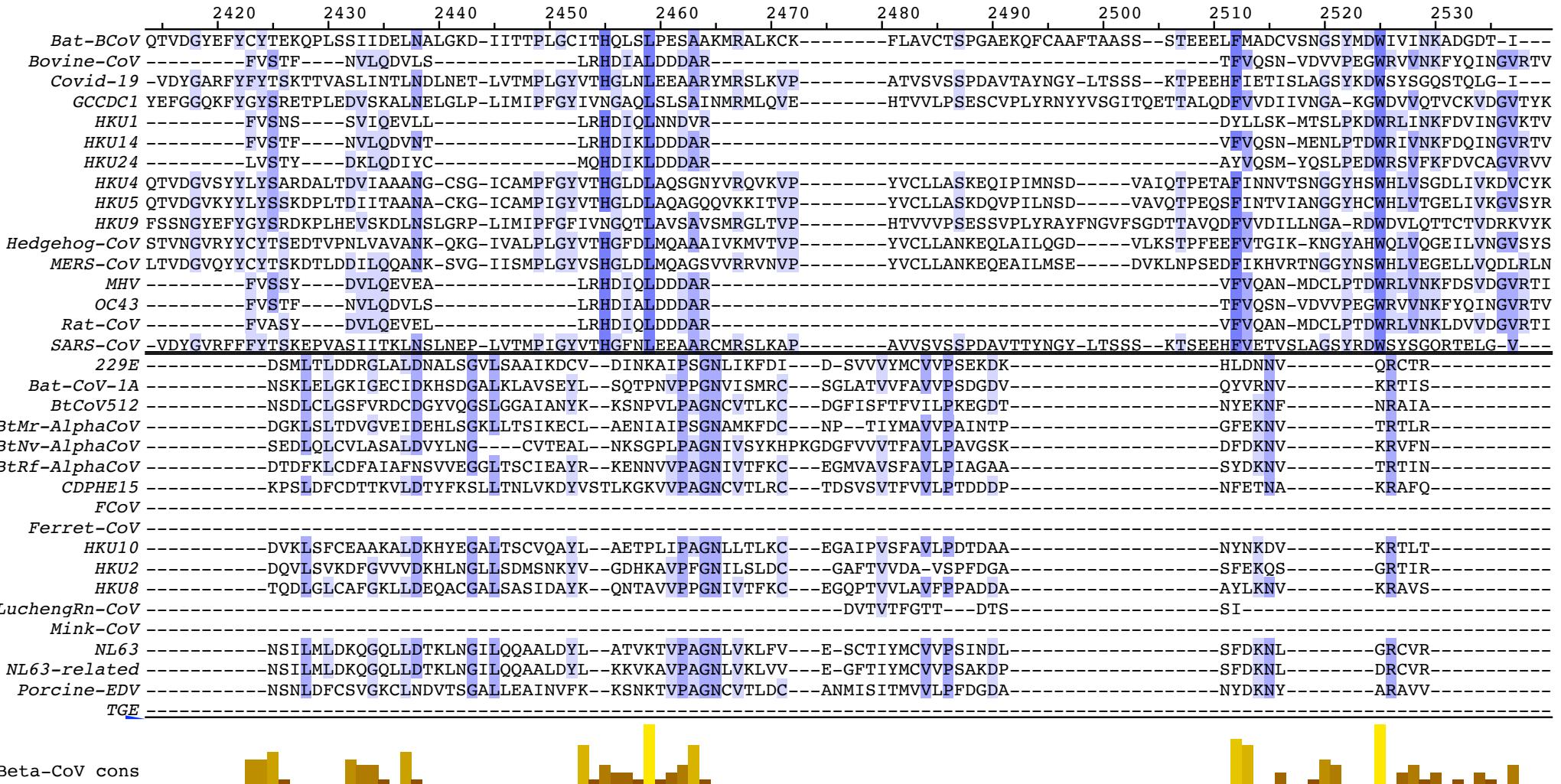


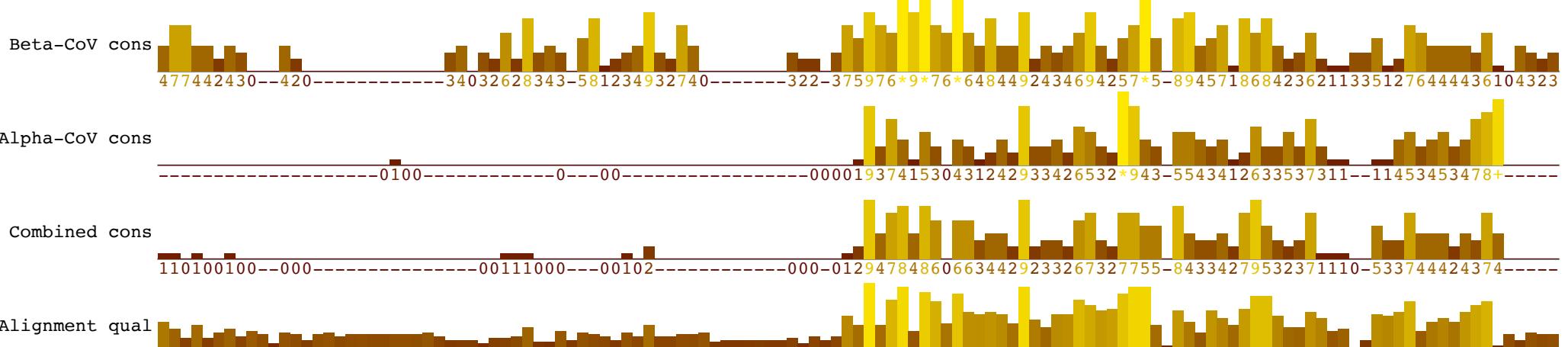
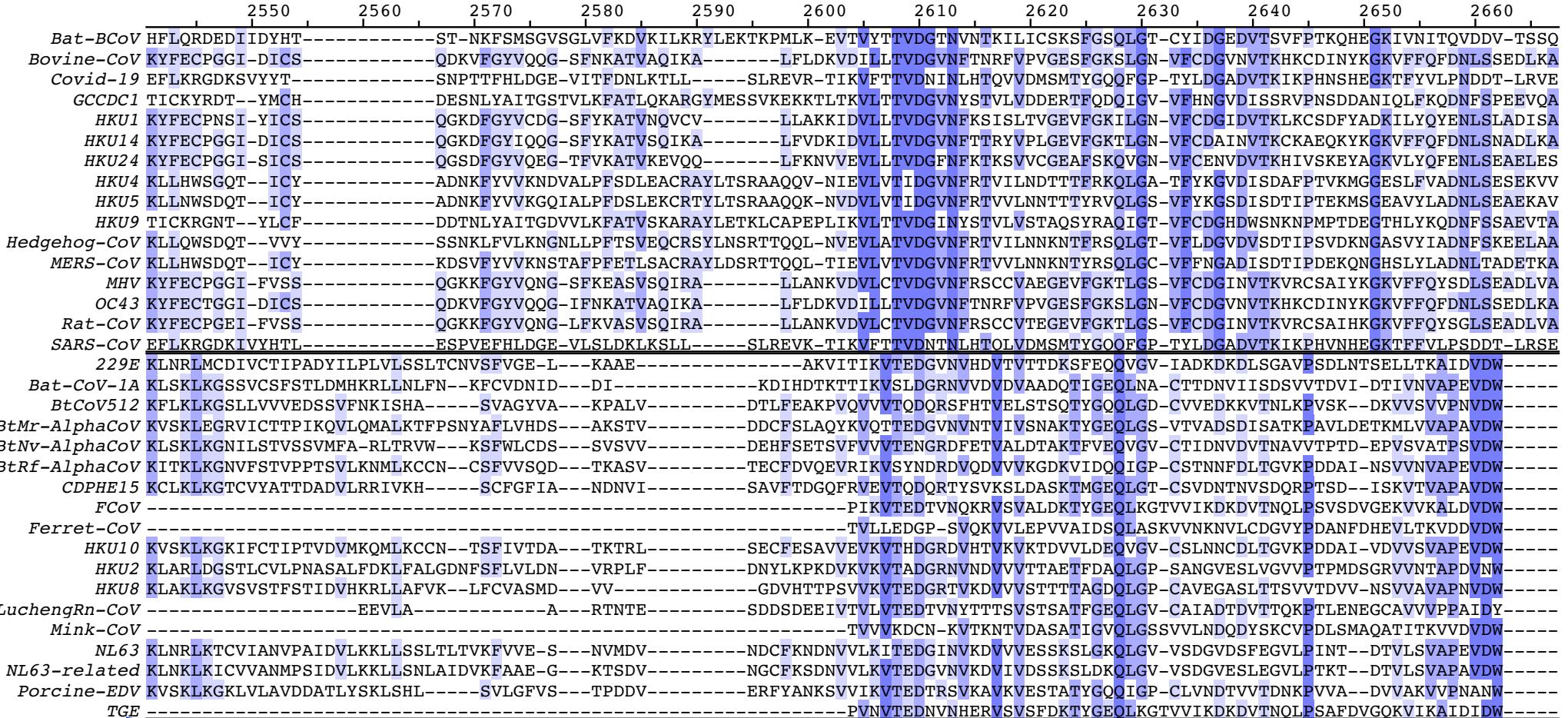












2670 2680 2690 2700 2710 2720 2730 2740 2750 2760 2770 2780 2790

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*Bovine-CoV* VRSSFNFQDQKELL-AYYNMLVNCSKWQVVFN--GKYFTFKQQANNCFVNVSCLM1QSL-NLKFKIVQWQEAWLEFRSGRPARFVSLVLAKGFFKFGDPAASRDFLRRVVFQSVDLT-GAICDFE--I

*Covid-19* AFEYYHHTDPSFLGRYMSALNHTKKWKPQV--NGLTS1KWAQNCCYLATALTLTQOI-ELKFNPNALQDYYRARAGEAANFCALILAYCNKTVGELCDVRETMSYLFQHANLD-SCKRVLN--V

*GCCDC1* ISDYYGVNDCTVVARALSVRRYVRTWPYSVV--DGRVLLAQRDNSCYLNAA1SVLQDI-EVEFSTPWISTAFDALKGGNPLPIVEVLITLGKTTIGEPDDAHMILSSLSSYASVK-AARRAIT--T

*HKU1* VQSSFGFDQQQLL-AYYNFL-TVCKWSVVNN--GPFFSFEQSHNNCYVNVA1CLMLQHI-NLKFKWKWQWQEAWYEFRAGRPHRLVALVLAKGHFKFDEPSDATDFIRVVLKQADLS-GAICELE--L

*HKU14* VKSSFNFQDQKELL-AYYNVLVSCGKQIVVN--GKYFTFKQQANNCFVNNA1CLMLQNV-NLKFKVSMQWQEAWLEFRAGRPLRFVALVYAKGAFKFGDPADSRDFIRVVLQTDLA-EAACDYE--F

*HKU24* VKQSFQFDQKQLL-AYYSY-RSCCRWEVVH--RKWFCKQADNNCYCNAACLMLQNL-NLTFKLQWQWQEAWLEFRAGRPLRFVALVLAKGAFVYQGPADARDFMRVVALSQADLS-SASCEYE--F

*HKU4* LKEYYGTSDVTFLQRYYSLQPLVQQWKFVWH--DGVKSLKLSNYNCYINATIMMIDMLHD1KVFVVPALQNAYLRYKGGDPYDFLALIMAYGDCTFDNPDEAKLLHTLLAKAELTVSAKMWR--E

*HKU5* LSEVYGTADTAFLHRYYSLLALVKKWKTWH--DGVKSLKLSNNCYVNVTMILMDMLKE1KFIVPALQAAYLKHGGDSTEFIALIMAYGDCTYGEPPDASRLLHTILSKAELTTQAKMVWR--Q

*HKU9* IREYYGVDDSNIARAMSIRKTQWTWYPTV--DGRVLLAQRDNSCYLNVA1SLLQDI-DVSFSTPWVCRAYDALKGNNPLPMAEVLIALGKATPGVSDDAHMVLSAVLNHGT--TARRVMQ--T

*Hedgehog-CoV* VKEMYGVEDPTFLYKYYISRAKVVWKWMAMC--EDAPSLCLNSNNCYLNAAVMM1DCLRD1KFN1PALQAAYMFKGGDFGDFISLLMAYGNCTYQGPDDASMLHTALSKAELLVSAWMWR--E

*MERS-CoV* LKELYGPVDPDFTLHRFYSLKAAVHGWMVVC--DKVRLKLSDNNCYLNAAVIMTL DLLKDIKFV1PALQAHFMKHKGGDSTDFIALIMAYGNCTFGAPDDASRLLHTVLAKAECCSARMWV--E

*MHV* VKDAFGFDEPQLL-KYYTML-GMCWPVVVC--GNYFAFKQSNNNCYINVACLMLQHL-SLKFKPKWQWQEAWNEFRSGKPLRFVSLVLAKGGSFKFNEPSDSIDFMRVVLREADLS-GATCNLE--F

*OC43* VRSSFNFQDQKELL-AYYNMLVNCFKWQVVNN--GKYFTFKQQANNCFVNVSCLM1QSL-HLTFKIVQWQEAWLEFRSGRPARFVVALVLAKGFFKFGDPADSRDFLRRVVFQSVDLT-GAICDFE--I

*Rat-CoV* VKDAFGFDEPQLL-KYYTML-GMCWPVVVC--GNYFAFKQSNNNCYINVACLMLQHL-NLKFKPKWQWQEAWNEFRSGKPLRFVSLVLAKGGSFKFNEPSDSIDFIRVVLREADLS-GATCDLE--F

*SARS-CoV* AFEYYHHTDPSFLGRYMSALNHTKKWKPQV--GGLTS1KWAQNCCYLSSVLALQOL-EVKFNAPALOEAYYRARAGDAANFCALILAYSNKTVGELCDVRETMTHLLOHANLE-SAKRVLN--V

229E-VEFYGFKDAVTFAT--VDHSAFAYESAVV--NG1RVLKTSDDNNCWVNAVC1ALQYL-KPHF1SQGLDAAWNKFV1LGDV1FVAFVYYVARLVKGDKGDAEDTLNKLSKYLVNEAQVQLEHY--S

*Bat-CoV-1A*-DSFYGFPHAAEFGHM--LDHSAF1AFDNDVV--DGKRALVGTDNNCW1AVCLQ1QFA-EVDFTESEG1KDMWNEFLVGNVAKFC1HWIYLWLRANKGDAGDAENALNMLNKVKAHTVTLTRE--T

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*BtNv-AlphaCoV*-DAFYGFENASVFT--LDHSGFEFESTVV--EGKRTLKSSDNNCWVNAVC1Q1QYL-DAKFVSEGLN1WDSYVTGNVSKFVHWLYWITDHSKNDRNDAESVLLRLSVYLCKSCVFTIERE--T

*BtRf-AlphaCoV*-ISHYGFPRADVWHT--LDHSAYDYESKEQ--DGVRVLKFSDDNNCWVNA1C1Q1QIA-K1TRTEGIQAMWDEF1VGKVA1FVHWLYWYQNSSKGDPGD1VTLNA1SRYMANKSTV1VISRV--A

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*FCoV*-NAYYGF1PNAAAFSA--SSH1DAYEFDVVTH--NNF1VHKQTDNNCWVNA1C1Q1QRL-KPTWKF1PGV1SKLWD1FLTRKTAGFVHWLYH1SGLTKQPGDAELTLHKLVDLMSDSAVTVHT--T

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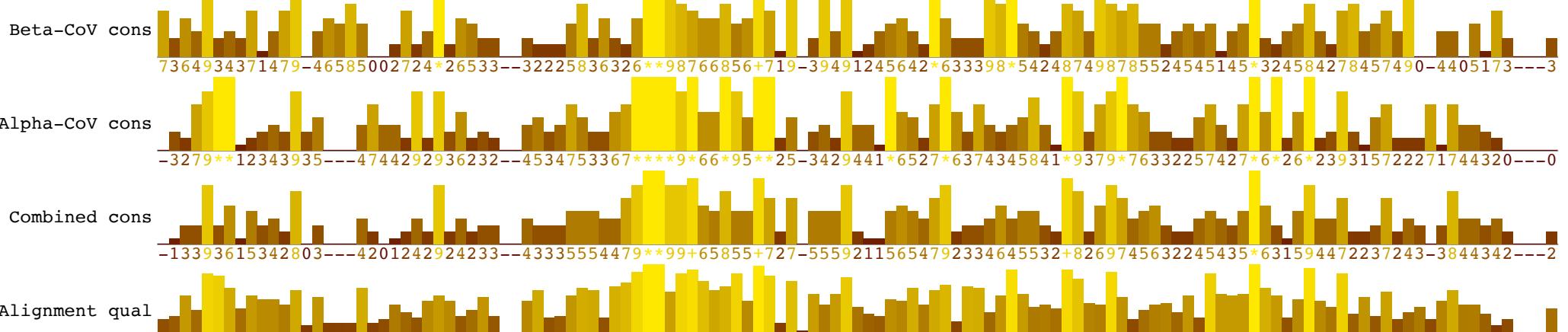
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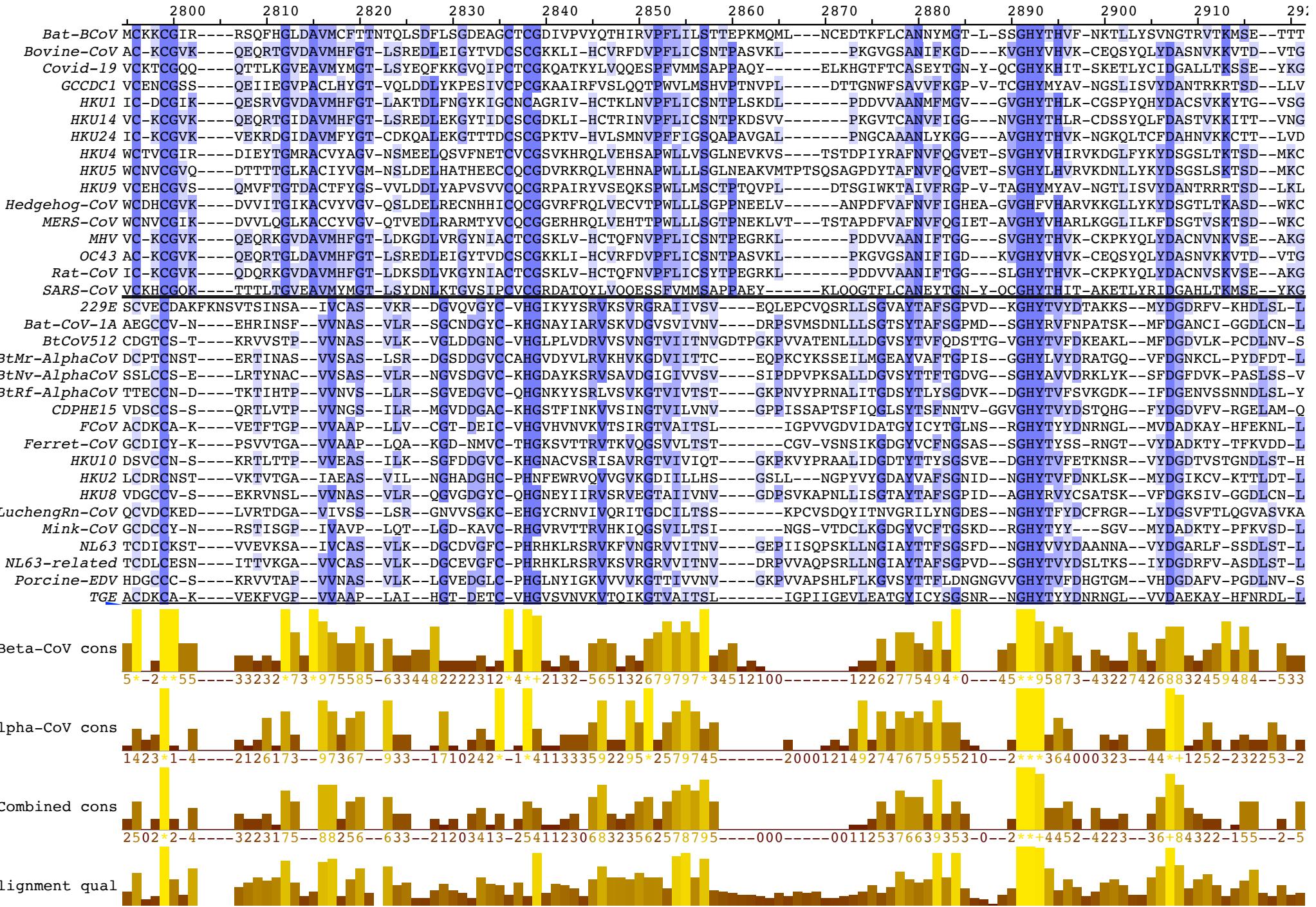
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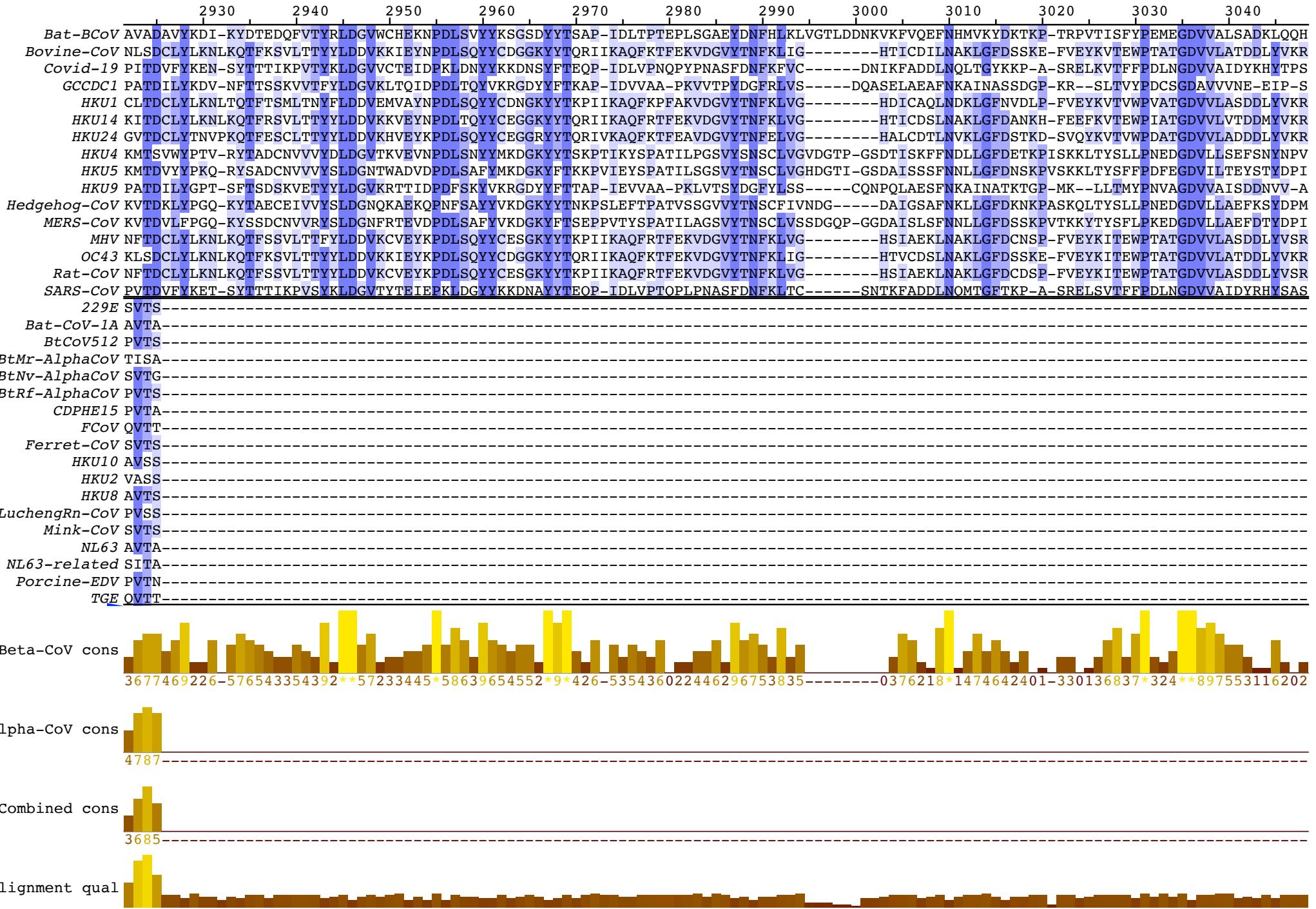
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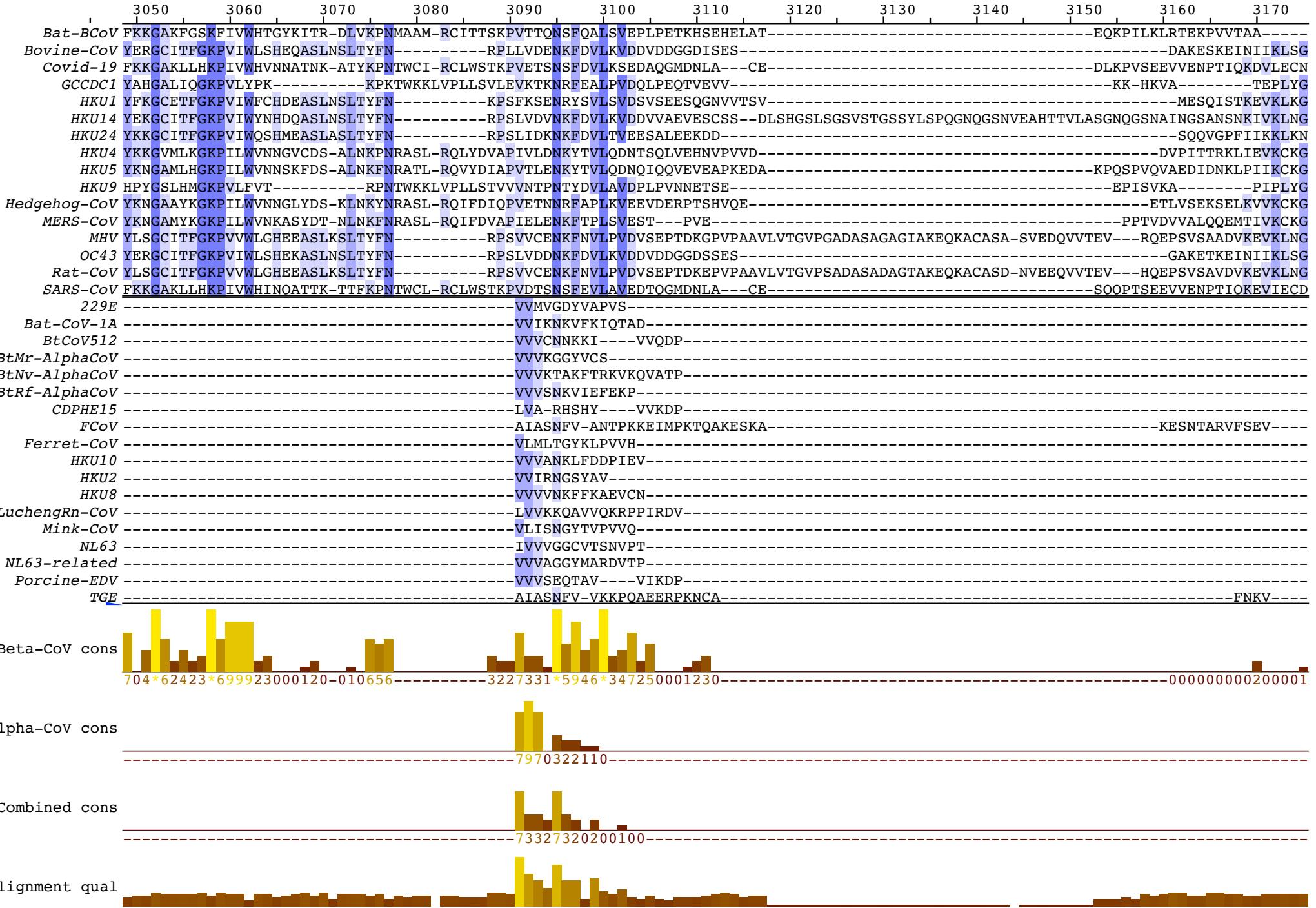
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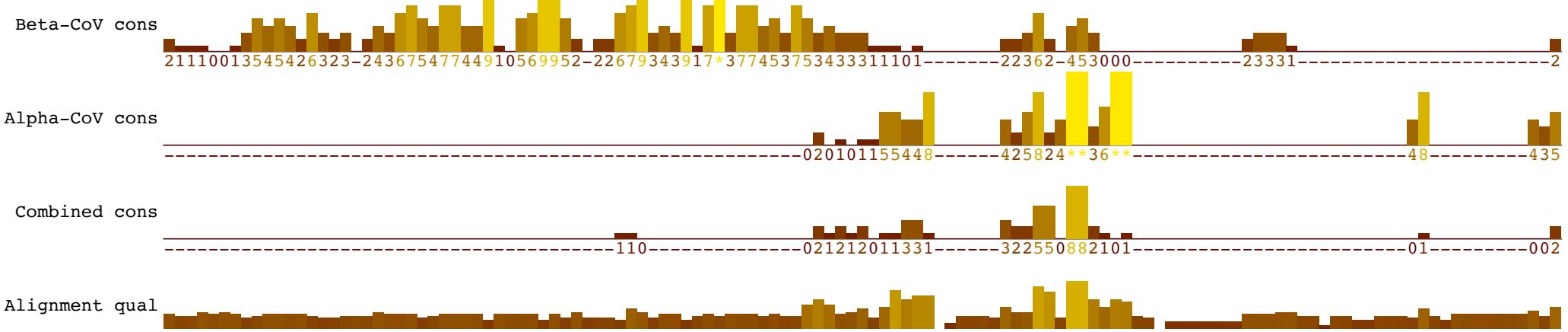
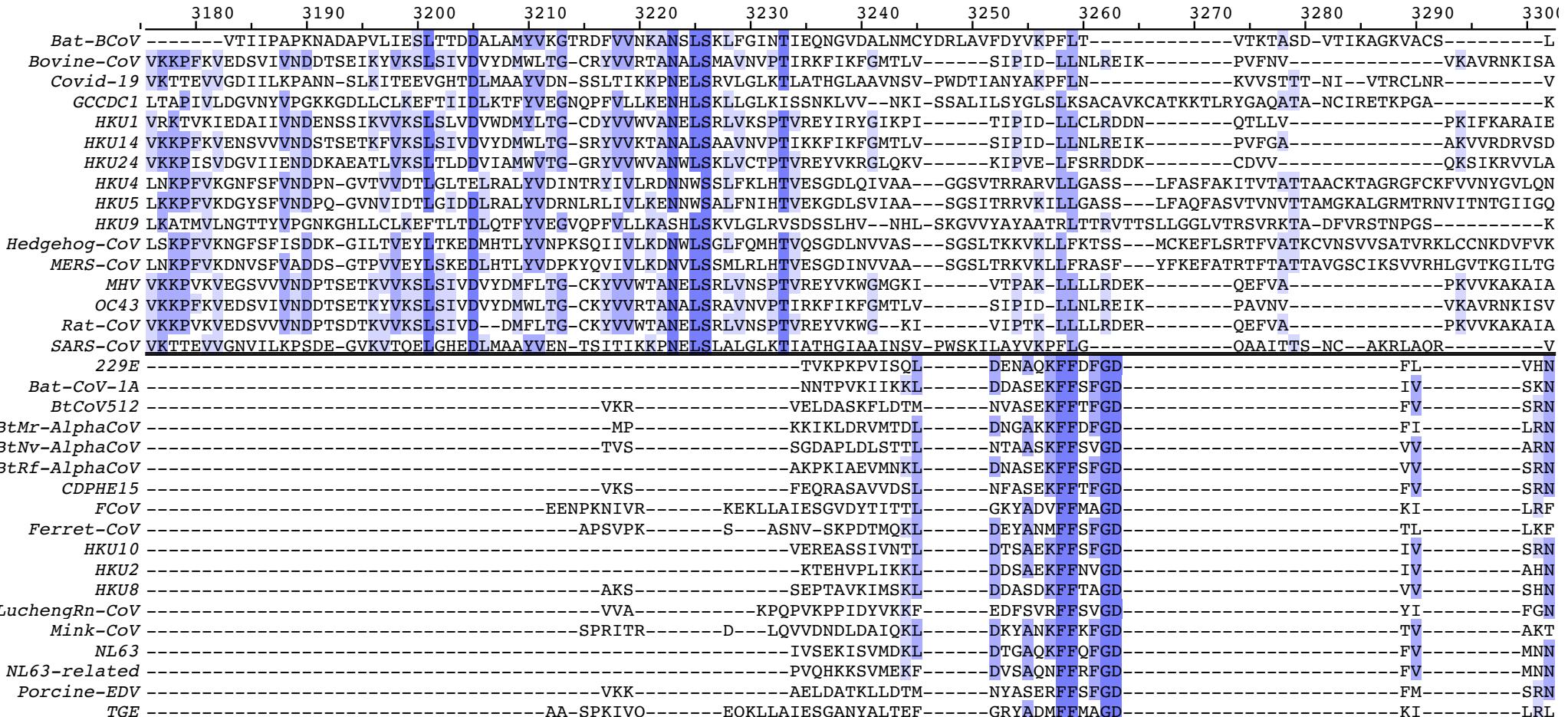
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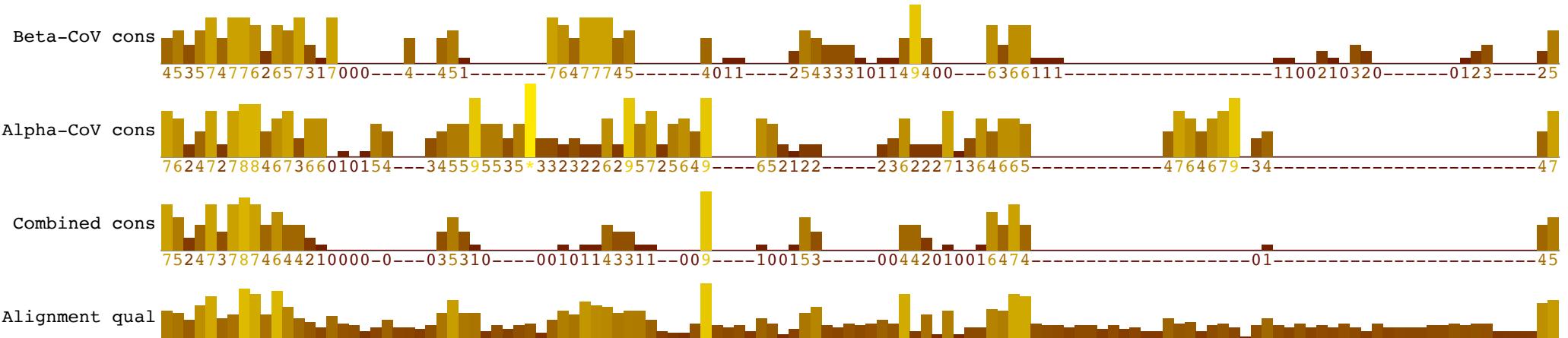
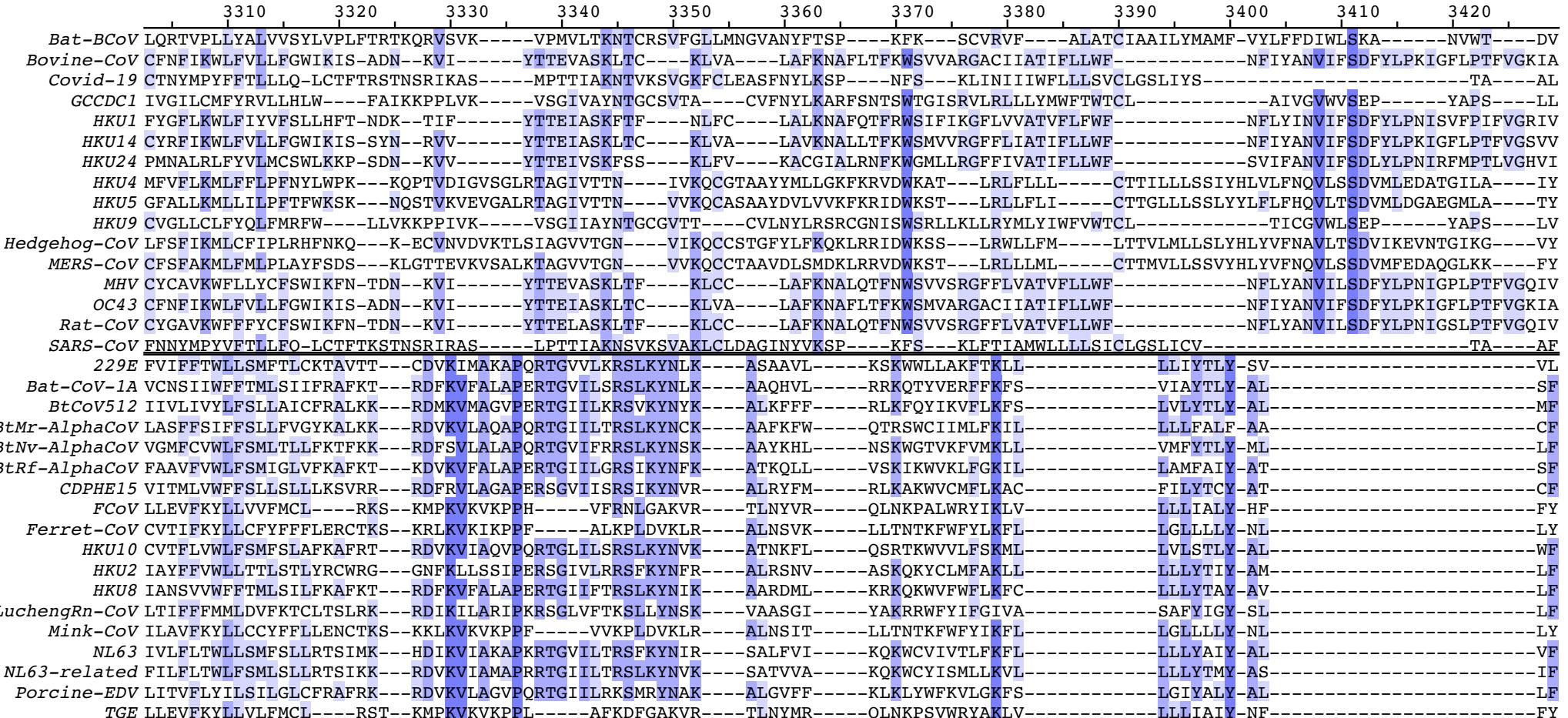


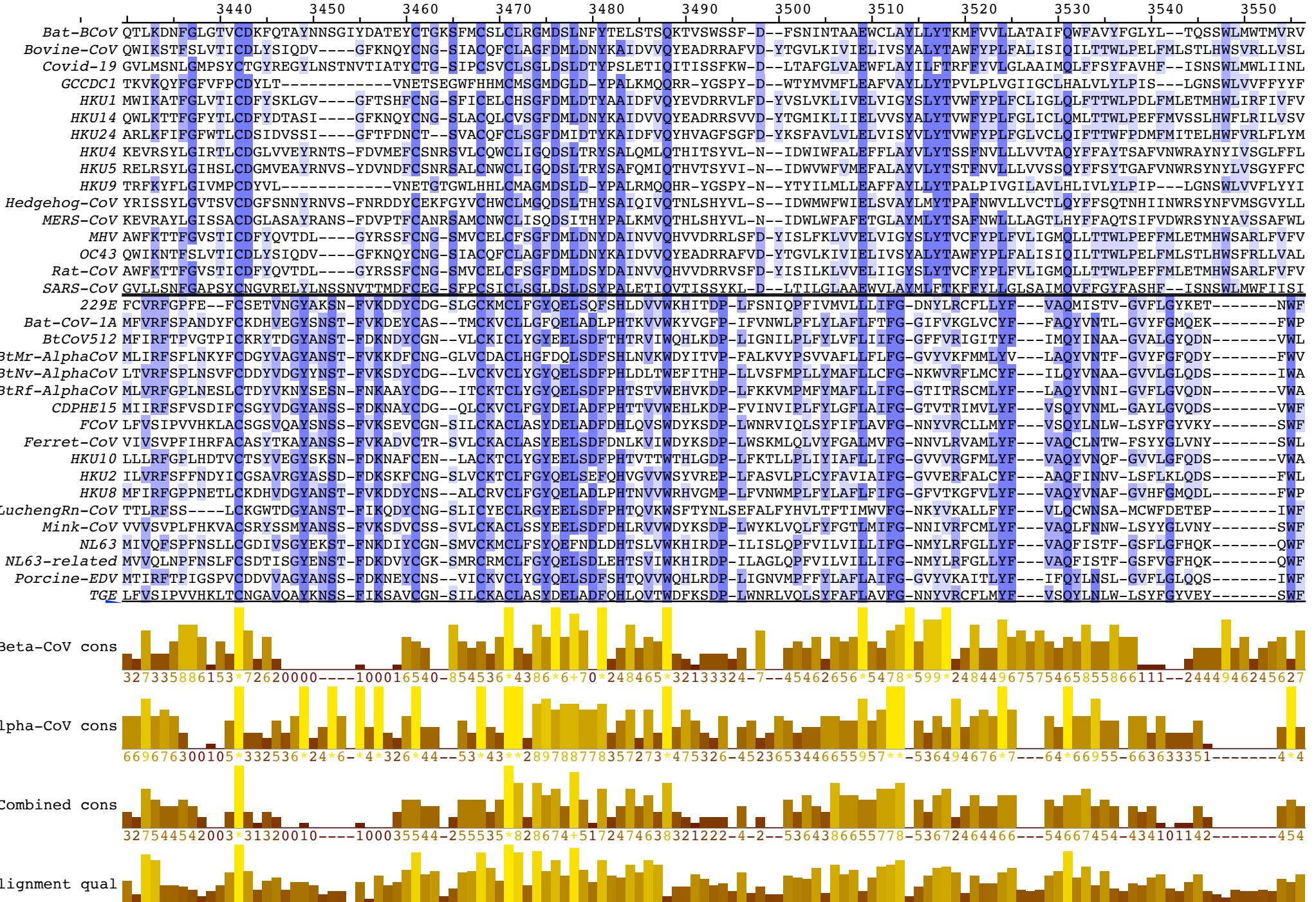


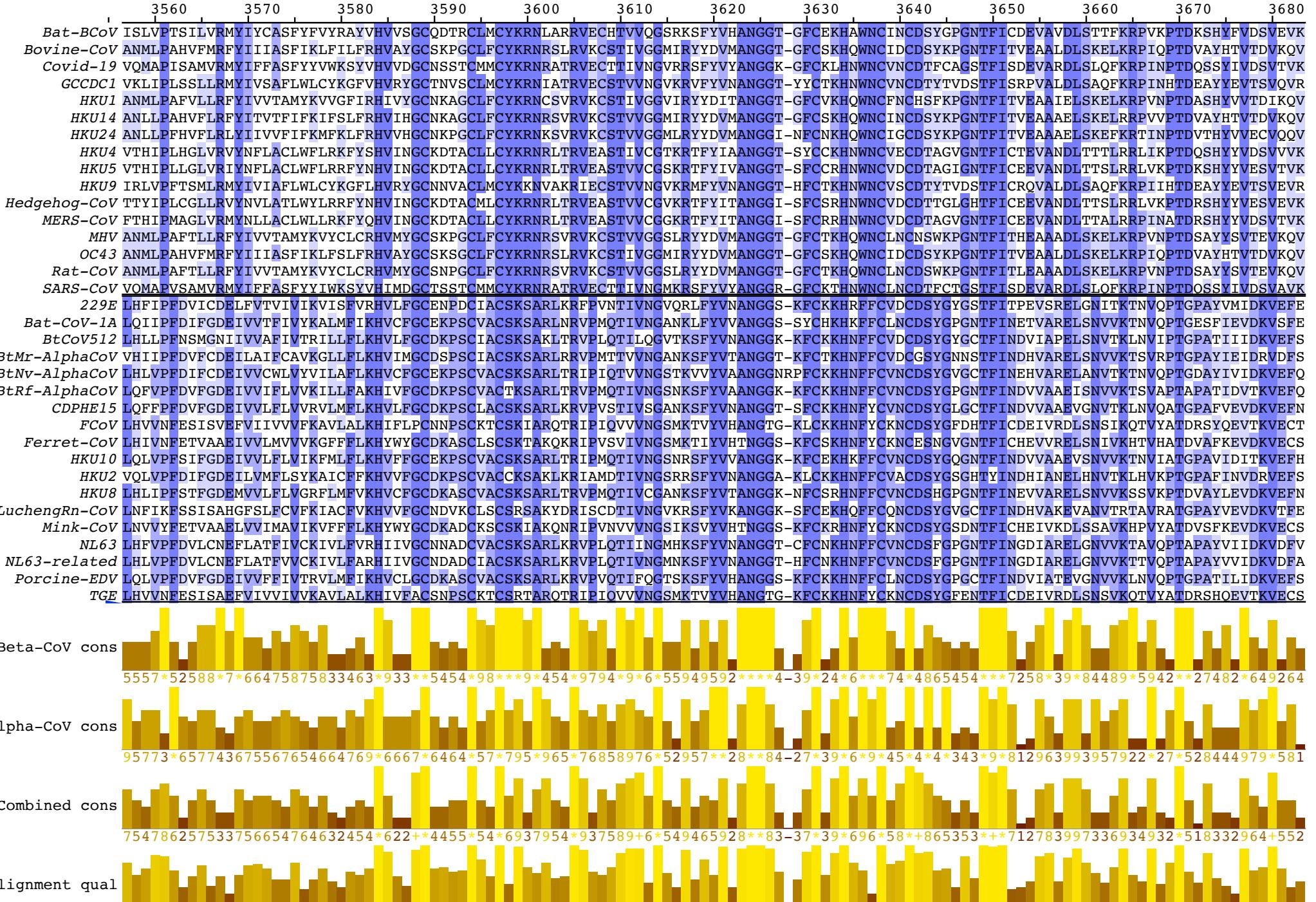


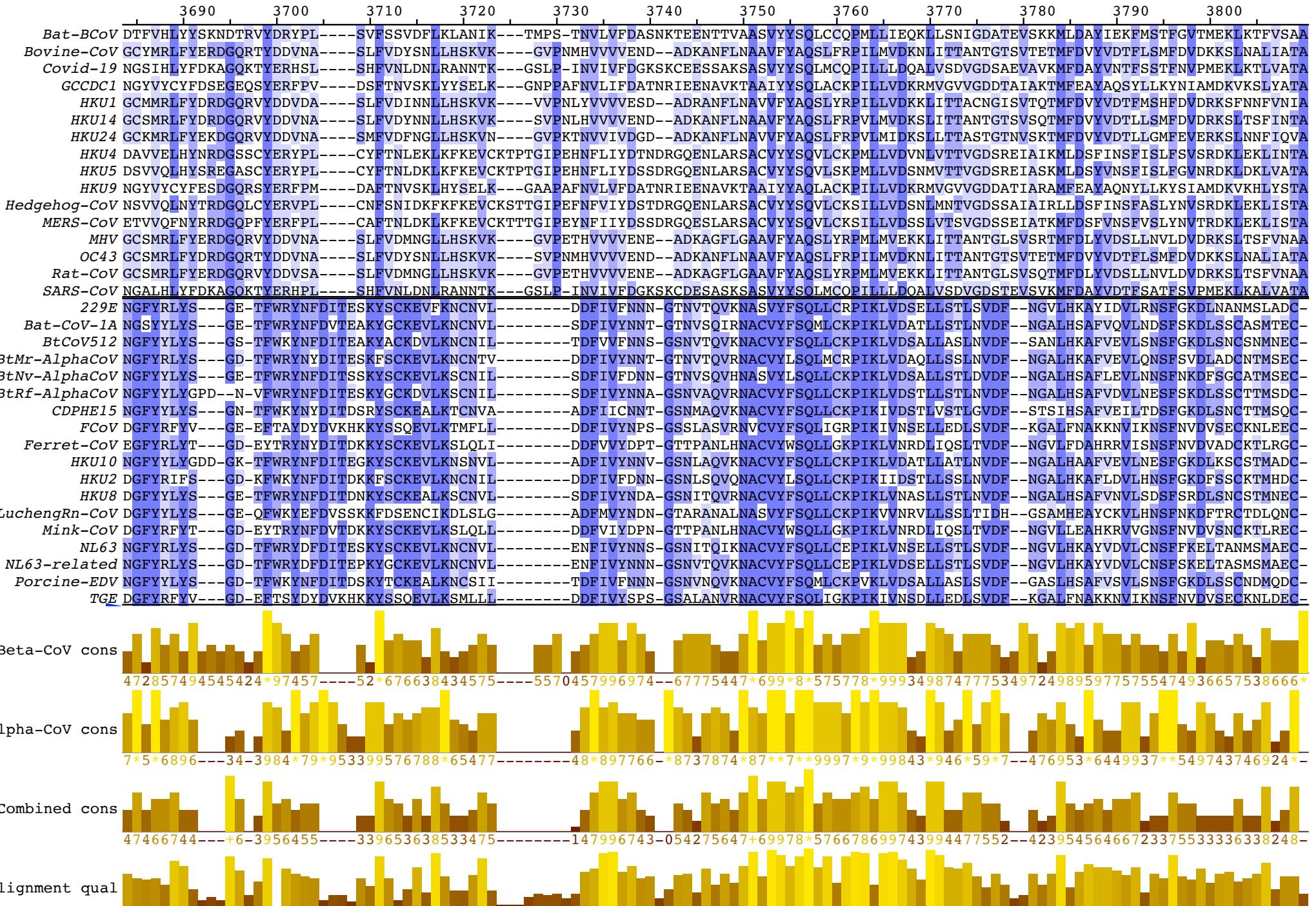


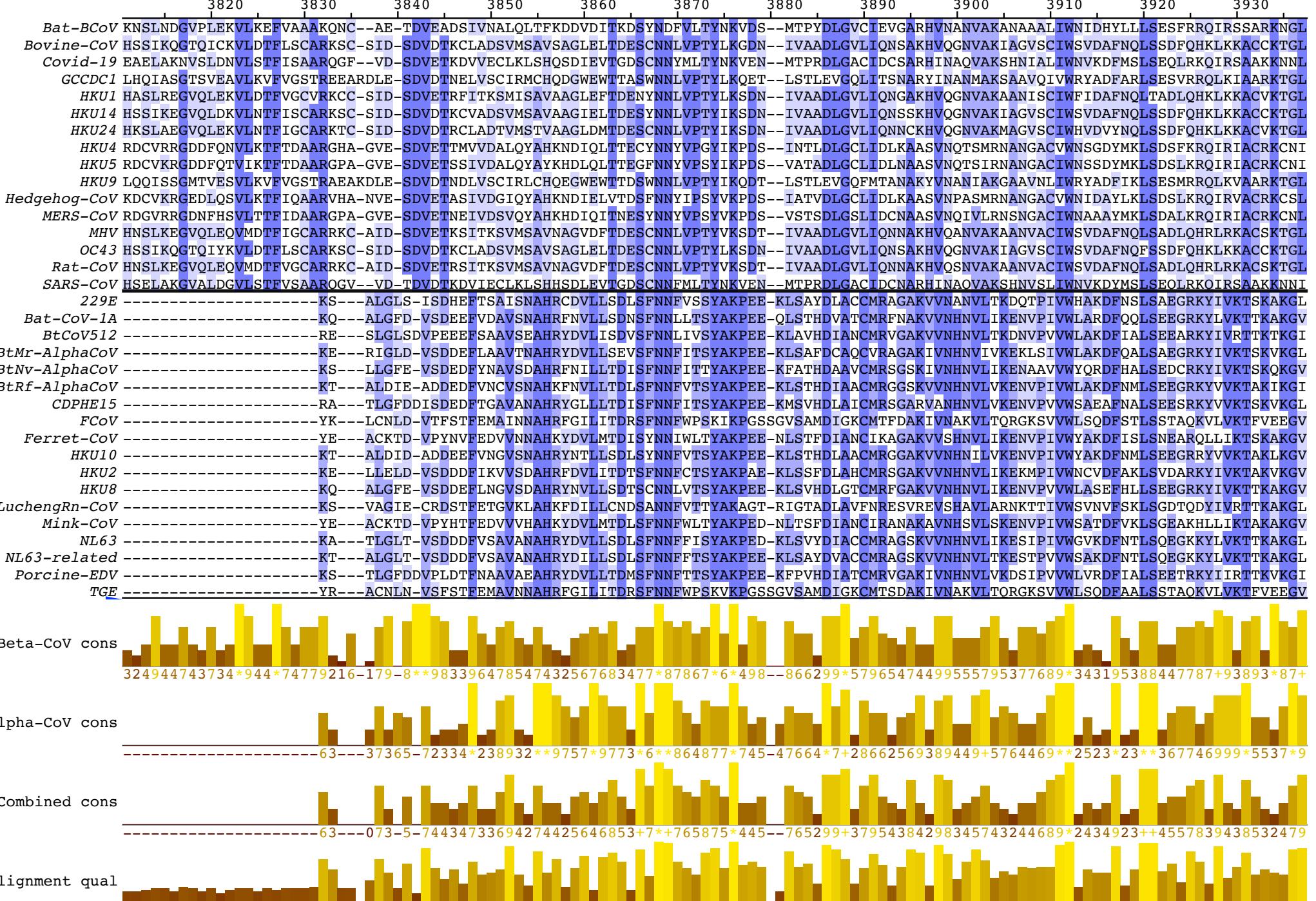


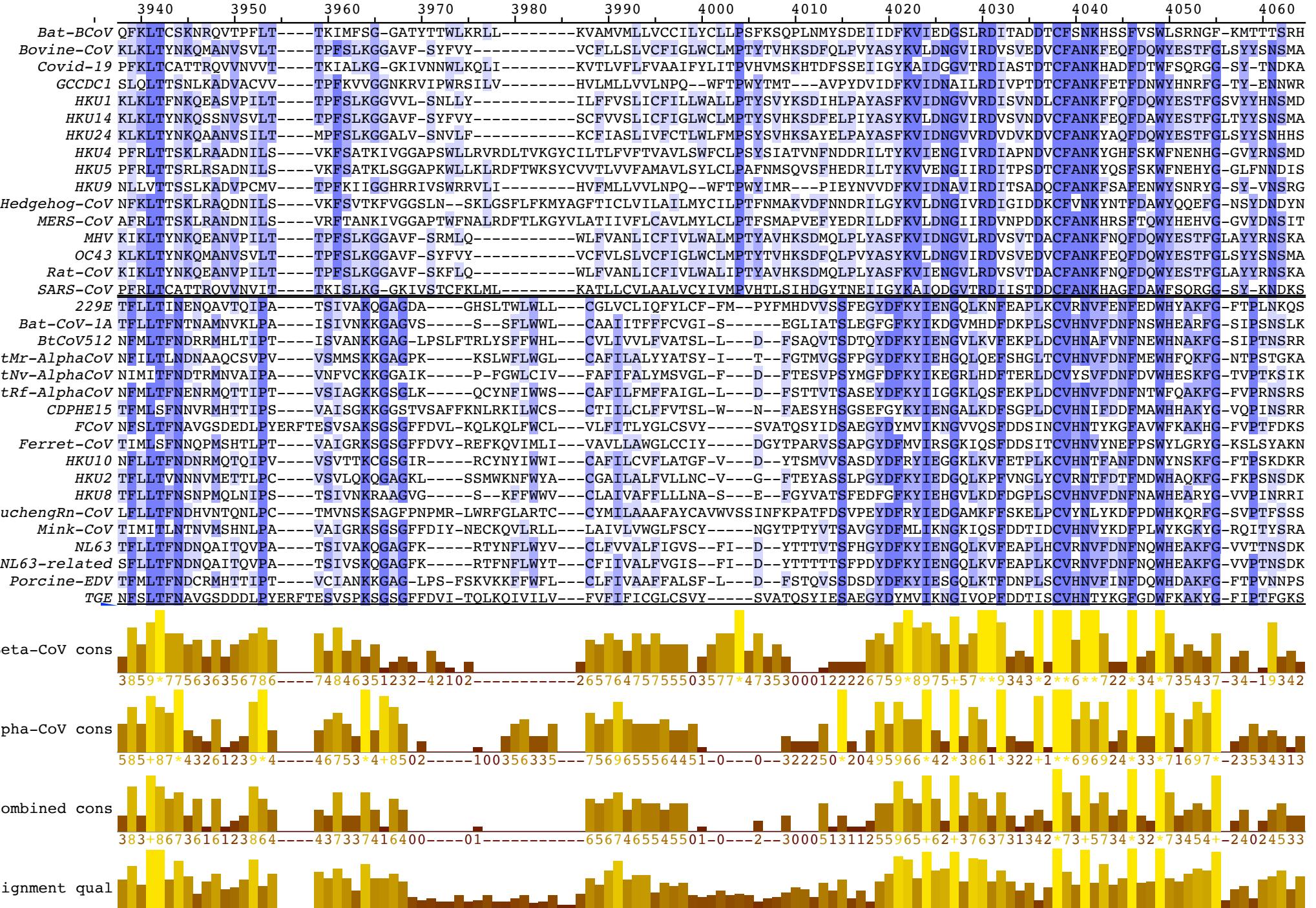


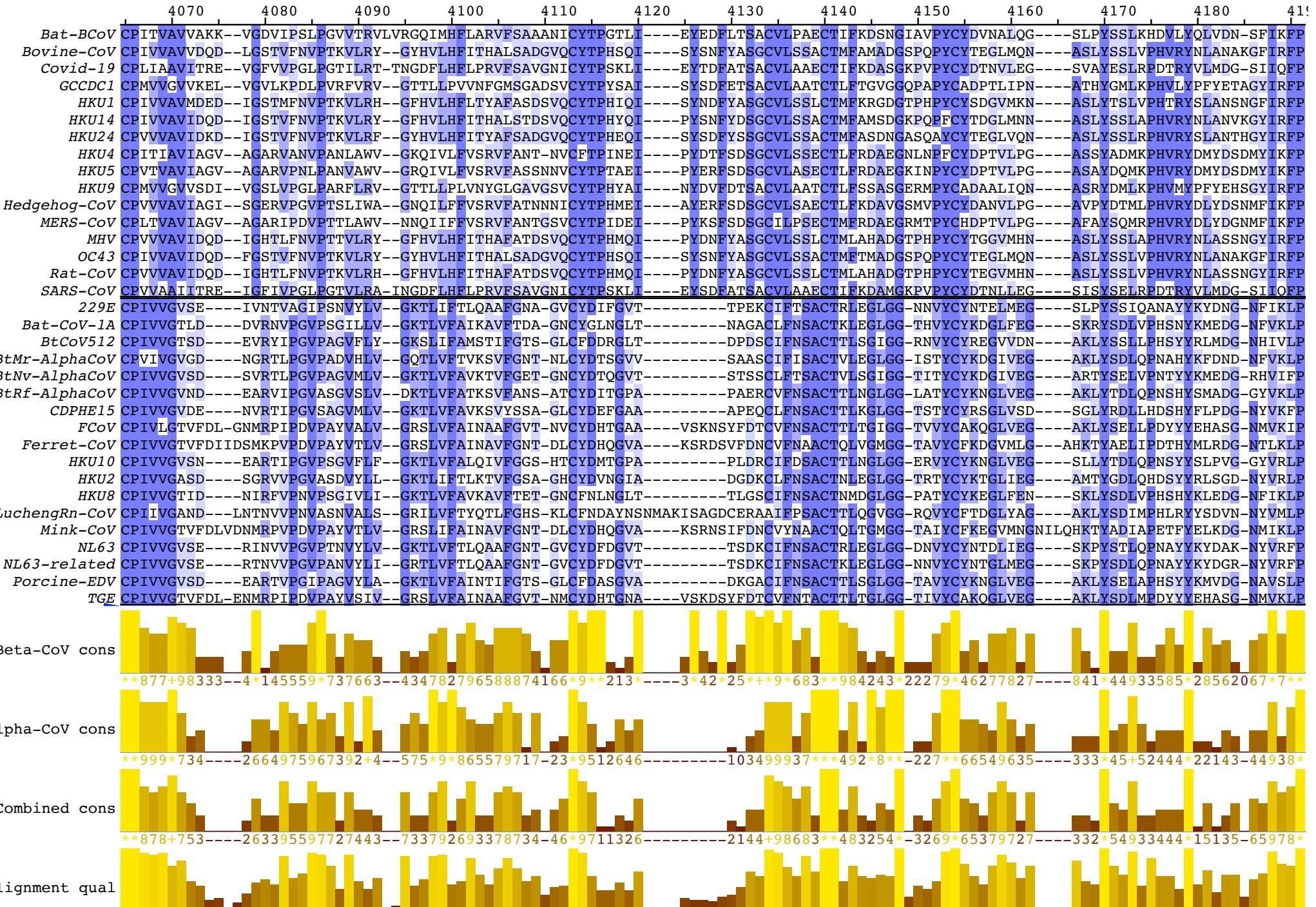


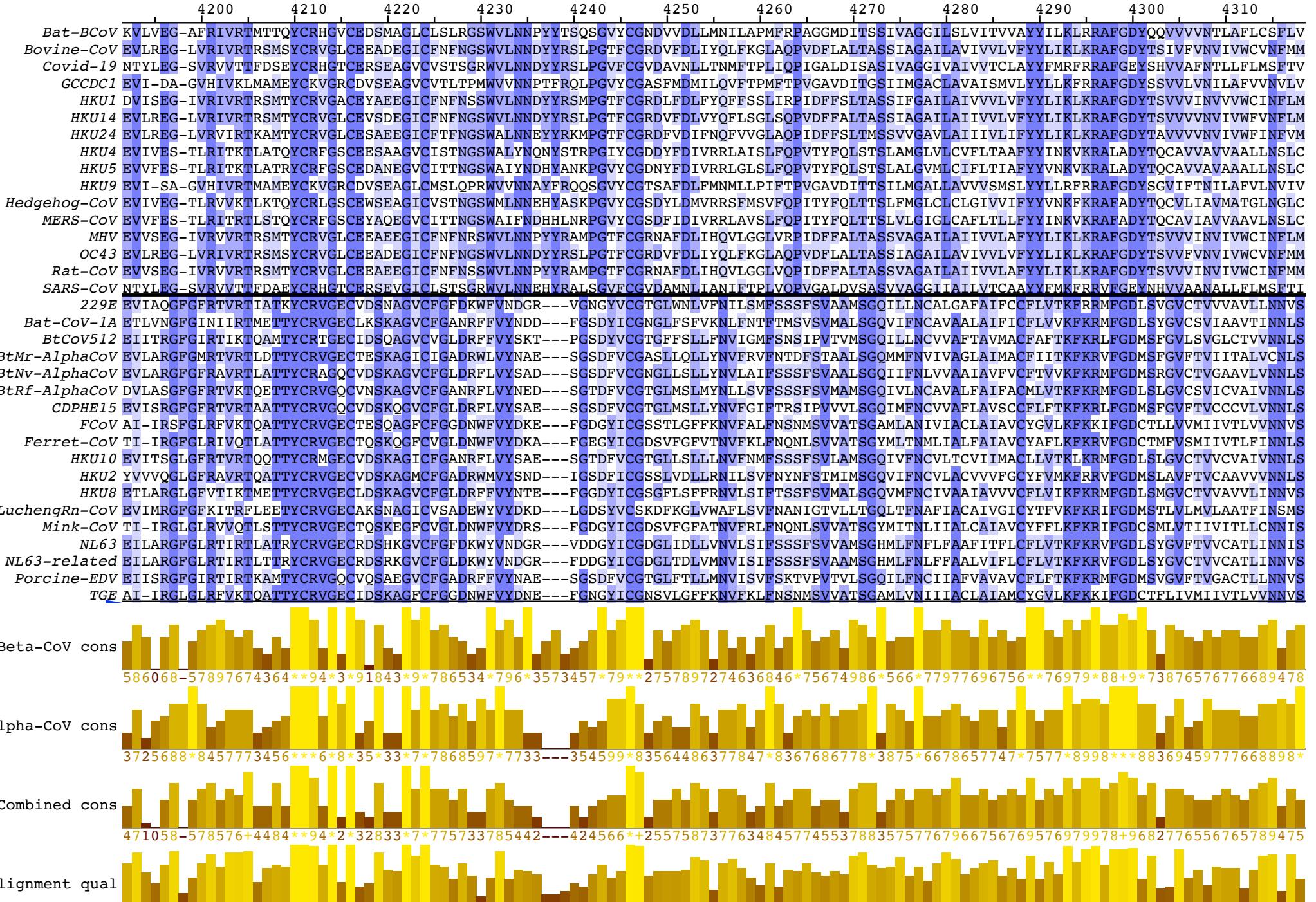


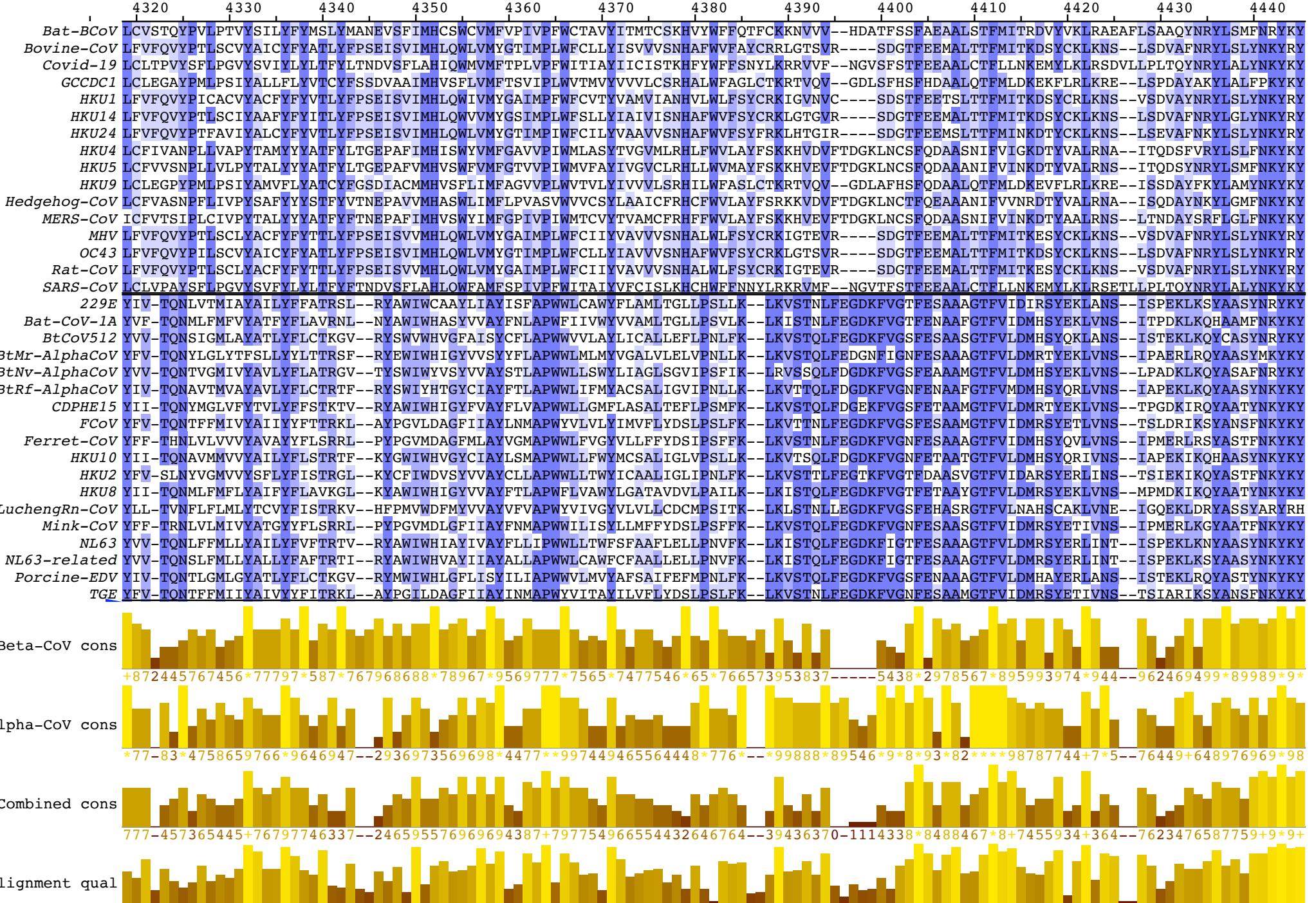


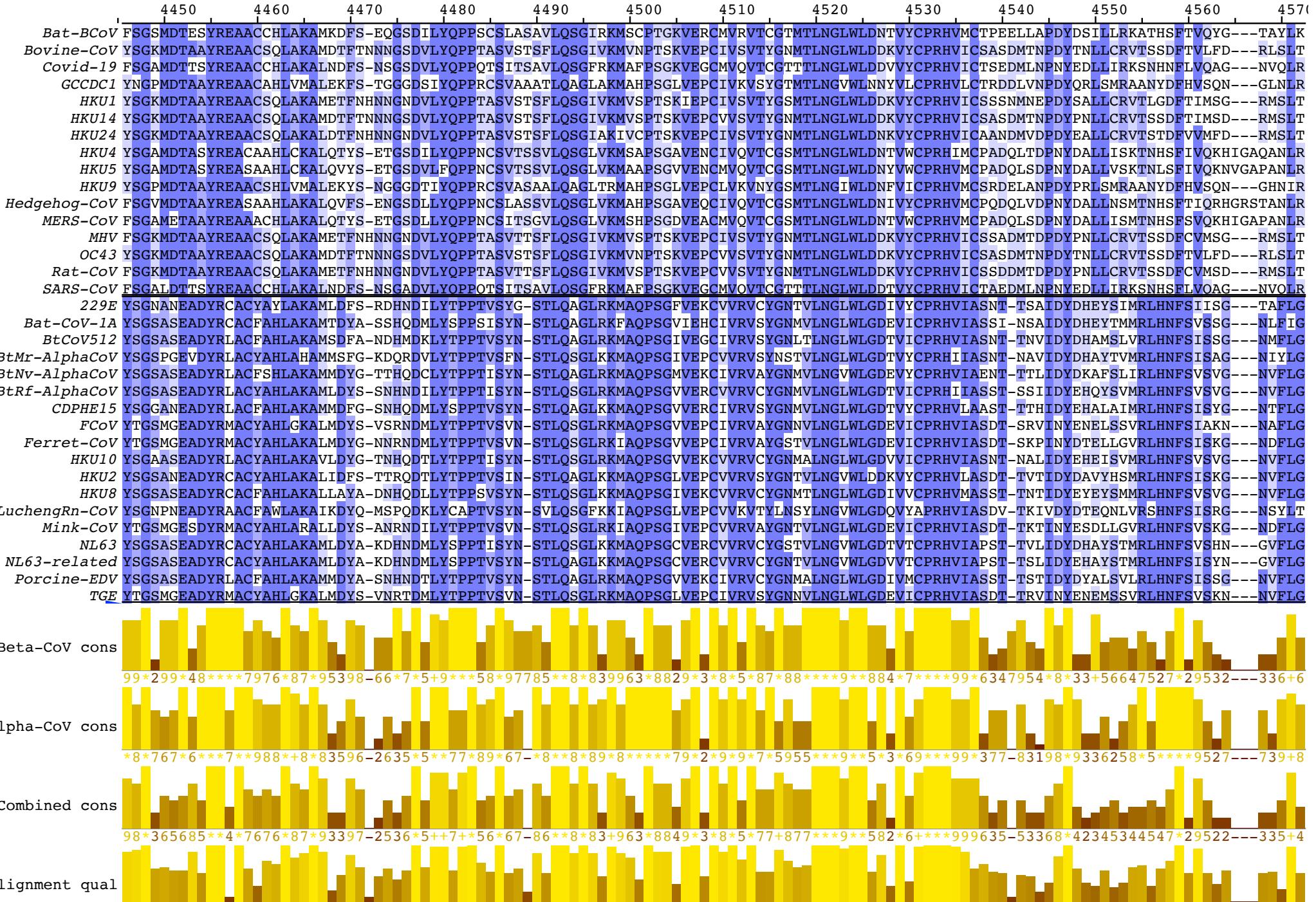


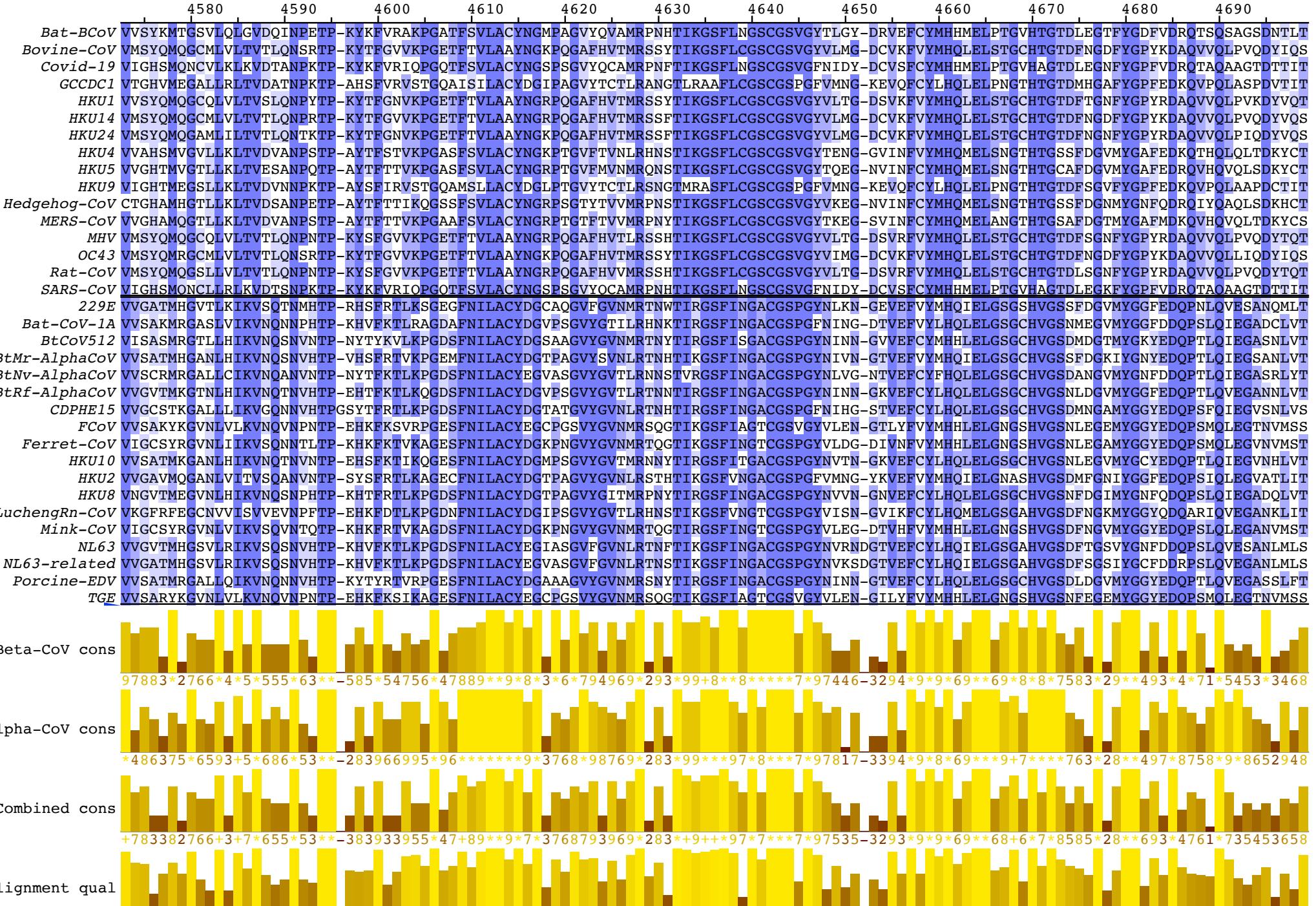


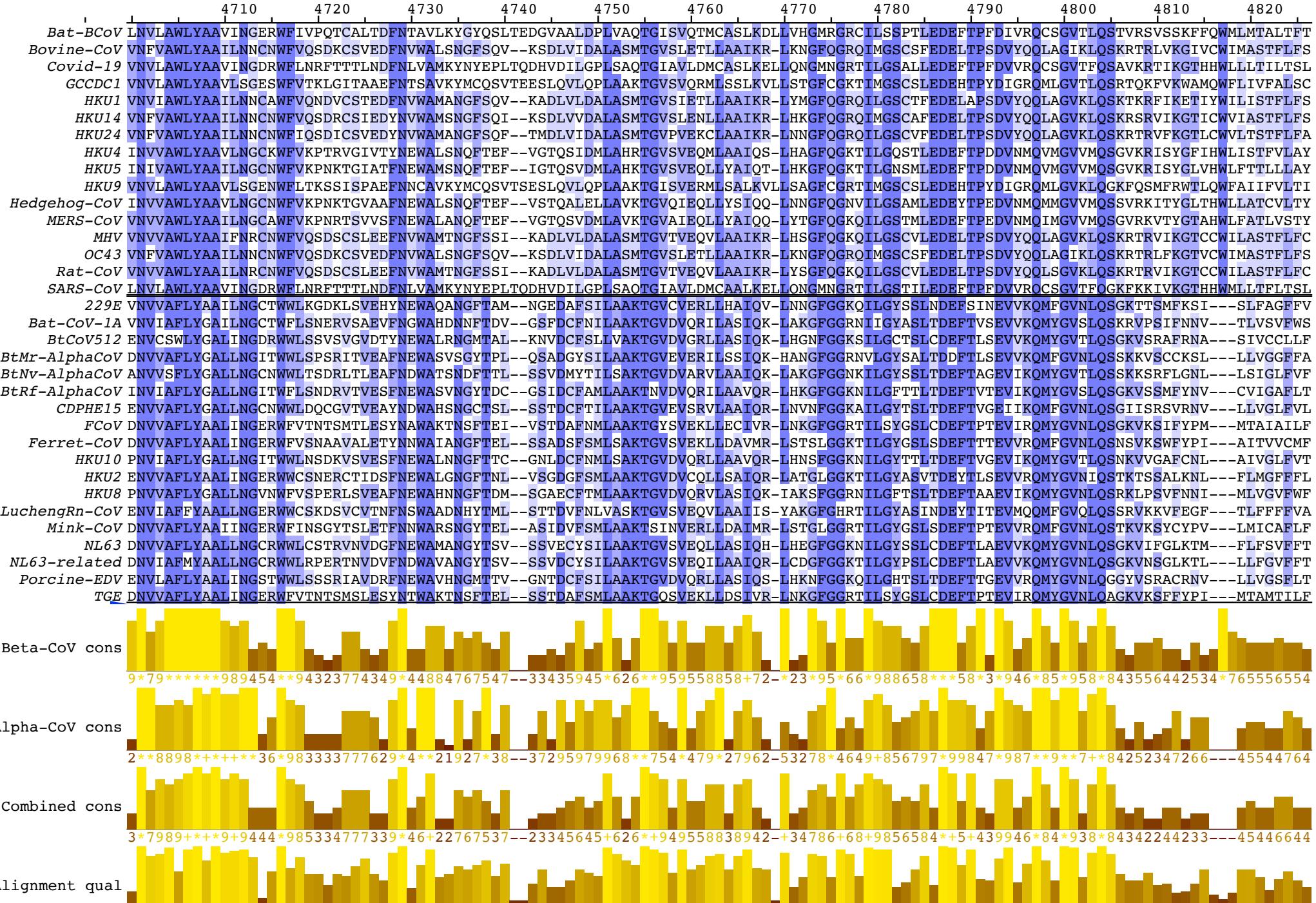




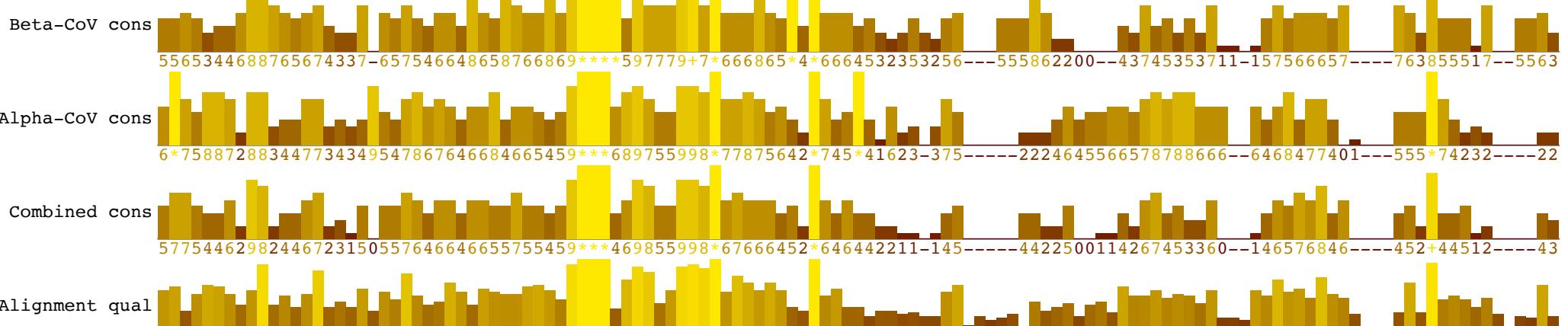


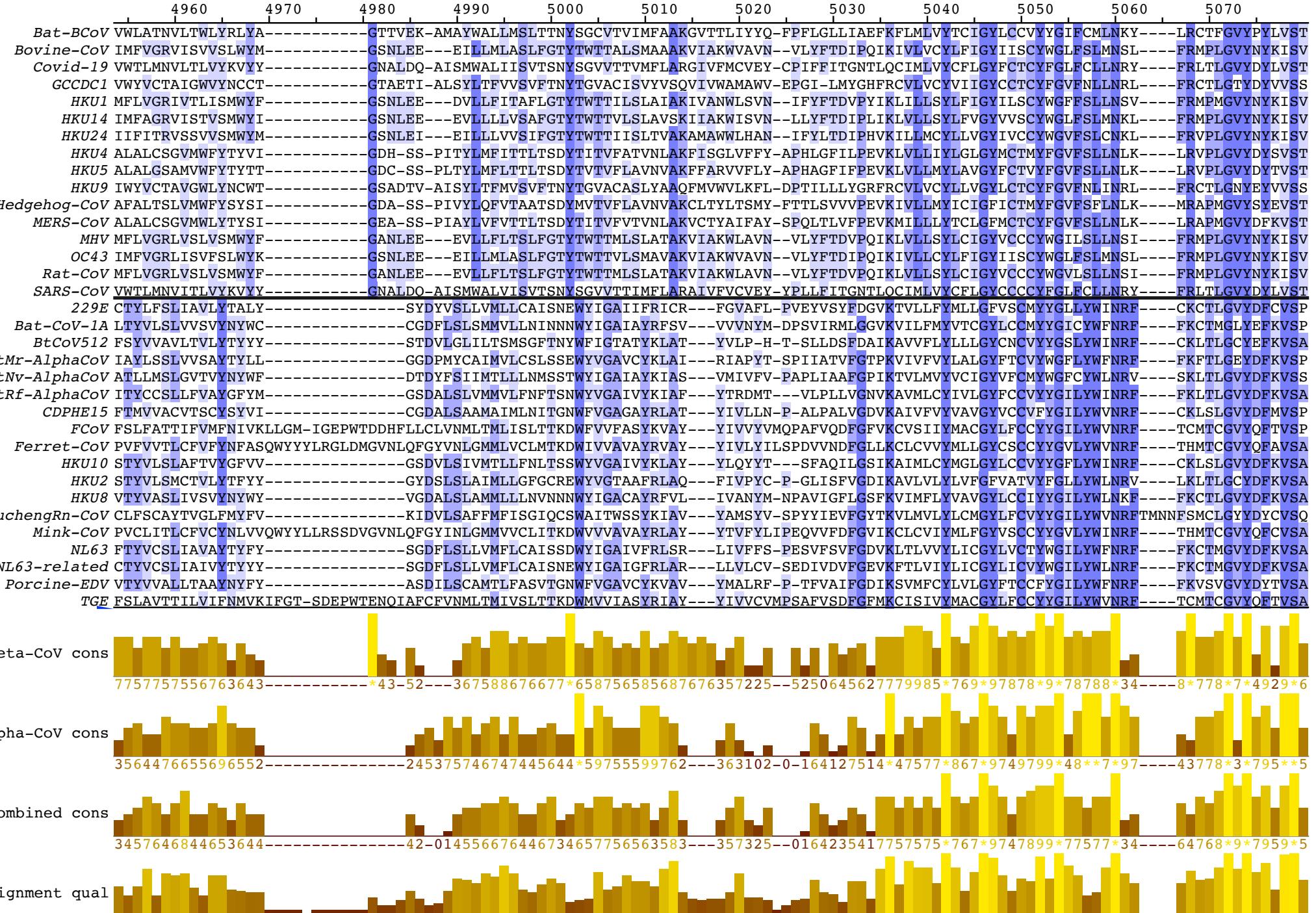


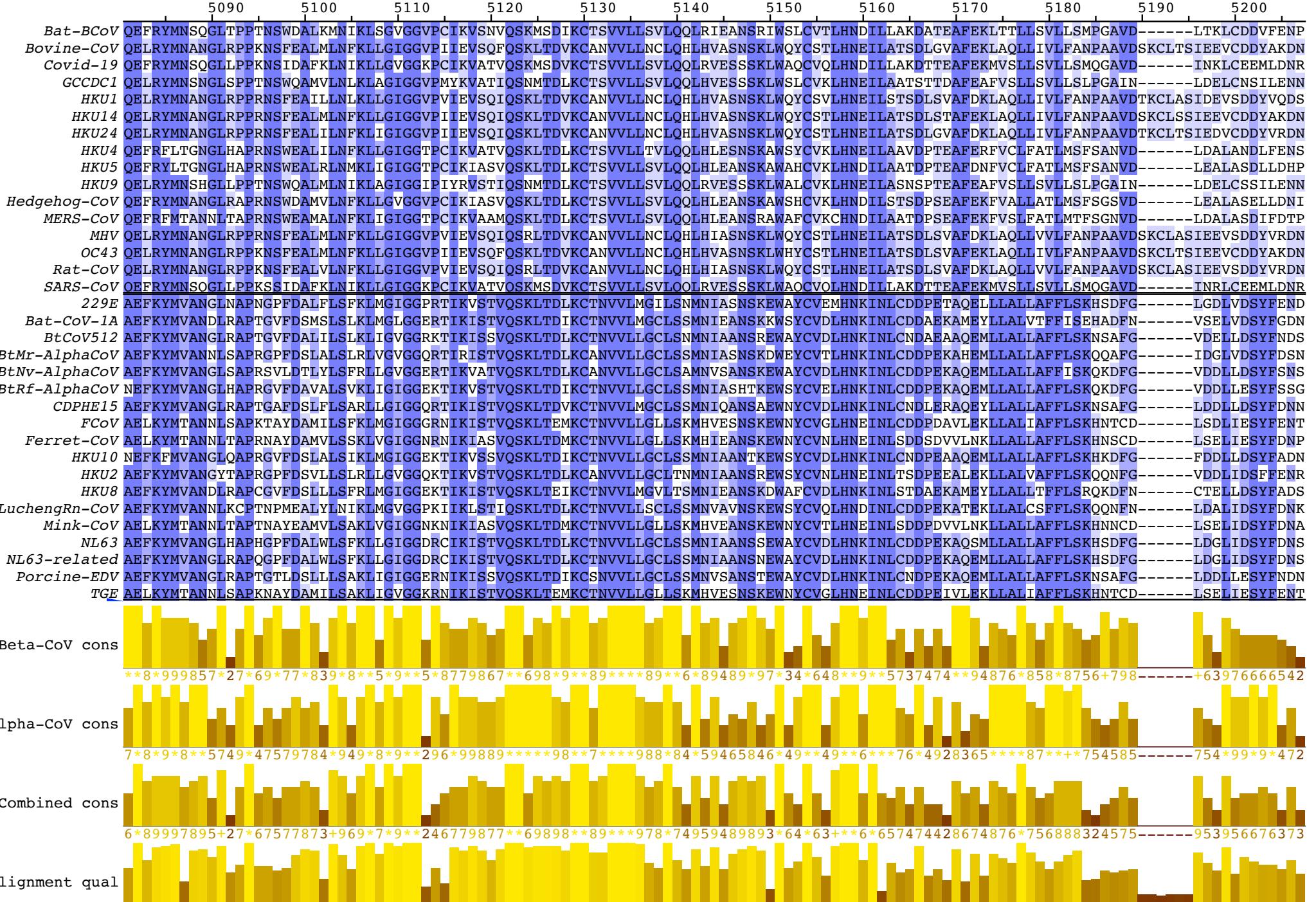


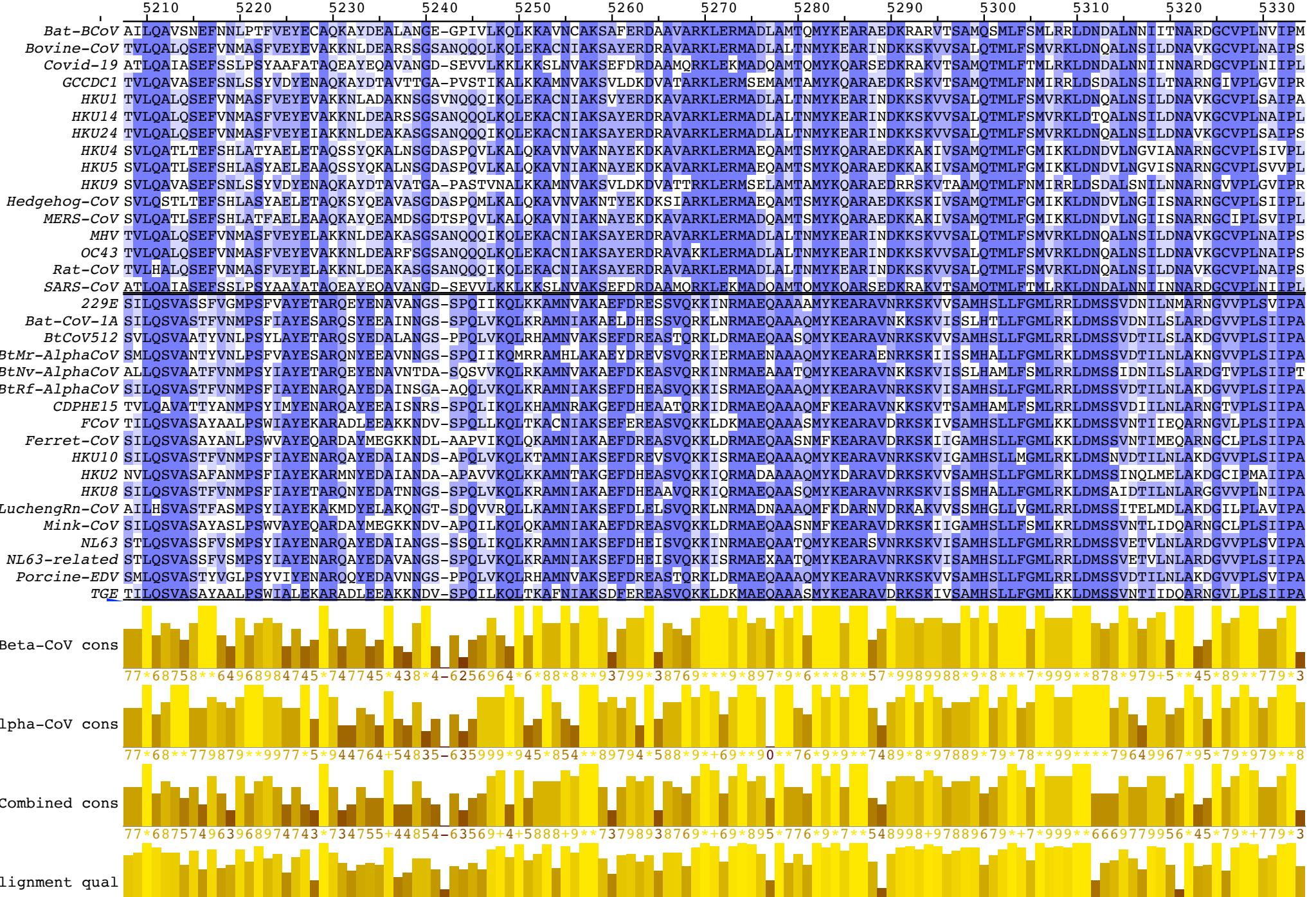


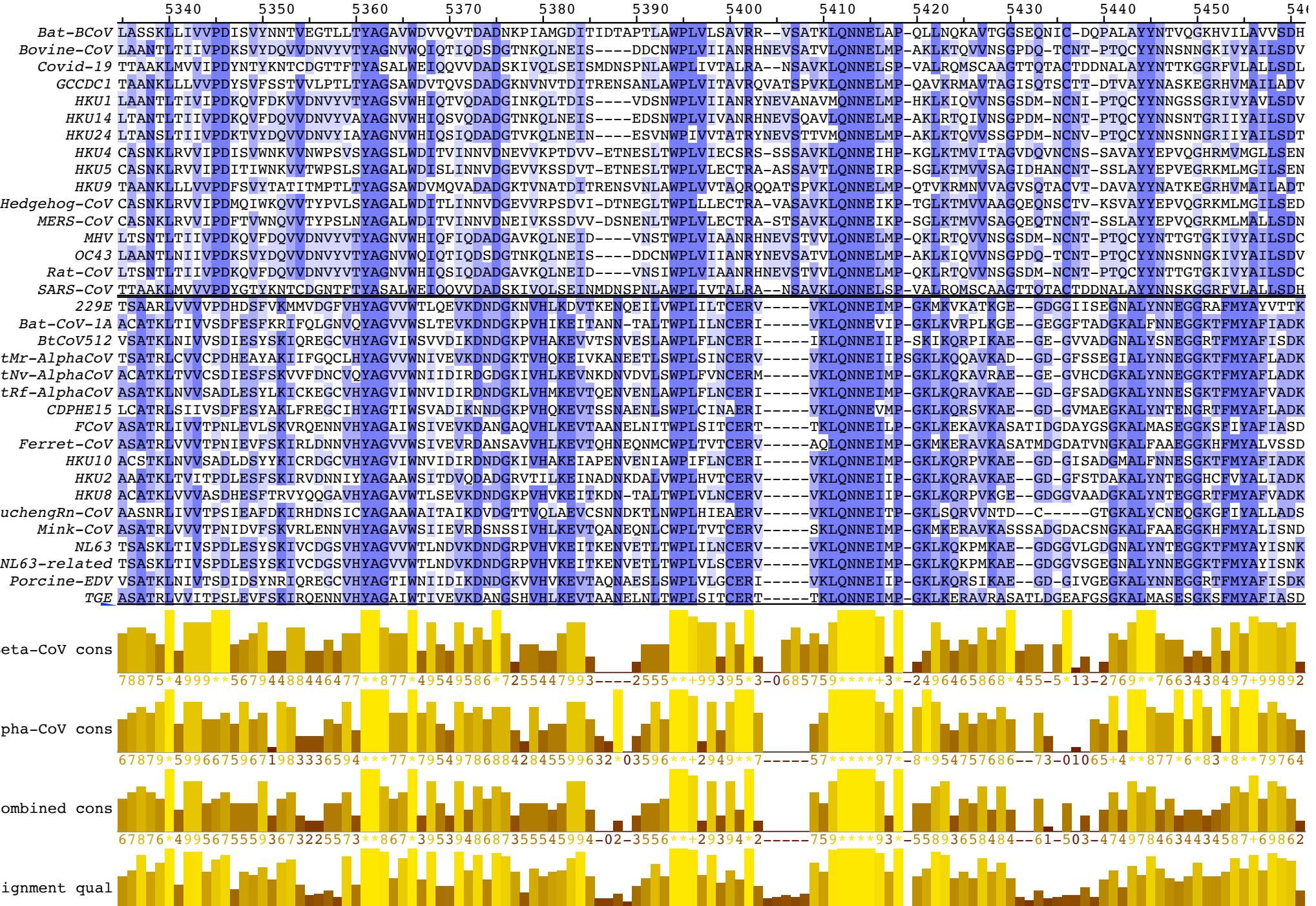
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*Covid-19* LVLVQSTQWSLFFFYENAFLPFAMGIAMSAFAMMFVKHKHAAFLCLFLPLSLATVAYFMNVMPASWVMRIM-T-WLDMVDT--SLSGFKLKDCVMY-ASAVVLLIL---MTARTVYDD--GARR  
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*Porcine-EDV* FFWSELVSYTKFFWVNPNGVTPMFACTSLLSLLMFTLKHKTFLQVFLIPALIVTSCINLAFDVEVYN-YLA---EHFDYHVSLMGFNAQGLVN--IFVCFVVTIHLGTYTWRFNTPA--SS  
*TGE* AFWLEEFFMYTPFTWINPTFVSVVLAITLVSLLVAGIKHKMFLFMSFVLPSPVILVTAHNLFWDFSYYESLOS---IVENTNTMFLPVDMOGVML--TVFCFIVFVT--YSVRFFFTCKO--SW

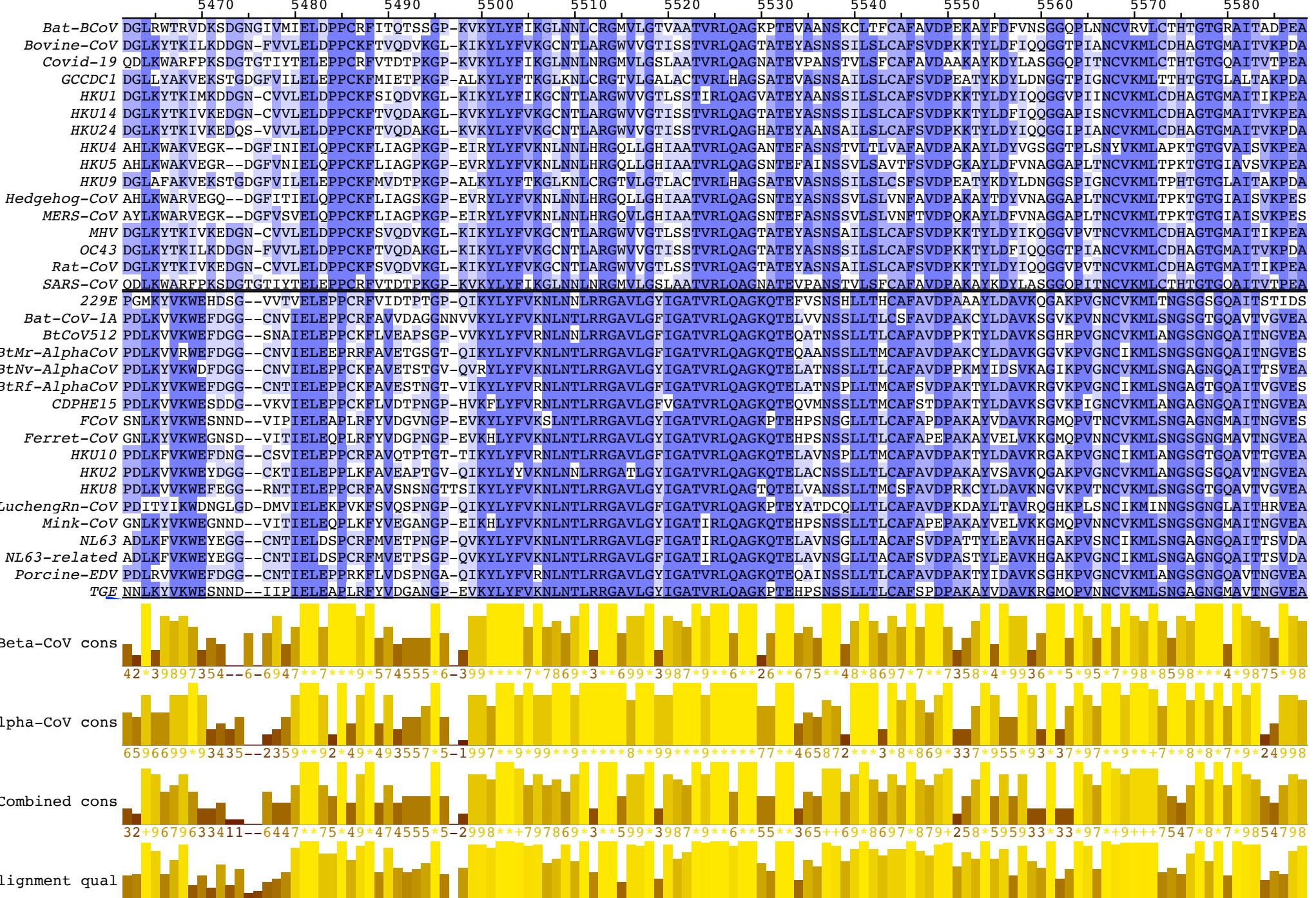


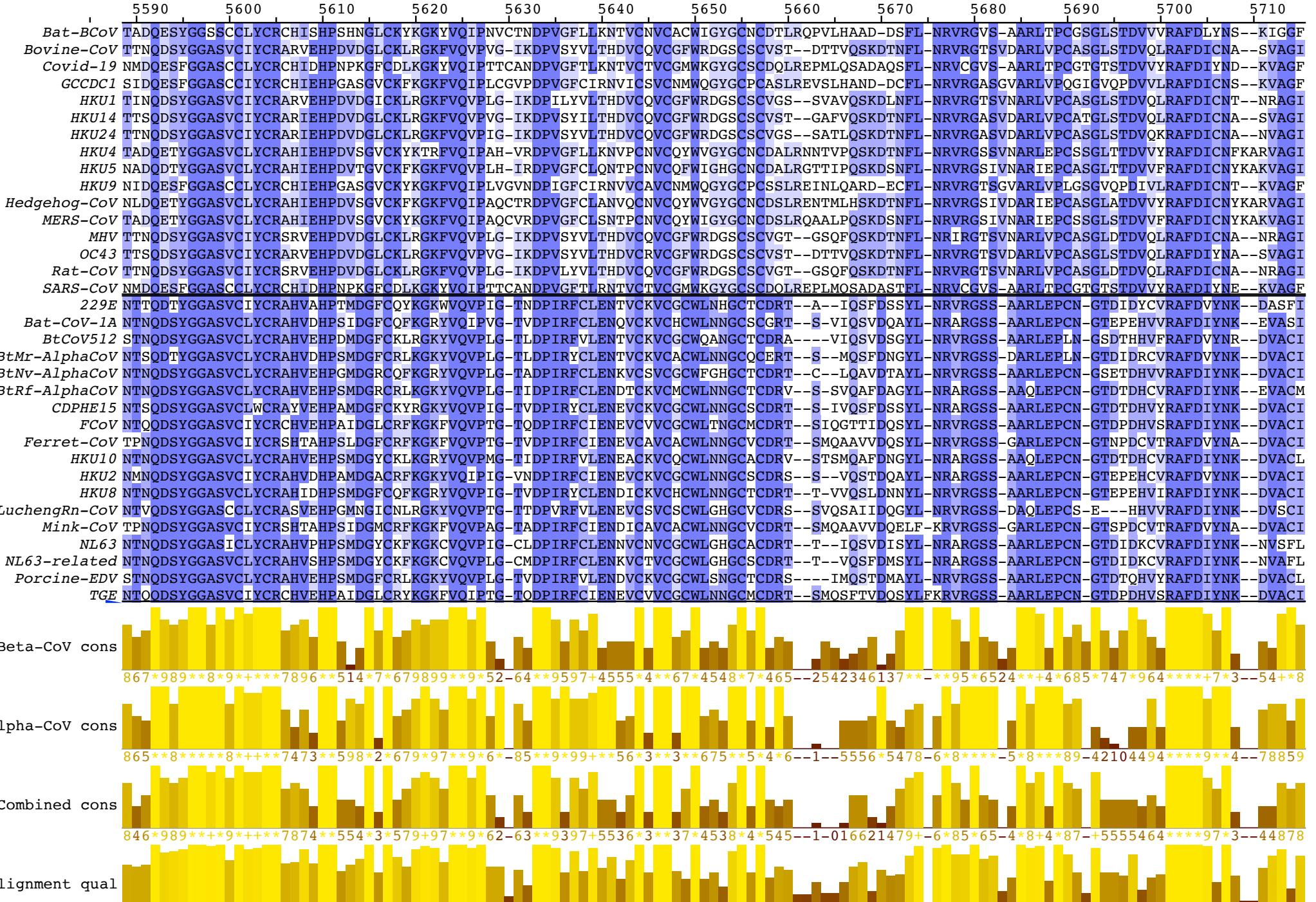


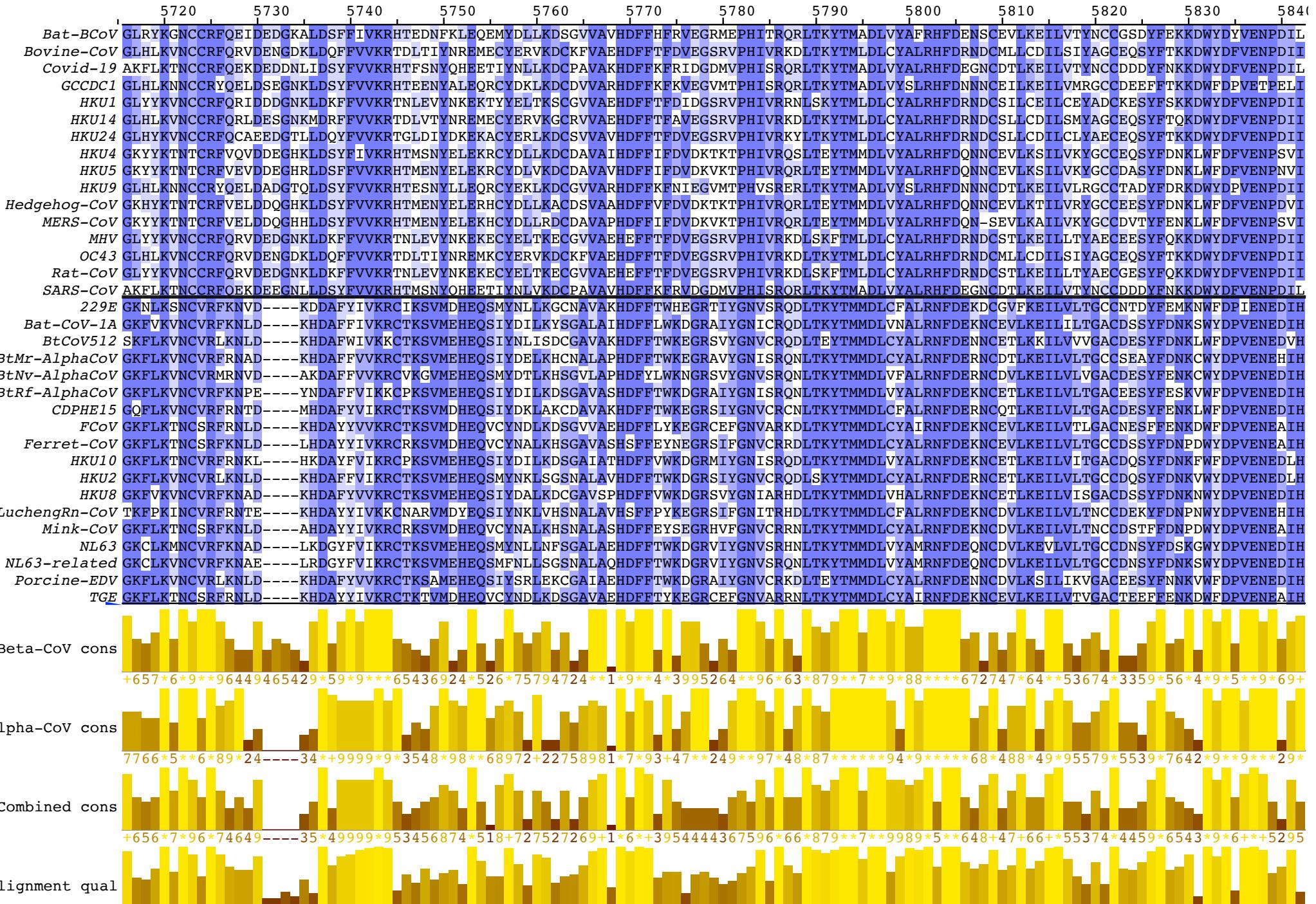


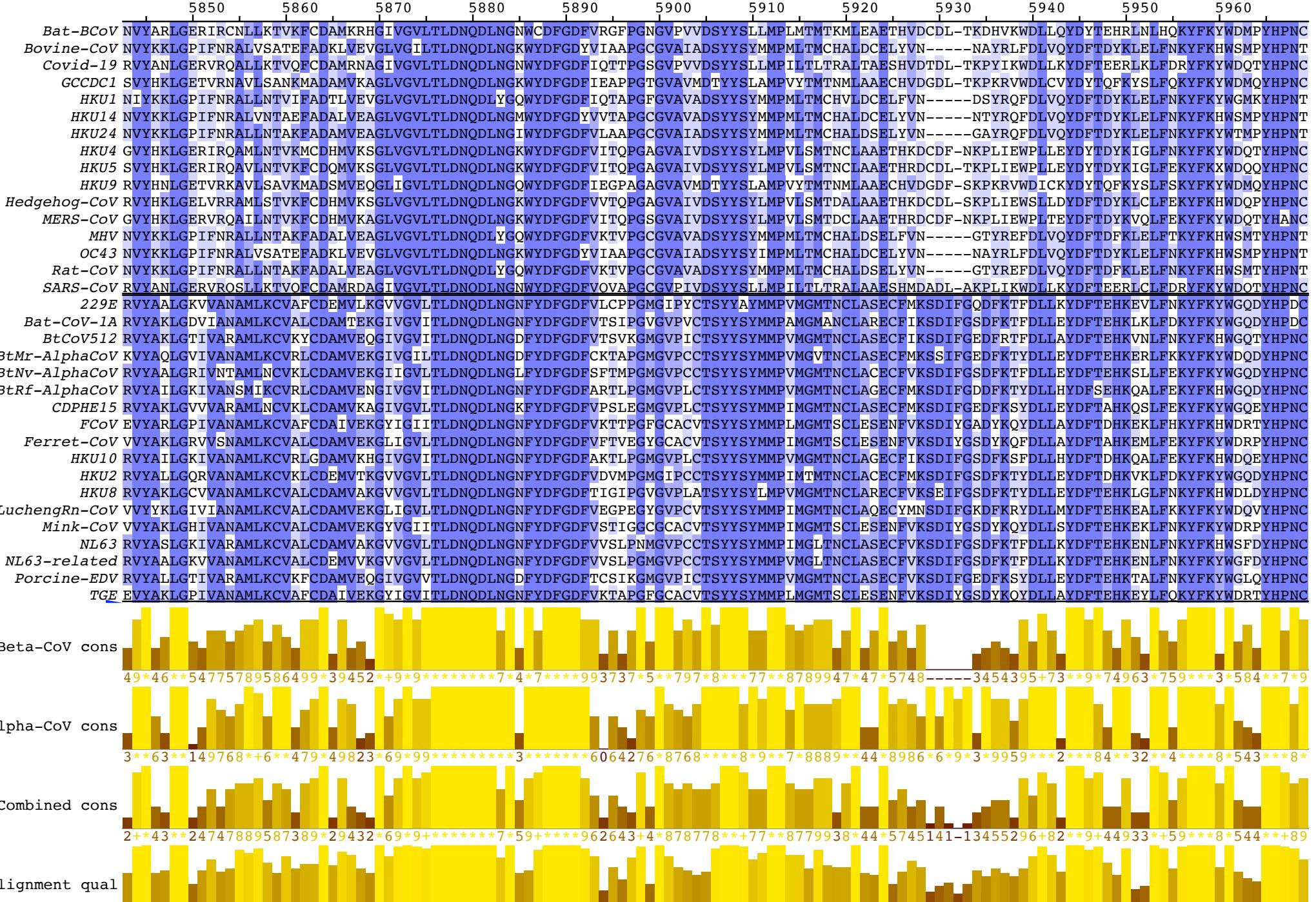


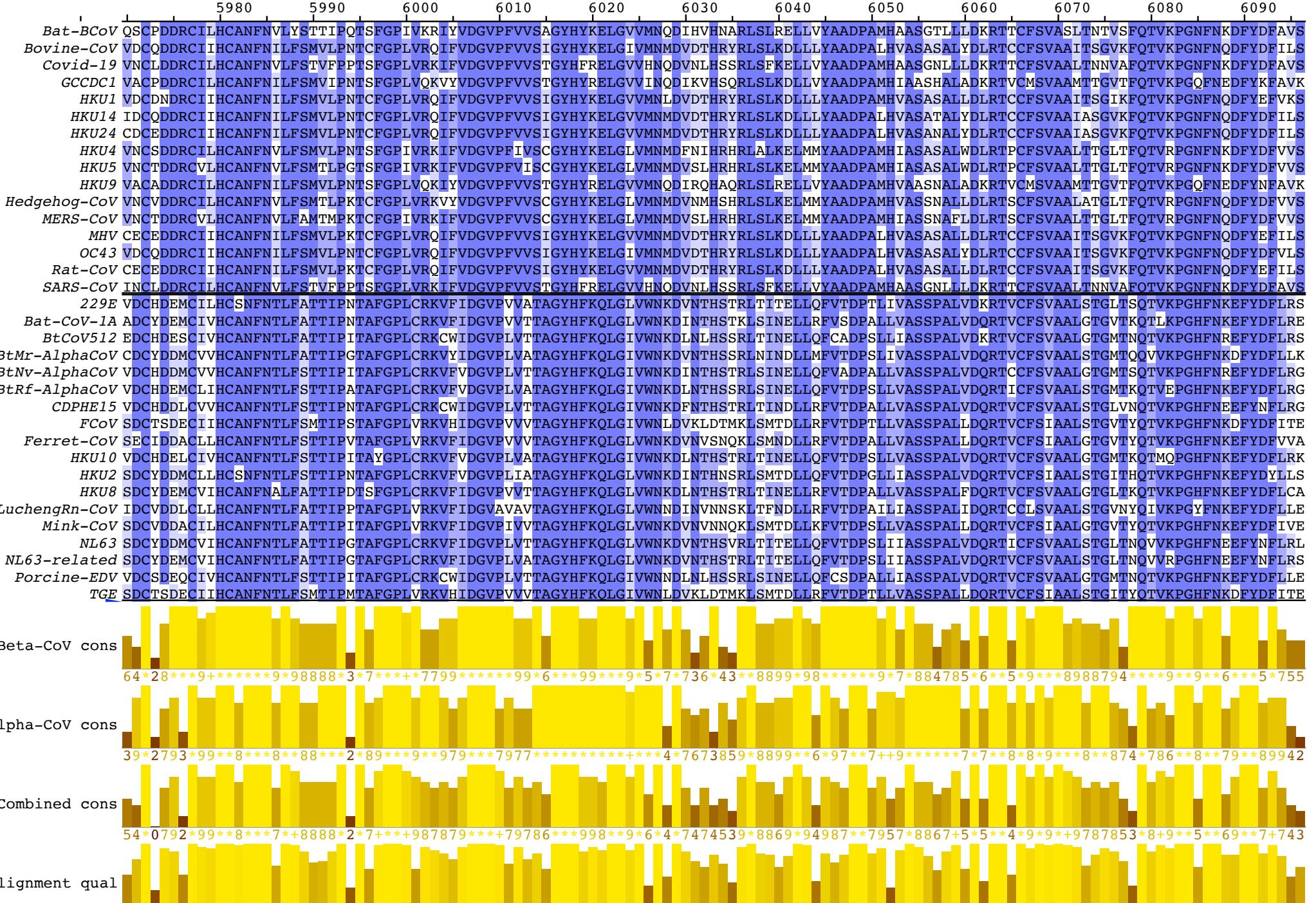


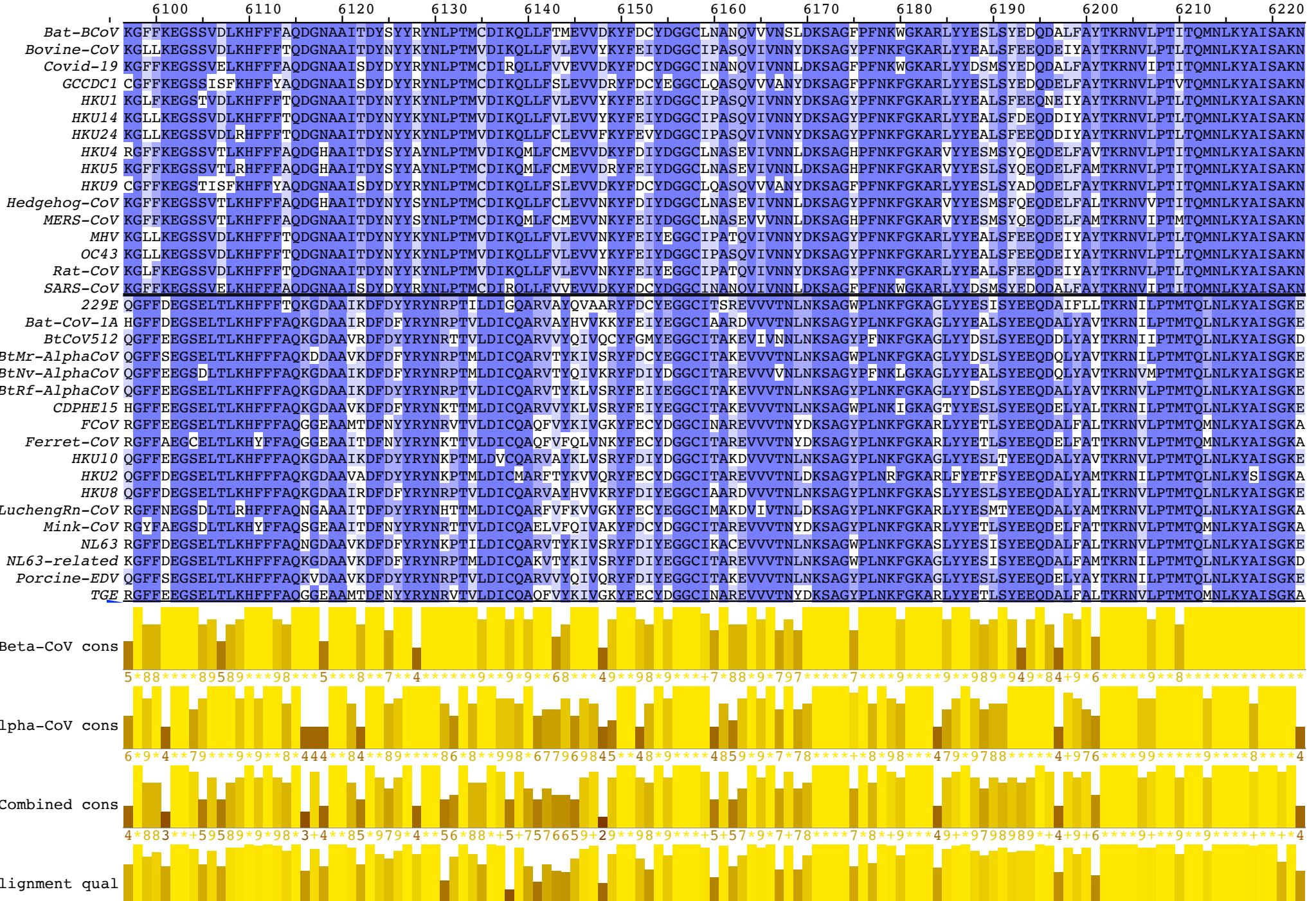


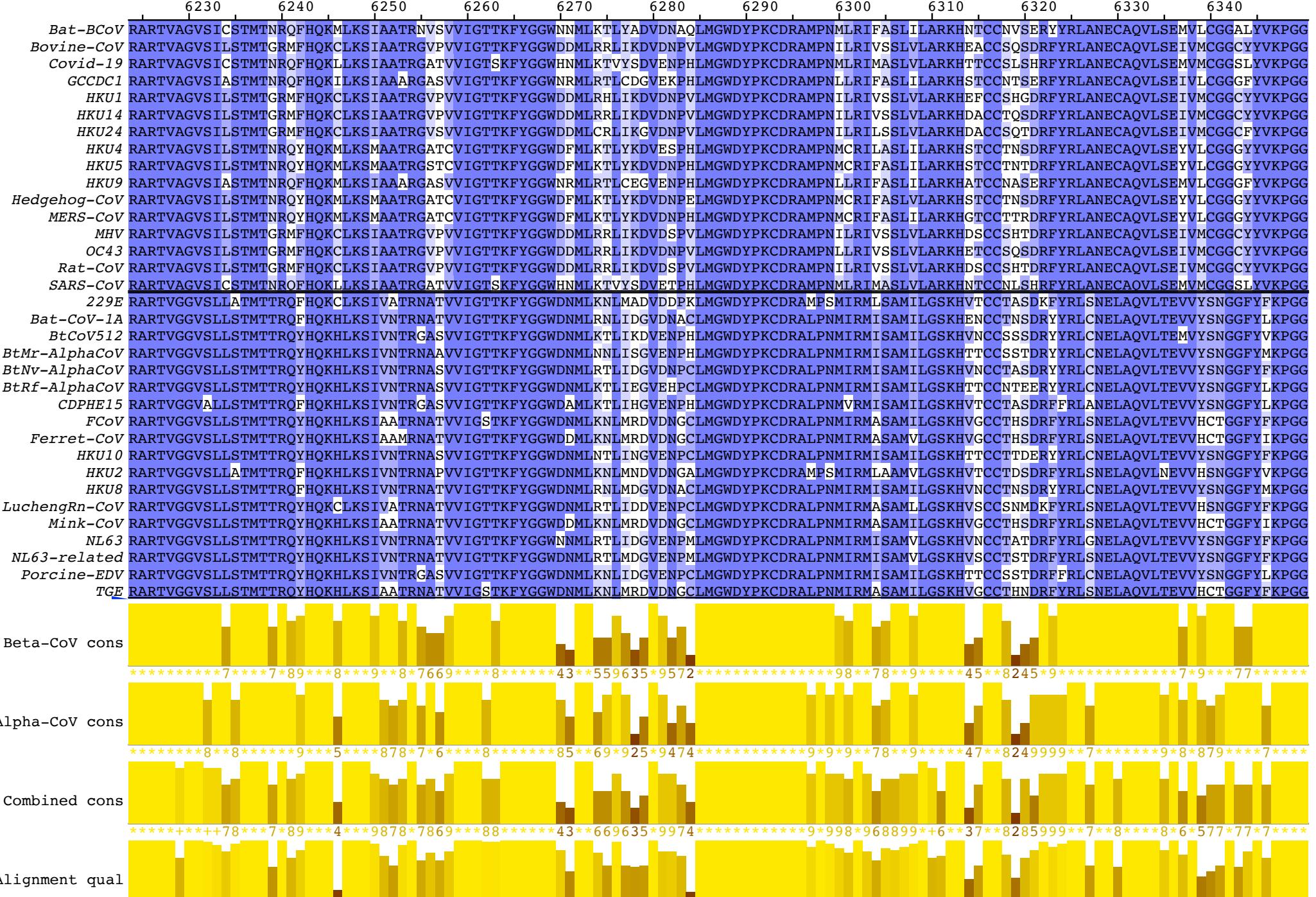


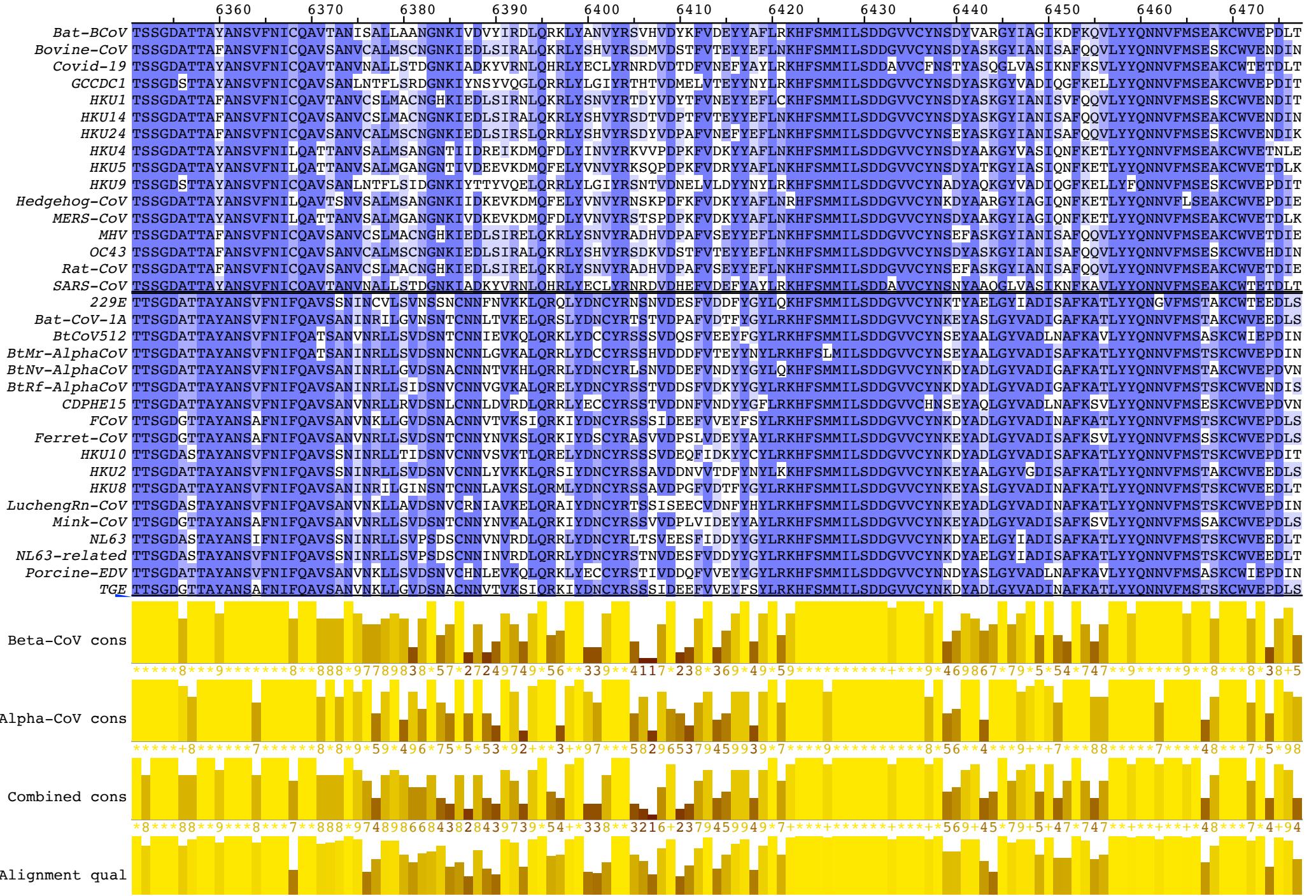


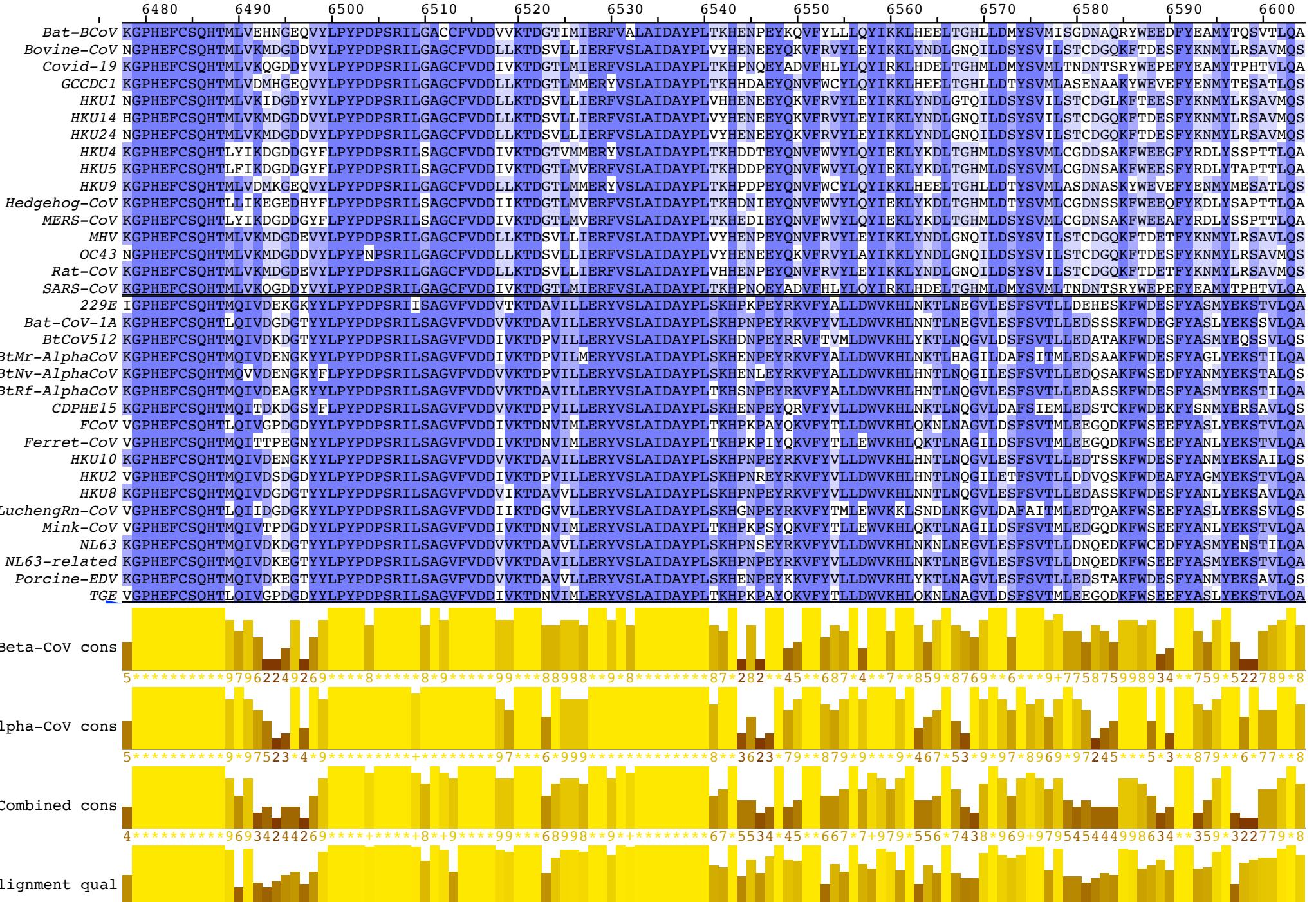


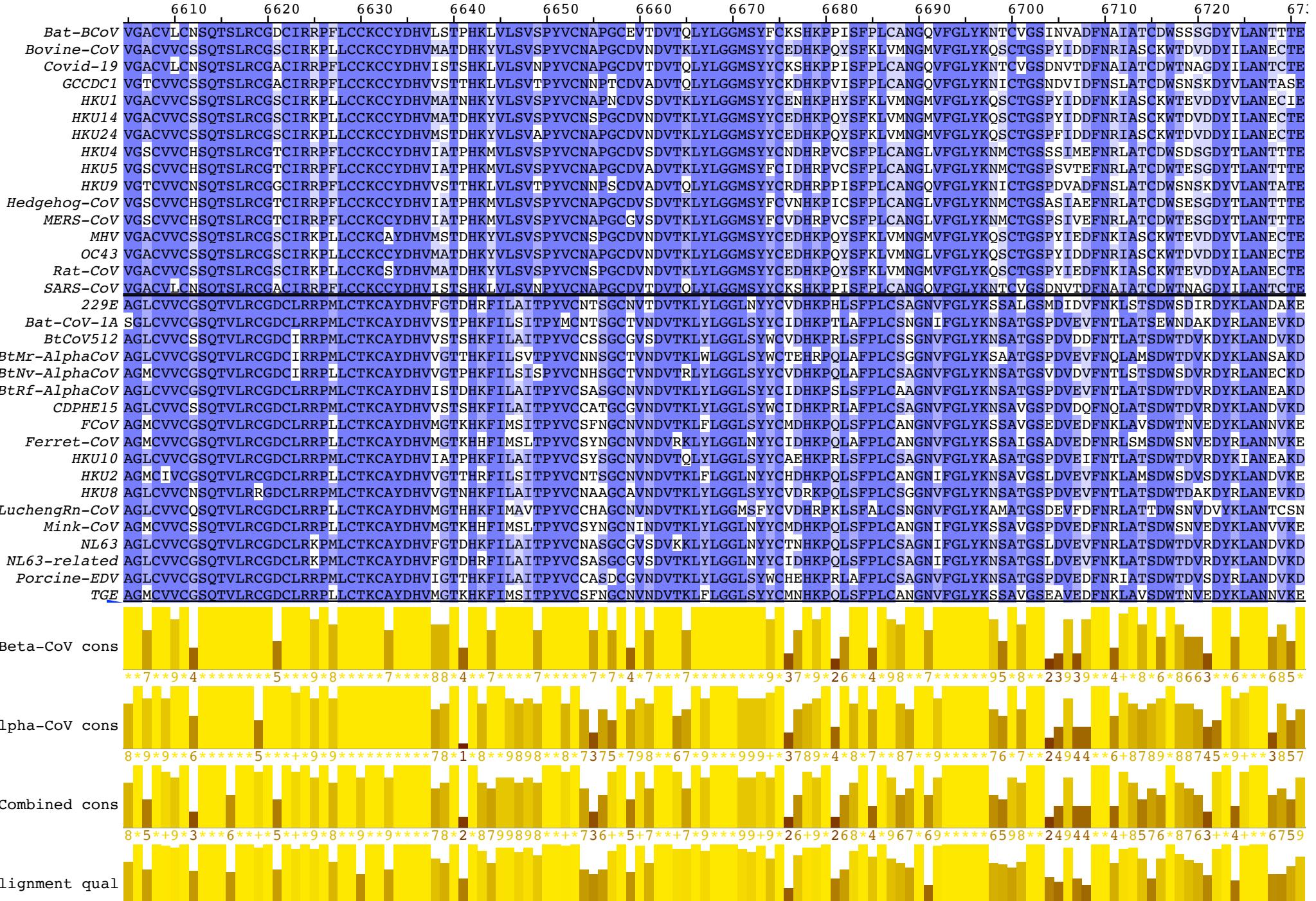


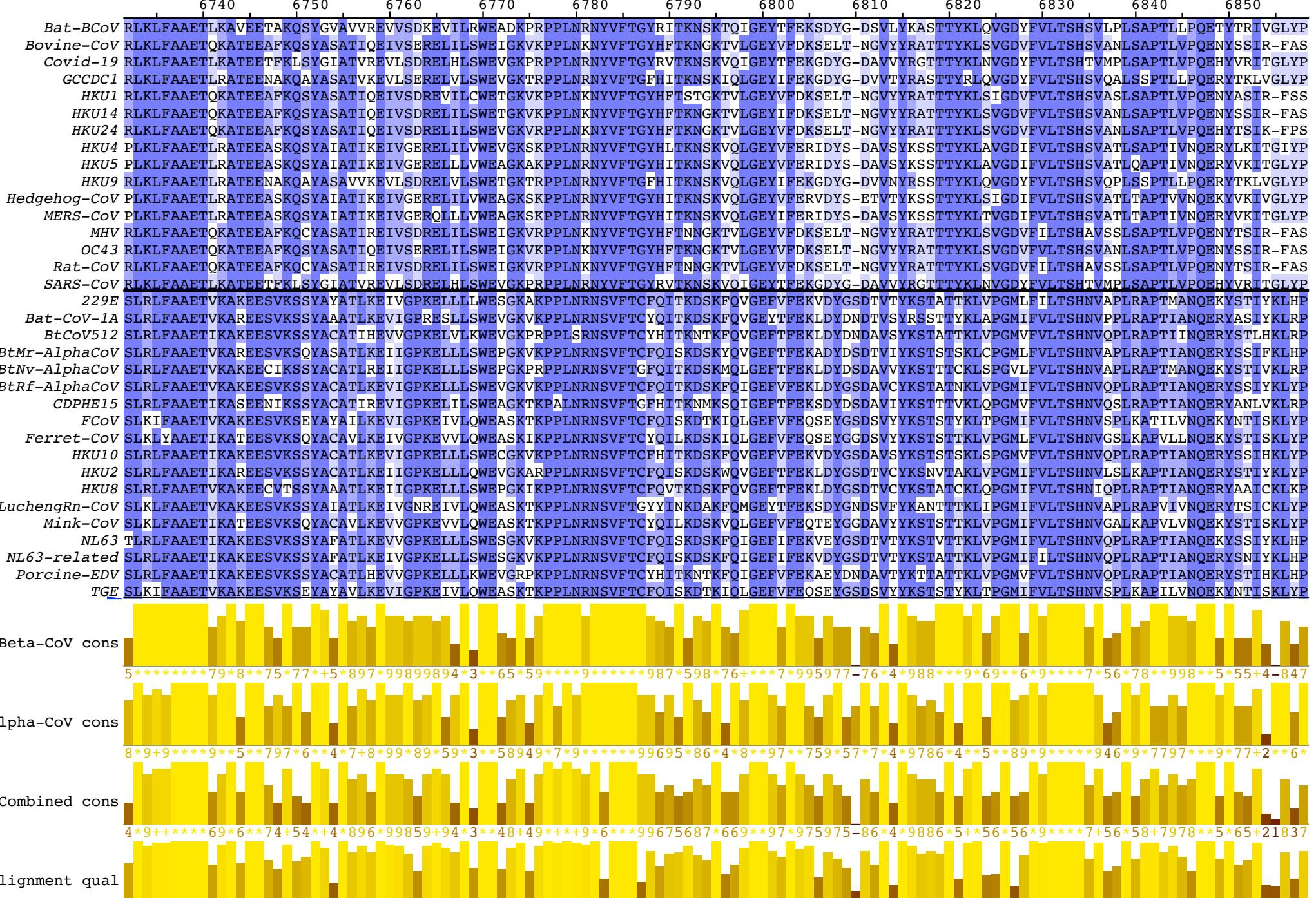


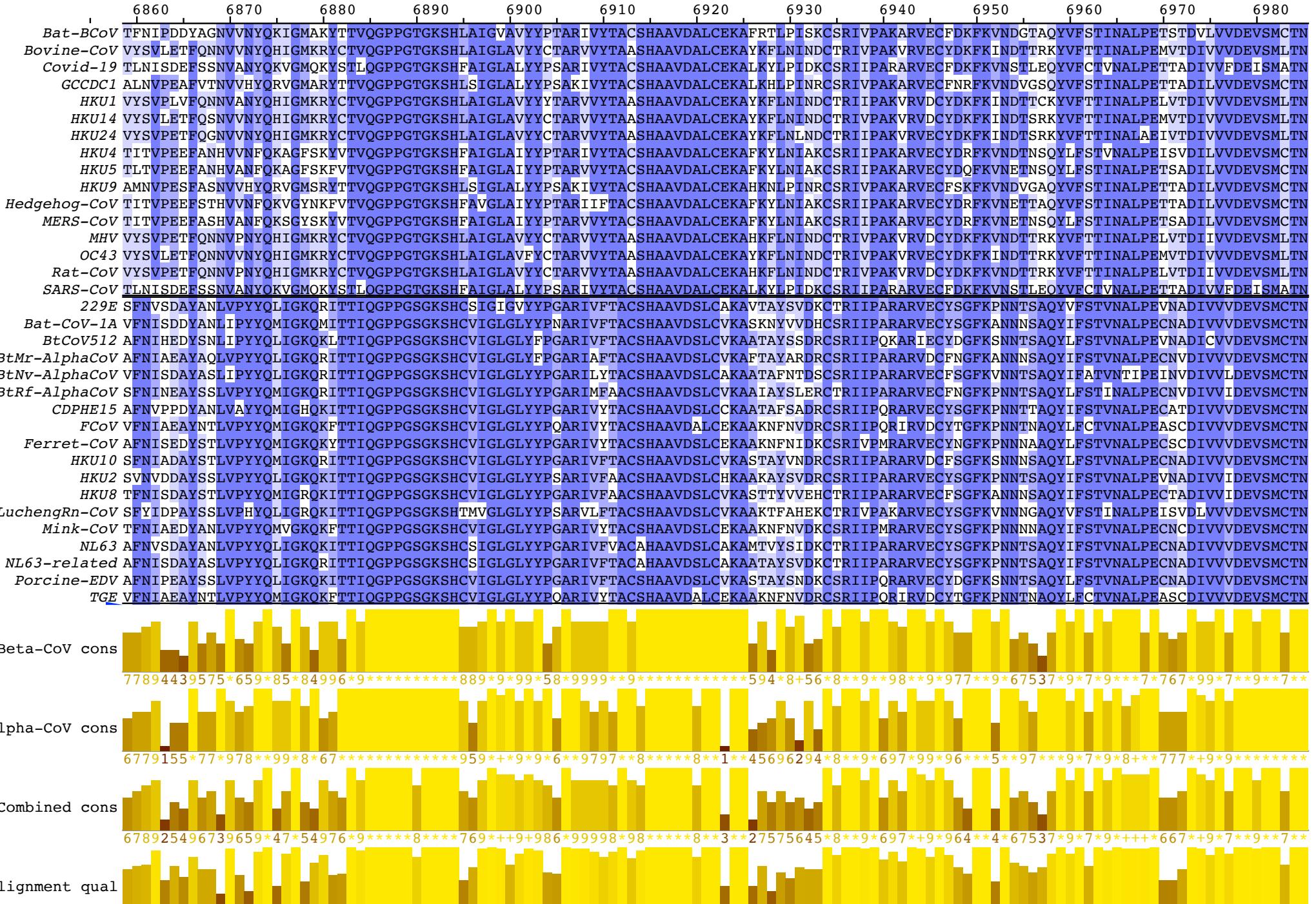


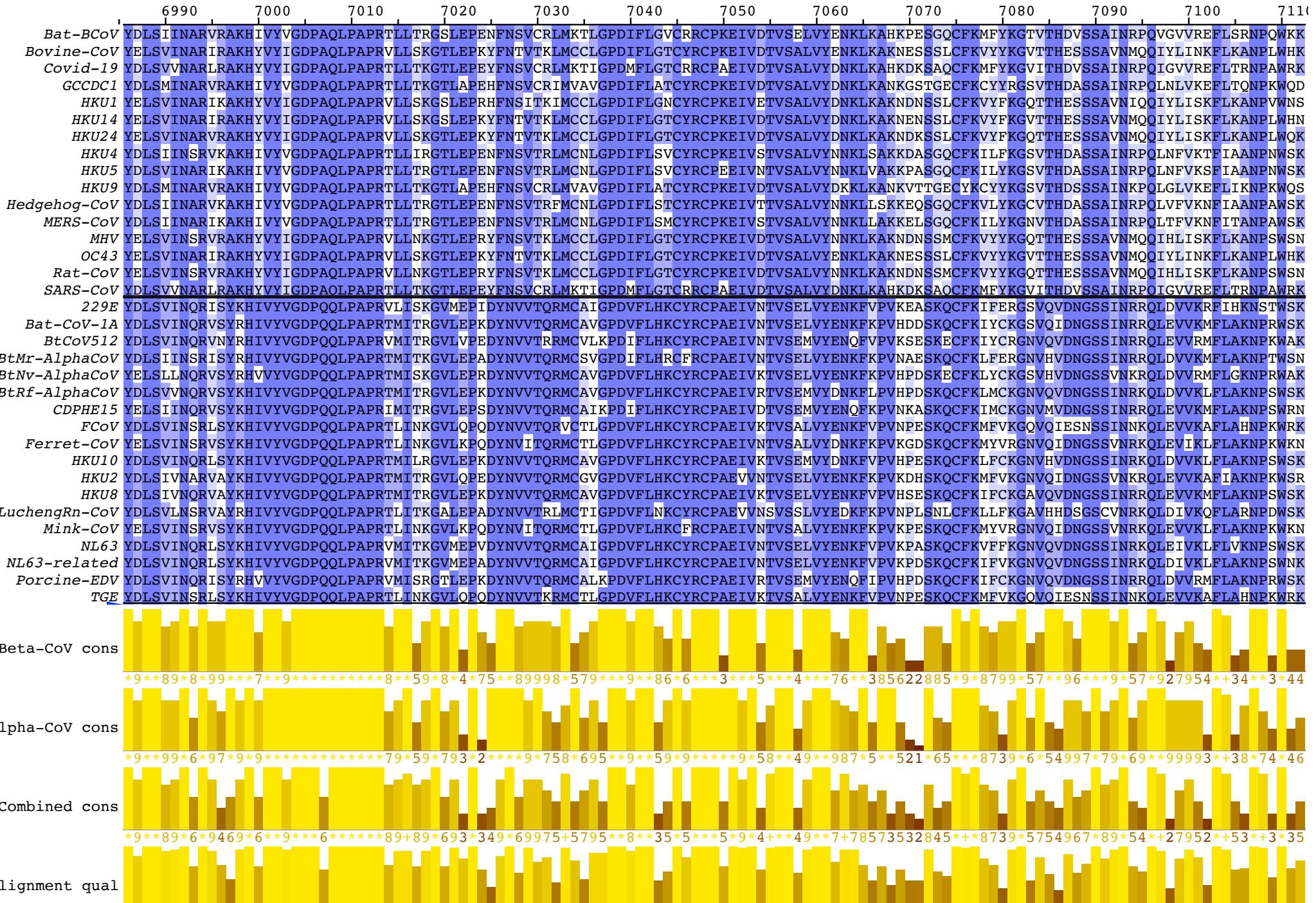


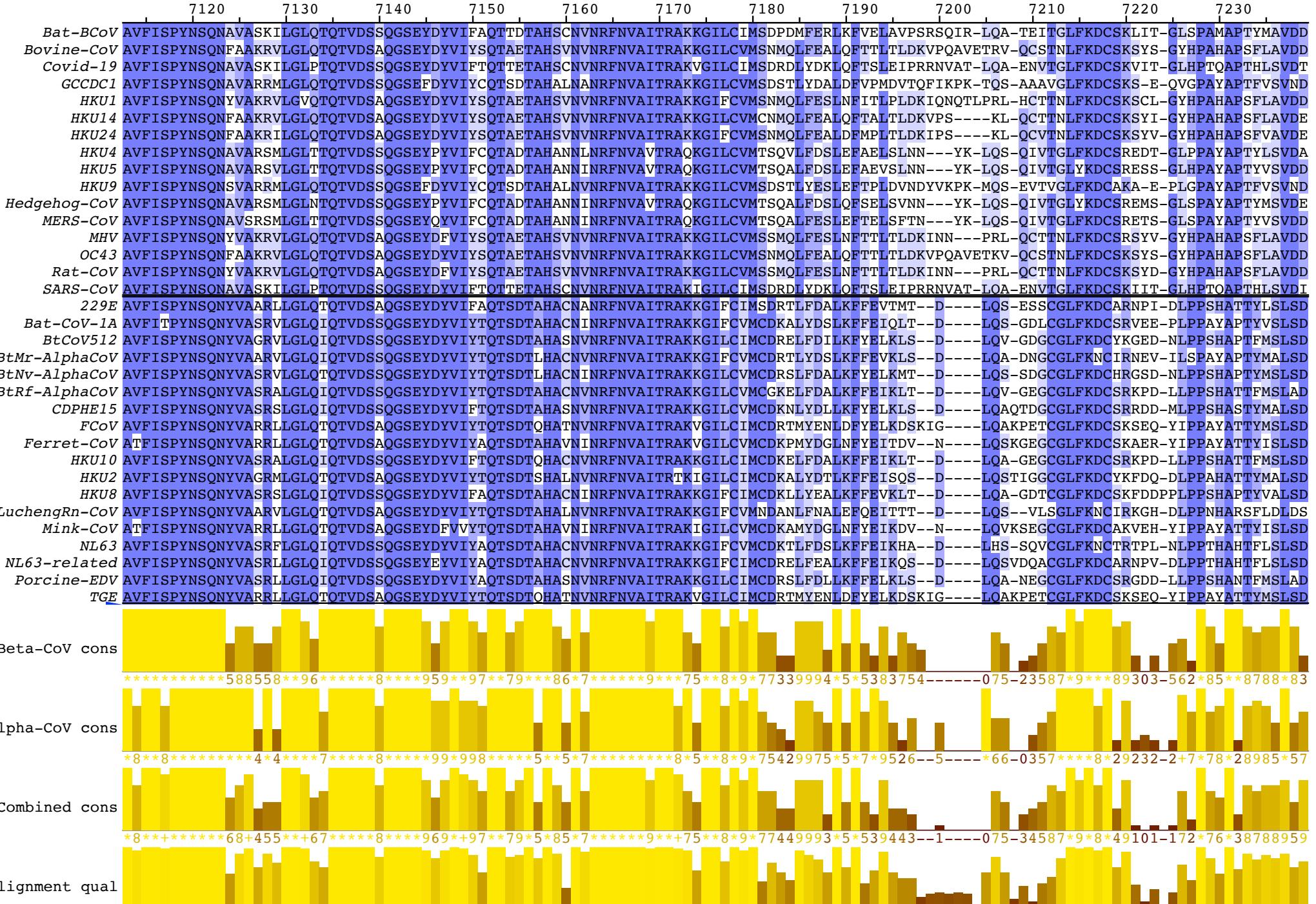


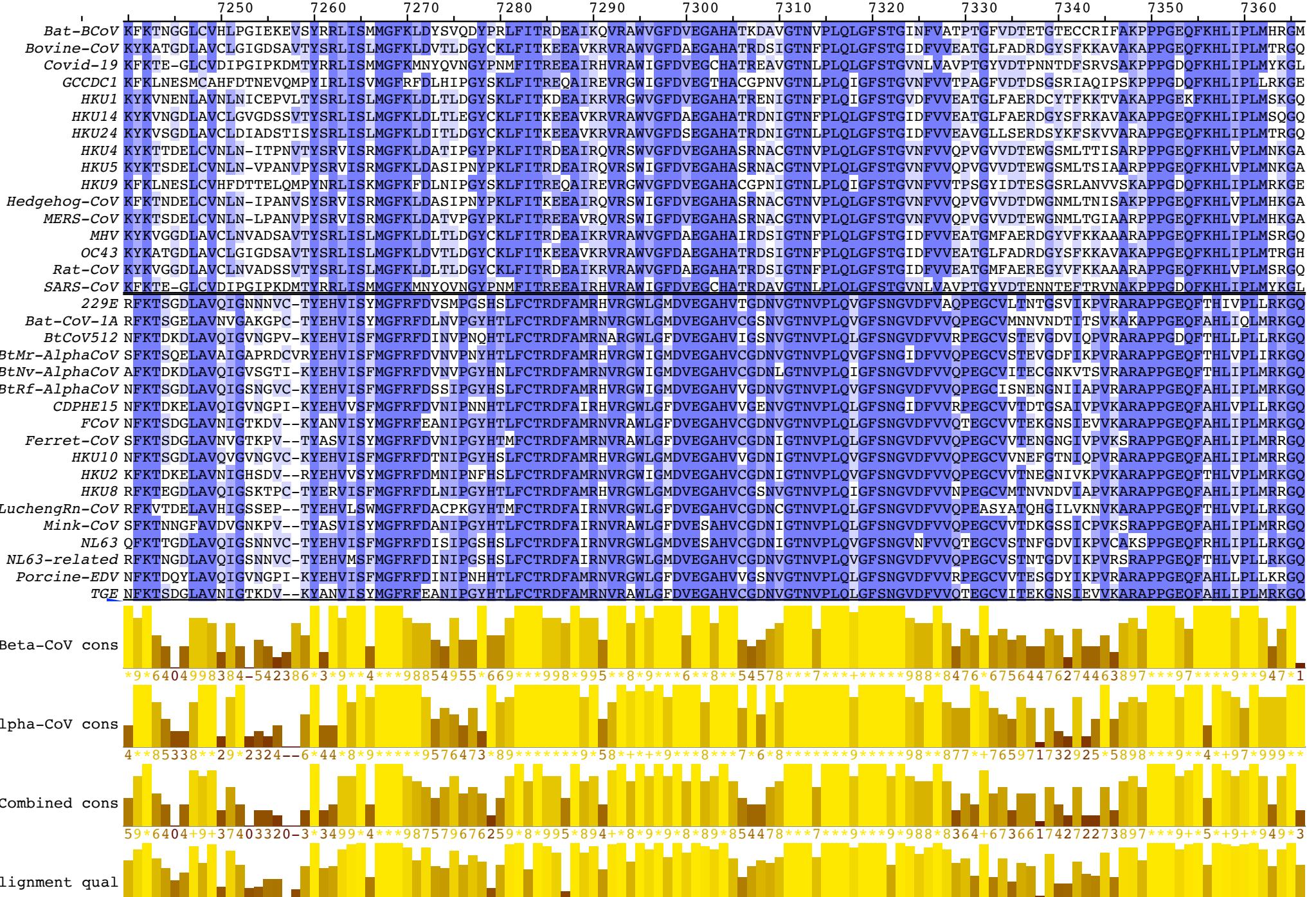


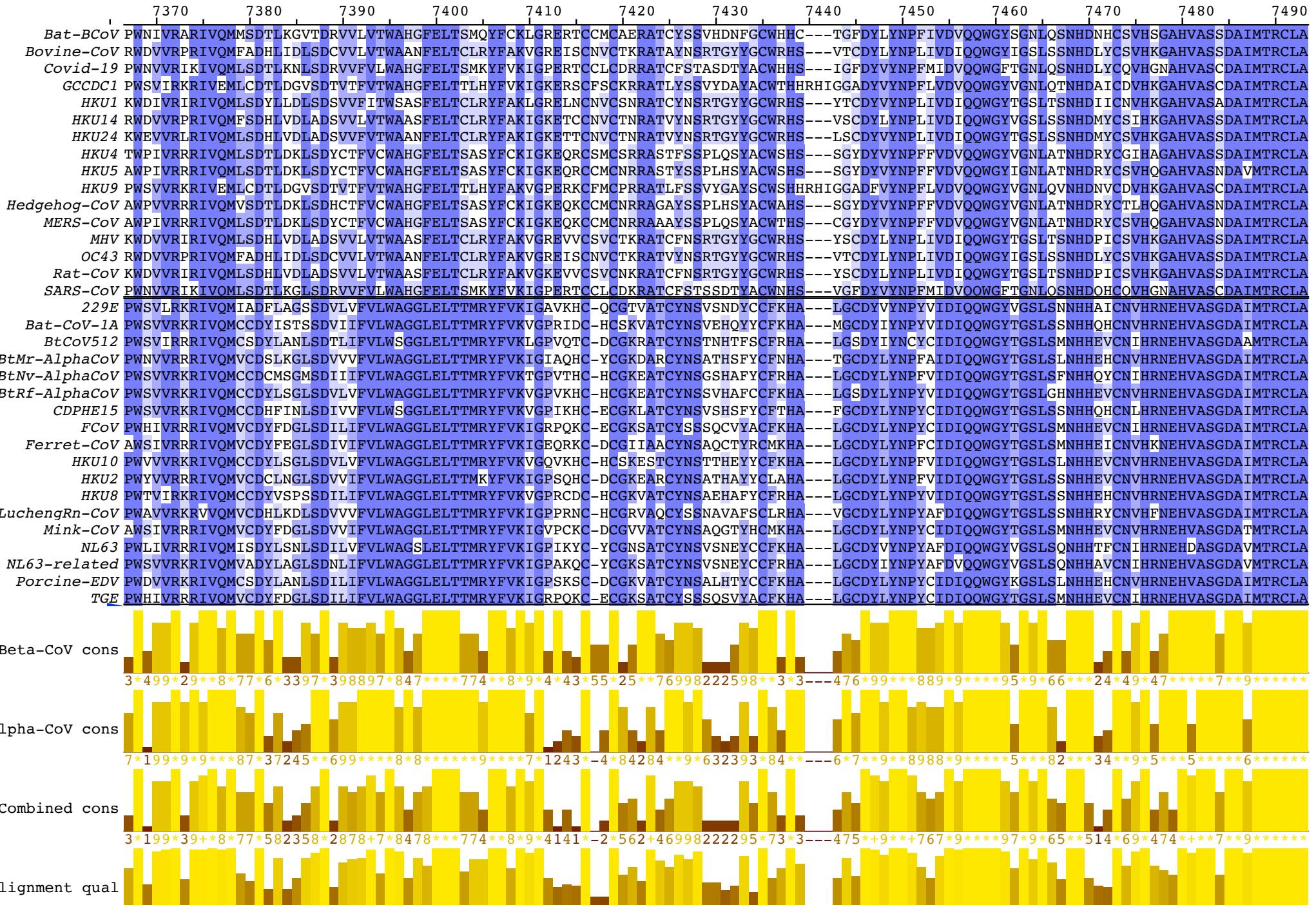


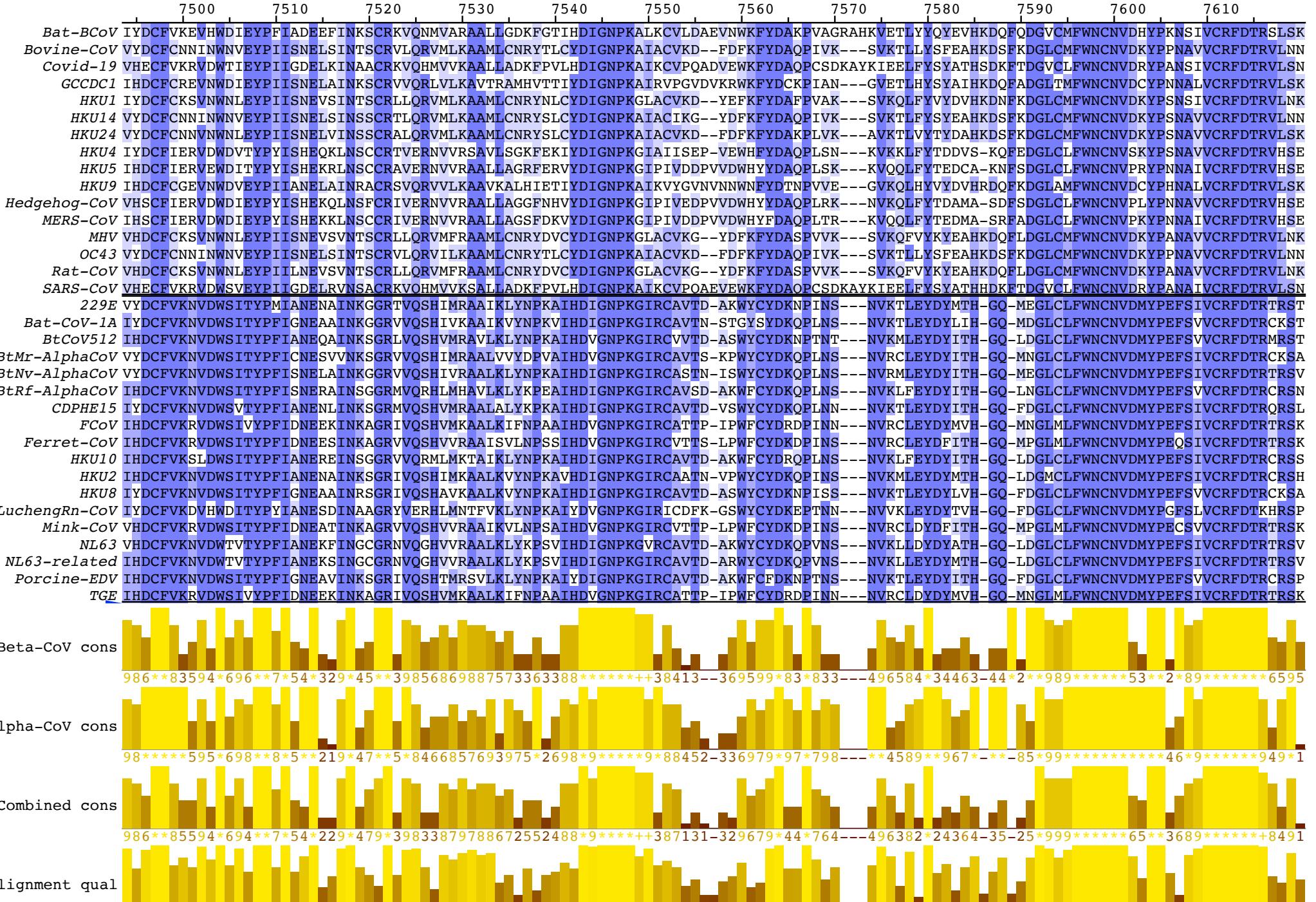


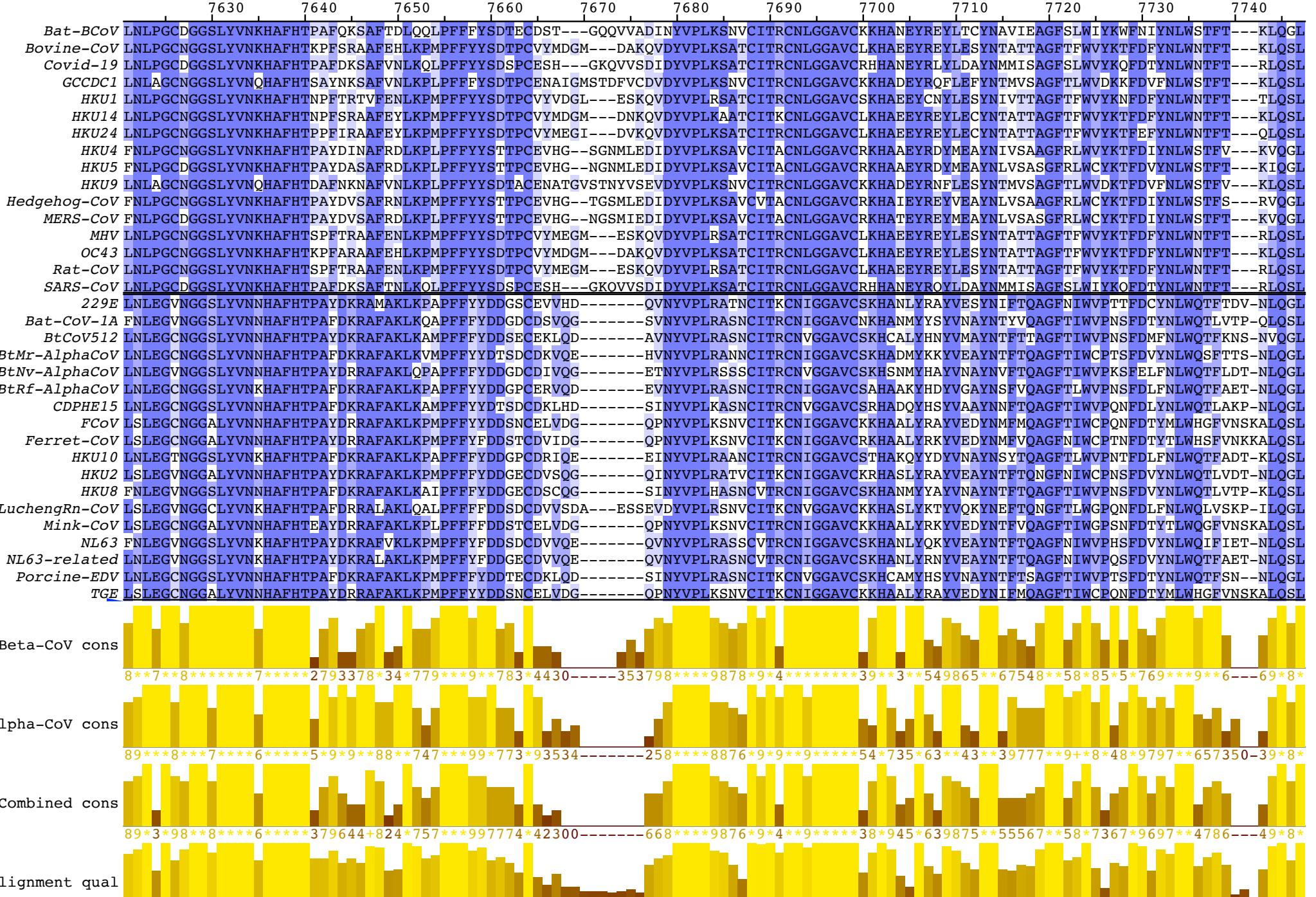


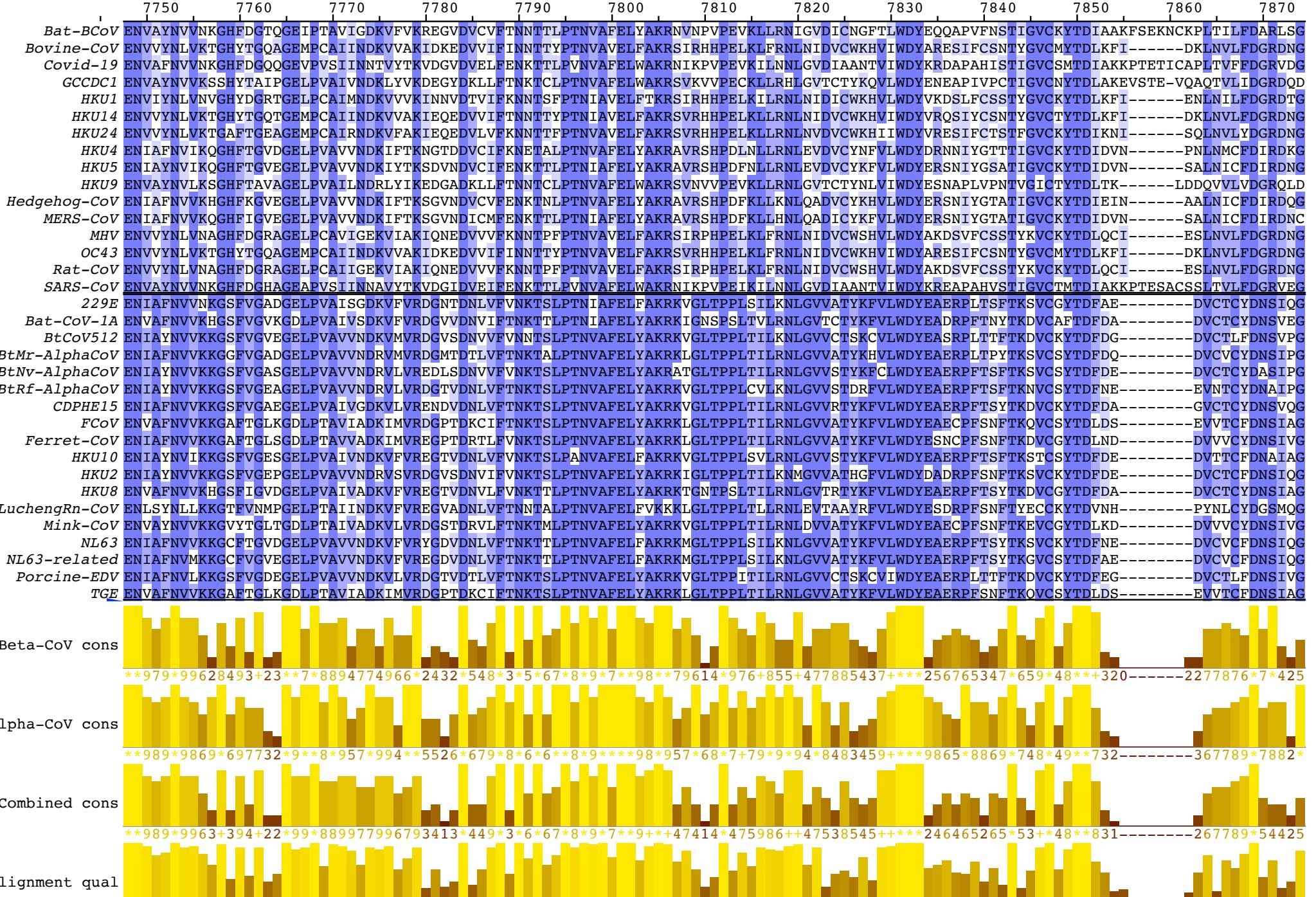


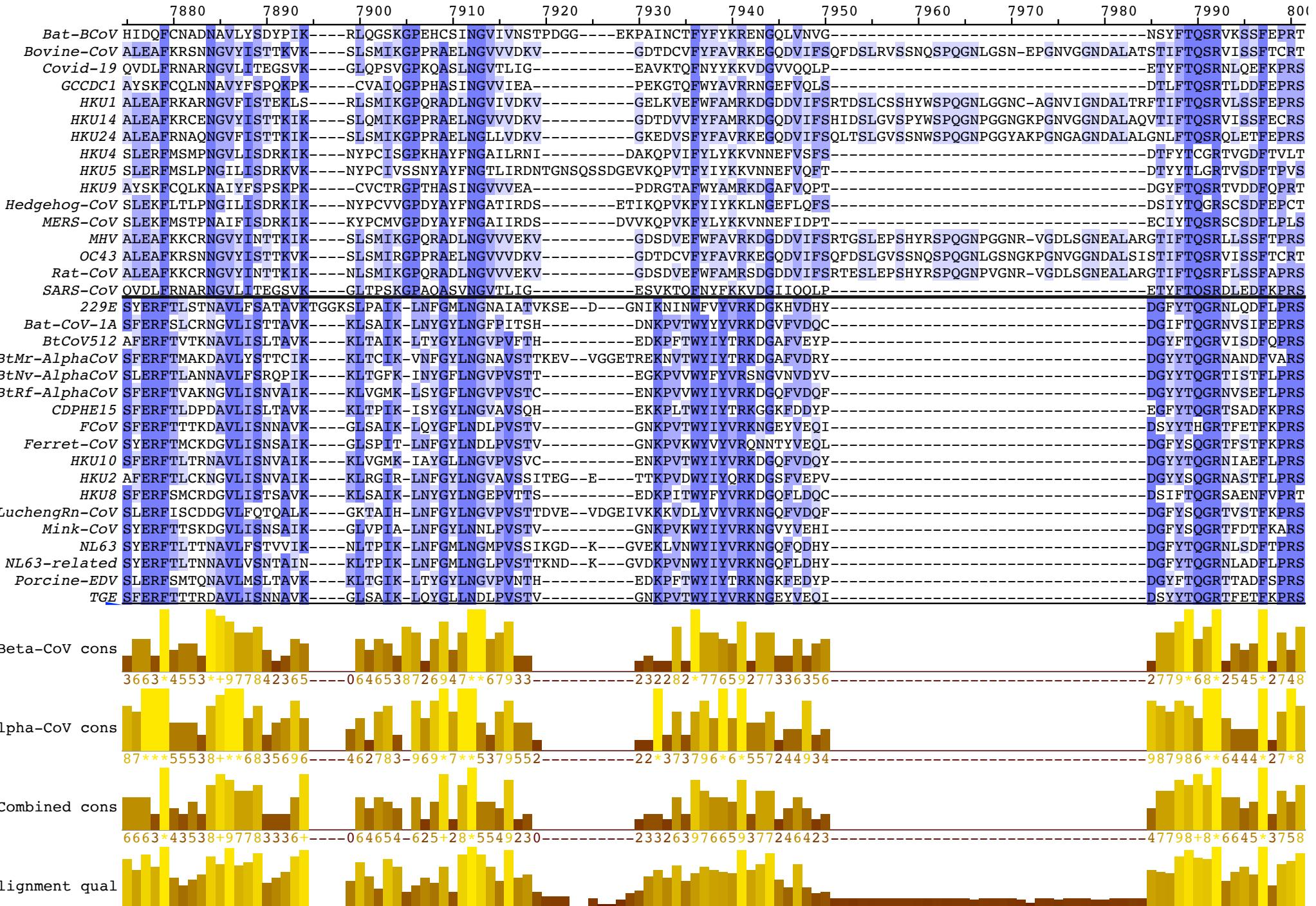


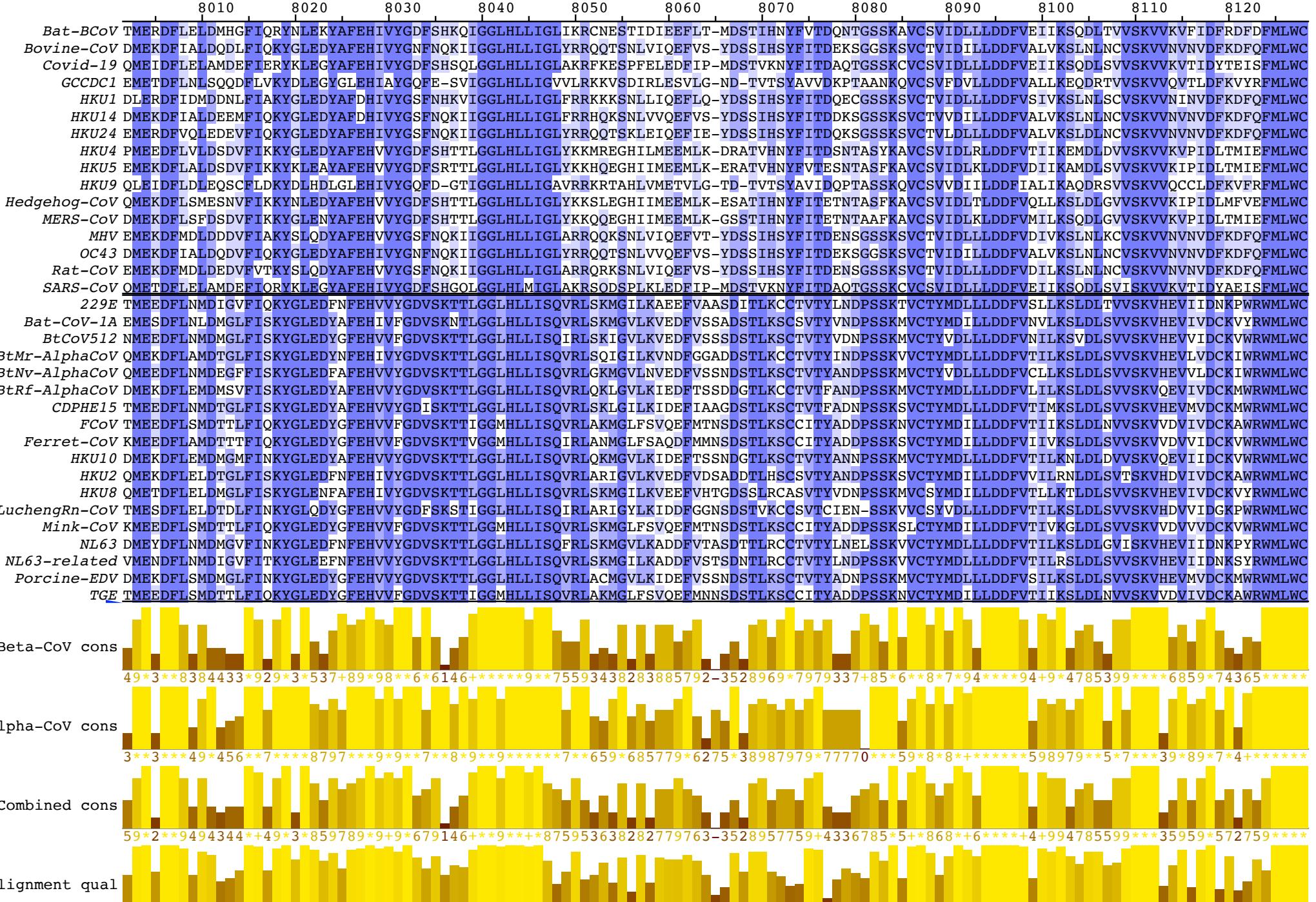


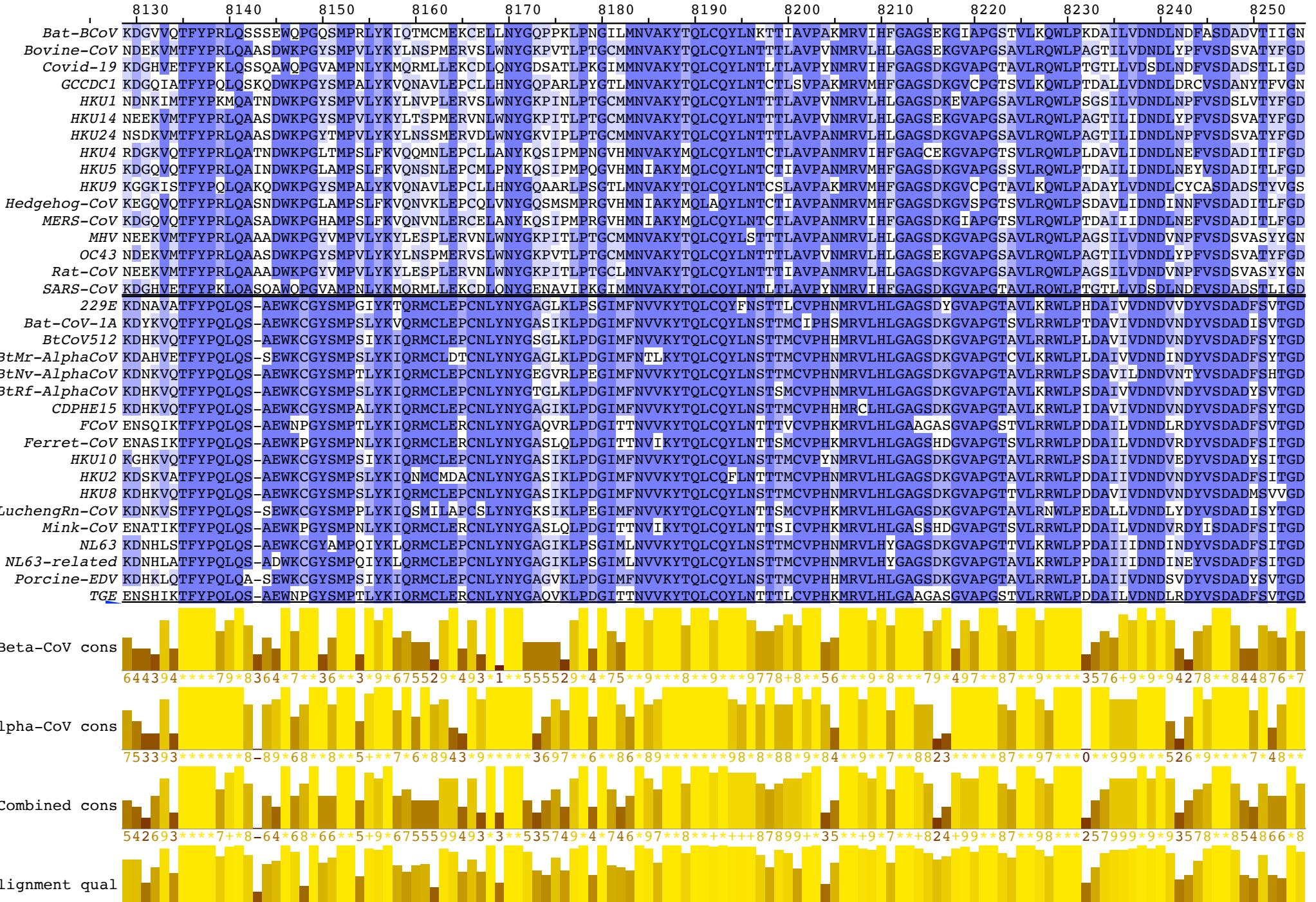


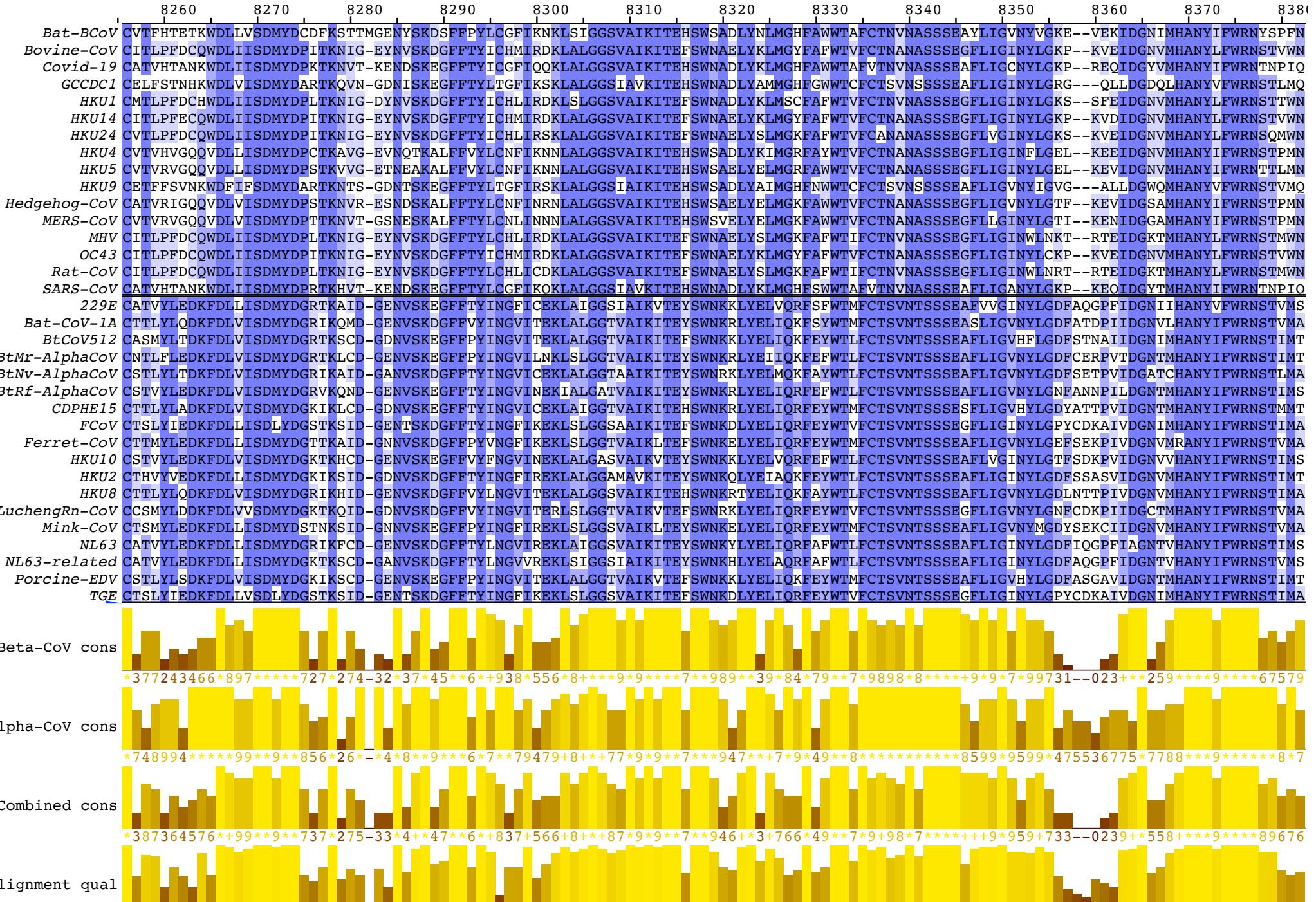


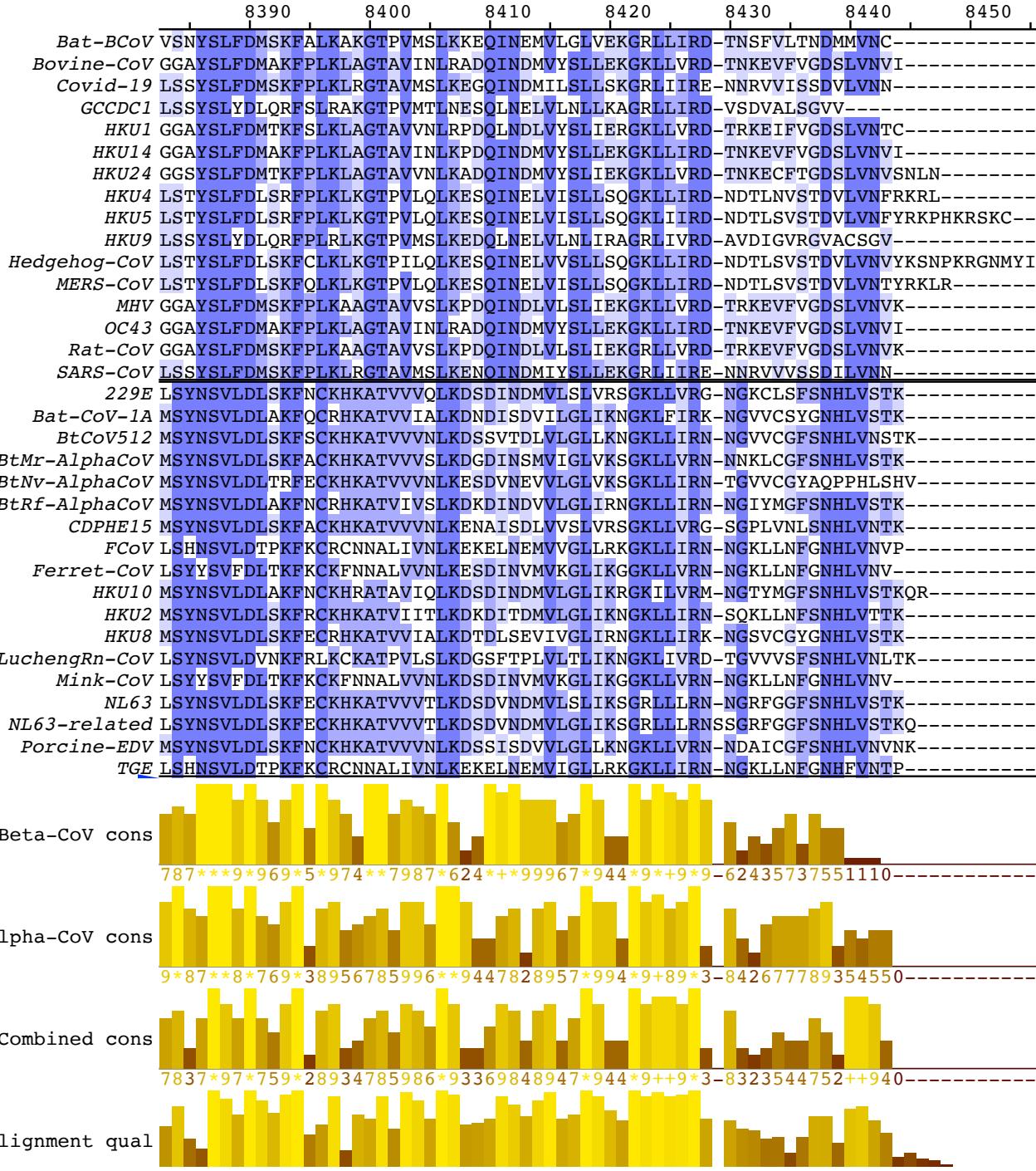






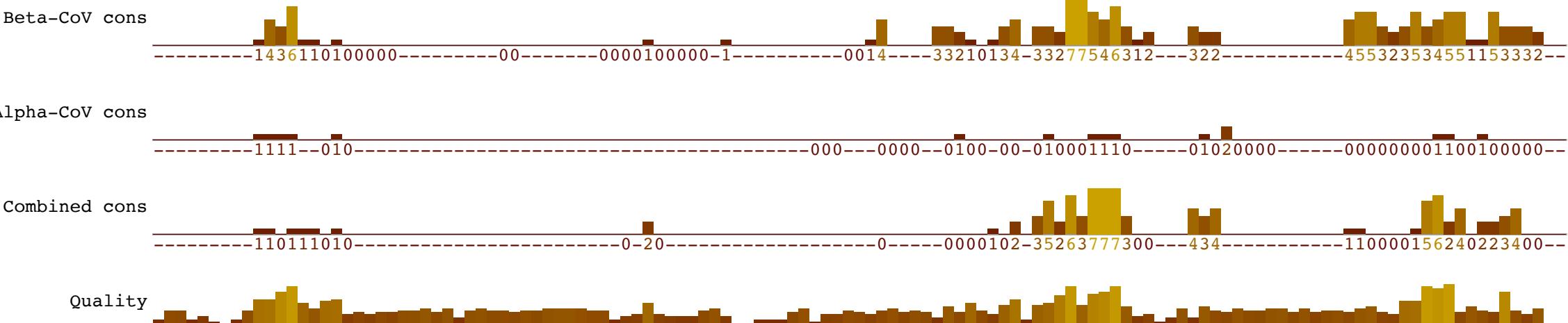
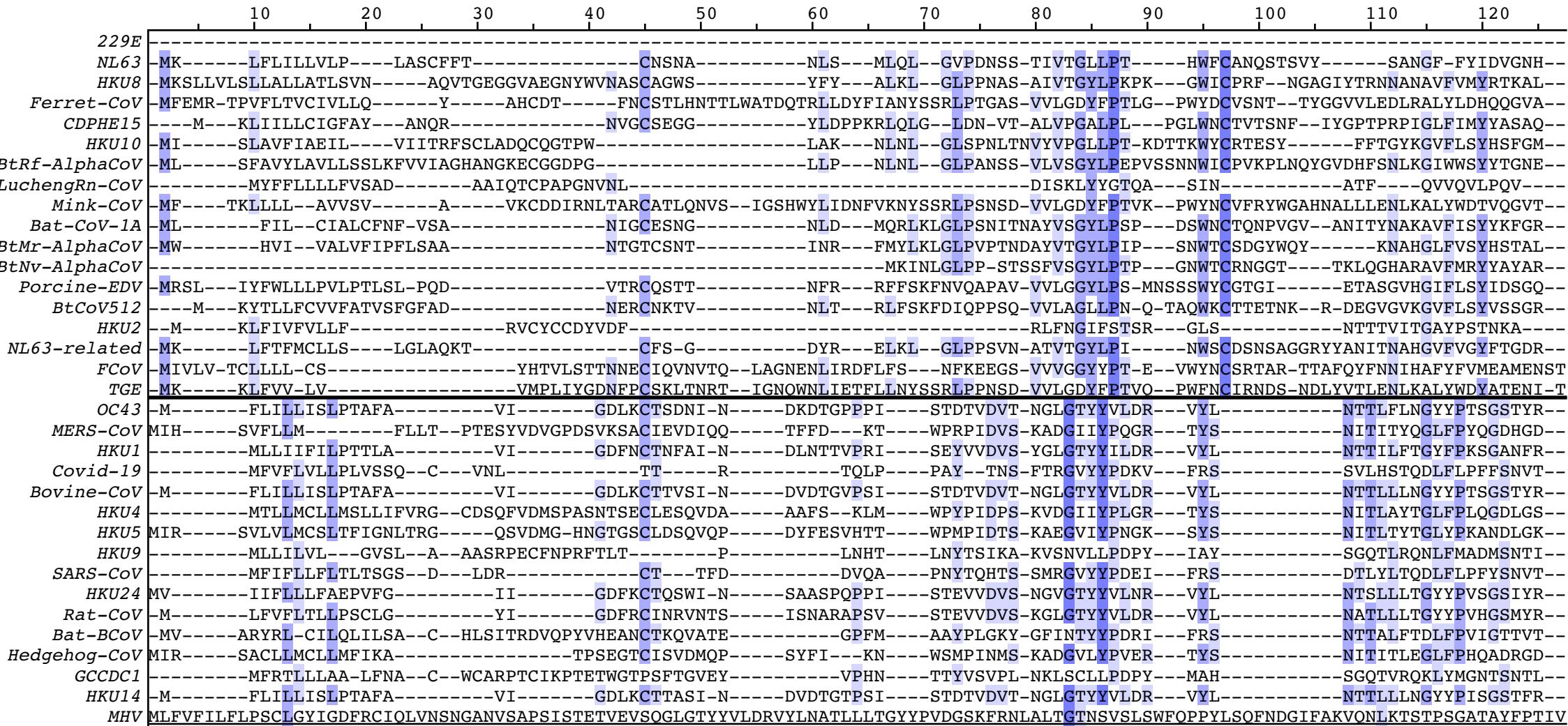


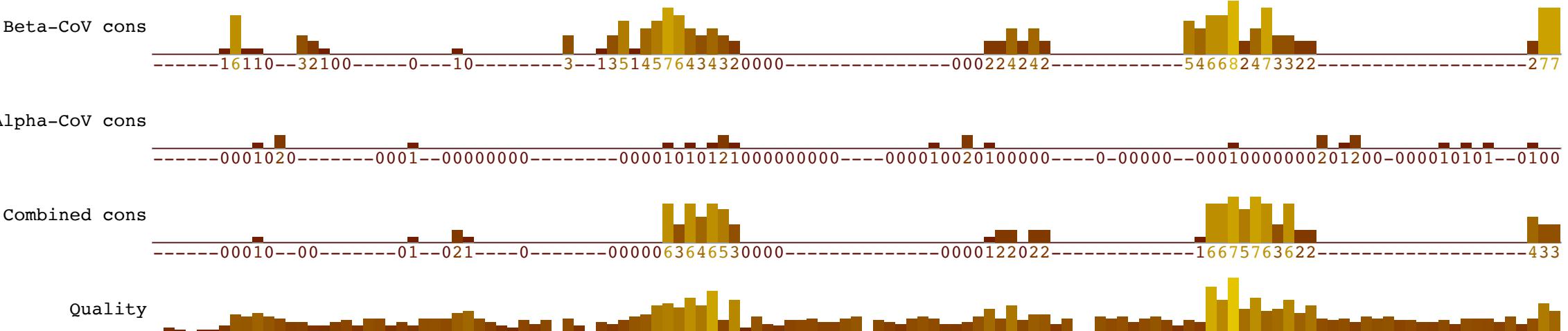
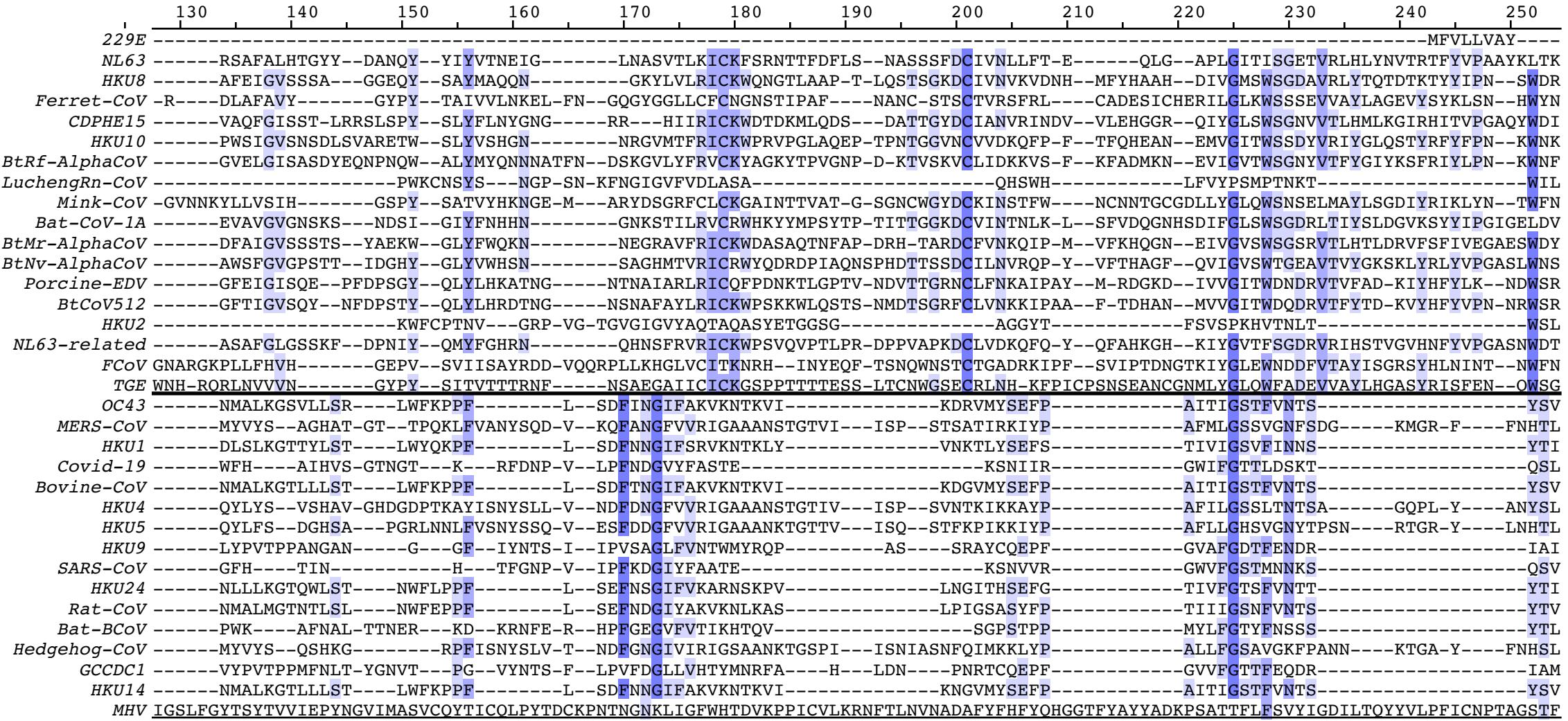




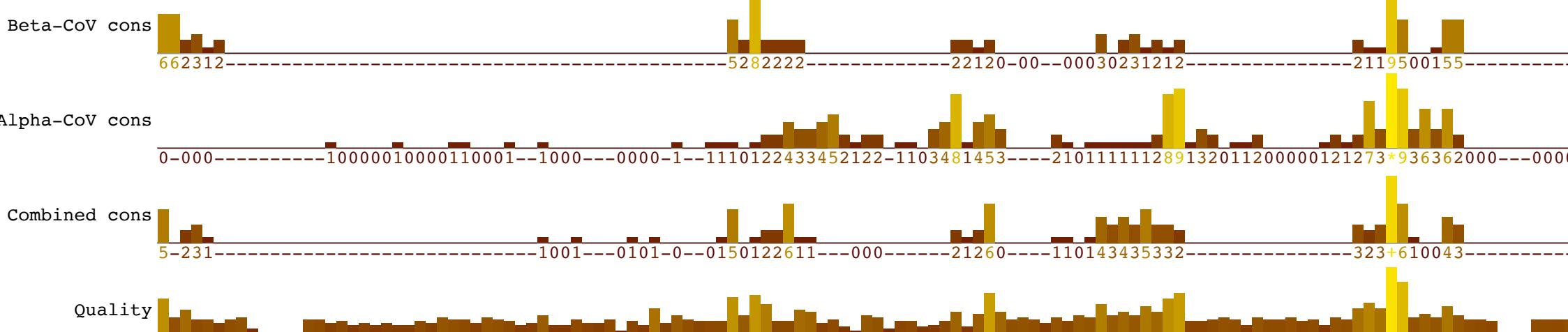
## Spike (S) Protein

SARS-CoV-2 1 ---MFVFLVLLPLV-S-----SOC---VN-L---TT---RTQLP---TNSFRGVYVYDDEPKFRSSRLSTQD  
 SARS-CoV 1 ---MFVFLVLLPLT-S-----GSD---LD-R---TCFTFDVVAQ---PNYQTHTSRRGVYVYDDEPKFRSSRLSTQD  
 HKU1 1 ---MLLIIIFILPT-TLAVIGDFNCTNFA---IN---D---LNTTVPRISEYVWVSYGLCTYVILDRVVLNTLTLFTGY  
 OC43 1 ---MFLLILISPLT-AFAVGDLKSLCSAN---IN---D---KDTGGPFPISTDVTWVTLNGLCTYVILDRVVLNTLTLFTGY  
 MERS-CoV 1 MIHSFVLLMLPLTPESTVWDVGPSDKSAC---IEVDI---QZTFDFKWT---PR-IDVSKAID---PQGRQTSNNTITYQGL  
 229E 1 ---MKLFLLIVLPLASCCF---TCNSANSLMSQLQGVDPDNSSITVGLLPTHWFC-ANQSTSWSANGCFP---IDVGNRHSRAFALHGTG  
 NL63 1 ---MKLFLLIVLPLASCCF---TCNSANSLMSQLQGVDPDNSSITVGLLPTHWFC-ANQSTSWSANGCFP---IDVGNRHSRAFALHGTG  
 90 100 110 120 130 140 150 160 170  
 SARS-CoV-2 55 FLPPFSNVT---WFHAIHVSGTNGTKR---FDNP-VLP---NDGVYFAST---E---KSNIIRGWIFGTT  
 SARS-CoV 59 FLPPFSNVT---GFHTIN---HT---FGNP-VIPFKDGYFAAT---E---KSNNVRGVWFGFST  
 HKU1 67 FKPGSANFRDL---SLKGTTYLST---LWYQKPFPLSDENGNIFSRVK-N---TKLYVNKTL---YSEFTSVIYGSV  
 OC43 68 YPTSGSTYRN---ALKGSVLSR---LWFKPFPLSDEINGIFSRVK-N---TKLYVNKTL---YSEFTSVIYGSV  
 MERS-CoV 75 FQYGDGHDMYVYSAHAGTGTQPKLVFVANSQD-VKQ-EANGFVWRIG-A---AANSTGTVIISPSATIRKTYIPAFMLGSS  
 229E 1 ---MFVLLVA  
 NL63 1 82 YDAN---QYYIYTNEIGLNASVTLKI---CKFSRNTTDFLSNASSSFDCIVNLLFTEQLGAPLGITISGETVRLHLYNVTTFYVPA  
 180 190 200 210 220 230 240 250 260  
 SARS-CoV-2 110 LDSKTQ---SLLI---VNN-A---TNVVIKVE  
 SARS-CoV 107 MNKQS---SVII---INN-S---TNVVIKVE  
 HKU1 130 FINNSY---TIVV---OPH---NGVLEITACO  
 OC43 131 FVNTSY---SVVV---OPRTINS-TODGDNKLQGLLEVSVO  
 MERS-CoV 153 VGNPSDGKMGRRFFNHTLVL---LPD-G---CGTLLRAF  
 229E 8 Y---A---LLHIAGCQTING---TNTSHSVNCNGVGHSENFAVESEGGYIIPSNFSFNWFLLTNTSSVVDGV  
 NL63 166 8 Y---A---LLHIAGCQTING---TNTSHSVNCNGVGHSENFAVESEGGYIIPSNFSFNWFLLTNTSSVVDGV  
 270 280 290 300 310 320 330 340 350  
 SARS-CoV-2 133 ---FQFNDPFLGVVYHKNN---KSWMESEFPRVYSSANNCFTFVSOPLFLMDLEGKOGFKNKLREFV  
 SARS-CoV 130 ---FELCDNPFFAVSKPMGT---QTHTMIFDNPANCFTFEMISDAFSLDVSEKSGNFKNKLREFV  
 HKU1 153 ---YTMCEYPHTICKS-KGSSRNESWHDFKEPLC---LFLKKNFTYTNVS-T-DWL-YFH  
 OC43 166 ---YNNCEQPYTCIHPNLNHRKELWHLDTGVSC---LYRKRNFTYDN-A-DYL-YFH  
 MERS-CoV 184 ---YCILERPSGNHGNPAGNSYTSFATHFTATCSDGNYNRRNASLNSFKEYP-NLRCNTMFTYNTIDEILEWFGITQZQV  
 229E 71 VRSQFPLLNLCLWSVGSQFTTGFVYNN---GTGRG-ACK-G-FYS  
 NL63 249 SRLYQPLRLTCLWVPGLKSTGTGVYNN---ATGSDVNCN-G-YQH  
 360 370 380 390 400 410 420 430 440  
 SARS-CoV-2 194 KNIDGYFKIISPLVNLDRLPQGFSALEPLVLDLPIGINITRFQ---LALHRSYL  
 SARS-CoV 187 FKNDGFLFVKGYQDFIVDRDPLSPFNTLKPFLPLGJINIFNRA  
 HKU1 202 FYQERGIFTYAYADS---GMP-TTFPLSFLYLTGLSHYYV  
 OC43 216 FYQEGIFTYAYFTDT---GVV-TKFPLFNVYMLAHSHYYV  
 MERS-CoV 226 HLFSSRYVLLDLYGN---MFOPATPLVYDITKYSI  
 229E 134 EKTYSGAVVYCTNNTLVLSG---DAHIPSCTVLGNFVCFVNNTIGNETTSASFVAFKTRVREFVSRTHFYINGVRYFS  
 NL63 317 EKTLQYDVLFCNSSSGVL---DTTIPFGPSSQPYCFCINSTINTHVSFTVGLLPPVREIVVARTGQFYINGFKYFD  
 450 460 470 480 490 500 510 520 530  
 SARS-CoV-2 251 PGDSS---SGWTAGAAYYNGYQOPRTFLLKYNEGTITDAVDACLDPLSETCTKLKSFTEKIVYQTSNFRVQTESIVRPPNIT  
 SARS-CoV 240 ---PAQ---DIWTSAAAYFNGYKTPFMFLKDYDGTITDAVDCSQNPLAEKCSVSEIDKGIYOTSFRVQESGDFVRPNTNL  
 HKU1 244 ---AIS---SNTDNTLQYVPLSKRQYLLDFTGDRVYIATNAVDCSSSFSEIICKTLLKPNPFTGVYDLSCTKVATVTHRIPDLD  
 OC43 258 ---S---KLTLEWVMTPTBSQYLLNAPDQGCI-NLNAVDCMSDFMSIEKCTQSIAPPTFCVYELNGYVTOIADYVYRKKRNLPN  
 MERS-CoV 303 IQSDR---KAW---AAFYWKQQLPFLLDFPSVGDYIIRRRAIDCGFNDLSQLHCYESFDSVDESEGVYVSSEPEAKSGSVVQEQAEG-VE  
 229E 211 LDGEVANFPNVTNAA-TIVCCTAASYADVYDNVSQTALENANITYCN-SVINKLRLQDSDLEPDVFYTSFSTIPQVPLPMSIVSPV  
 NL63 394 LFGFEAVNFTNTASATDFTWVTFDVLVNVSATNQIYLLDQD-SPEFKEKLCQEHLOGLQDFYSANFLDDNLVPETYVALPI  
 540 550 560 570 580 590 600 610 620  
 SARS-CoV-2 336 CPEGEVNFATRFSVAVYNNRKRISNCVADYSVLNNSASFSTKVGSPVTLNLDLCTKNTVYADS---VIRGDEVRIQAPGOTKIAIDYNYK  
 SARS-CoV 323 CPEGEVNFATRFPVSYAVWERKNCVADYSVLNNSASFSTKVGSPVTLNLDLCTKNTVYADS---VIRGDEVRIQAPGOTKIAIDYNYK  
 HKU1 327 CIDKWLNNFNPVSPNLRWERKJFISNCNFDLSTLRLVHTDTSFCNCNFDESKYIYGCFSKISVLDKFAIPNSRSDLQLGSSGFLQSYNSK  
 OC43 335 CHIEANLNDKSVPSPLNWERKTFNSCNFMSSLMFIQADSFCTNNDIAKYGMCFSSTIDKAIAPNGRKLVDOLNLGYLQOFNSY  
 MERS-CoV 383 CDFSPLLSG-TPPQVNFVLRKFTWCNLYNCKRFTSQCISPAIAANCSYSSLILYVSPYLSMKSDLSVSSAGPISOFNYK  
 229E 295 ---Y---LHKTFIV---LHKVKEH-QRGPKGKYNCRPSVIN---ITLANNE-  
 NL63 479 ---Y---YQHTDIN---PTAT---ASFGGSYVCKPBHQVN---ISLNG-  
 630 640 650 660 670 680 690 700 710  
 SARS-CoV-2 475 LPDDFTGCVIAWNNSNLDSKVGNN---YNYLRLFRKS---NJKPFERDISTEIYQA---GSTPCNGV  
 SARS-CoV 412 LPDDFMCGVLAWNTRNIDATSTGN---YNYKXRYLHRG---KLRPFVSDNNSP-F-DGKPCPT---P-ALN  
 HKU1 416 IDTTSQSLQYLSPLSAPINTVNNYNPSPSWNRRYGPNNF---NLSSHSFVSDNNSP-F-SCKSHKPPAS  
 OC43 424 IDTTATSCOLYNNPAANVSPRNPSTWNKRFQFIEDSFVFKPRPAGVLTNDHVVYAAQCHFKAQPNFCKVLDOLNLGYLQOFNSY  
 MERS-CoV 303 IQSDR---KAW---AAFYWKQQLPFLLDFPSVGDYIIRRRAIDCGFNDLSQLHCYESFDSVDESEGVYVSSEPEAKSGSVVQEQAEG-VE  
 229E 211 LDGEVANFPNVTNAA-TIVCCTAASYADVYDNVSQTALENANITYCN-SVINKLRLQDSDLEPDVFYTSFSTIPQVPLPMSIVSPV  
 NL63 394 LFGFEAVNFTNTASATDFTWVTFDVLVNVSATNQIYLLDQD-SPEFKEKLCQEHLOGLQDFYSANFLDDNLVPETYVALPI  
 720 730 740 750 760 770 780 790 800  
 SARS-CoV-2 488 ---YF---PLQSYGFQPT---NGVQYQPY---RVVVLSP  
 SARS-CoV 474 C---PLNDYGFYTT---TGICQYQ---RVVVLSP  
 HKU1 495 CPGITNYRSCESTVLDHTDNCRLSCLPDITAYPDRPSCSQKSLVVGVCHEAGCAGFVGDDEEKCGVLGDLGSYNSVNCCLSCTDAFLWVSYDTCV  
 OC43 513 PAGTNYLTCD---NLCTPDITFTGTYKCPQTKSLVGLIGEHCSCLAVKSDYCGG---NSCTCRPQAFLWVSYDADSC  
 MERS-CoV 534 VWEQGDYVYRQL---SLEGGWLV---SCTVAMT---EQIQMFG  
 229E 353 ---V---KLRWSASINTG-NC---PFSFGKVNRF-VKFGSVCFSL---KDIPG  
 NL63 530 ---RVKSGPGDSSWHLKSG-TC---PFSFSKNNF-QKFTKICEST---VEVPGS  
 810 820 830 840 850 860 870 880  
 SARS-CoV-2 516 ---ELL---HAPATVCGP---KKSTNLVKNQCVNFNFGLIGTGVLTESNKKEL-PFQFGRDIA-DTT-DAVRDPOTLEI  
 SARS-CoV 502 ---ELL---NAPATVCGP---KLSTDLIKNCVNFNFGLIGTGVLPSSKRFQ-PFQFGRDVS-DPT-DSVDRDPKTSEI  
 HKU1 584 SNRNCRNIFSNFIINGTSCTN---DLQPNPTEVFTDVLVCDYDLYGIGTQKIFKEVSAVYYNSQWNLLYDSN-GNII-GFKDFDVTNK  
 OC43 584 QGDCKNIFANIFIHLDVNSGCLCT---DLQKANTD1LICVNCVNDLYLQIGCIGEVFVNATYNSWNLLYDSN-GNII-GFKDFDVTNK  
 MERS-CoV 572 ---GTVQYGDTDNSVCPKLEFANDTKIASLQNCVNEVSLXGVSQFCVNFONTAVGV-RORFVEYDQVY-OLVQGYSD---DGNY  
 229E 424 TGV---PKPV---EGVFSMNTLNKTCYKINIDVMSGVVQIYRISNDTFLNG---ITYSTGSLN-  
 NL63 607 TGV---PKPV---SGIREFSNLVNNCTKYNIDYVCTGIIRSSNQLLAGG---ITVVSNSGNL-LGFKDVTWNTG  
 900 910 920 930 940 950 960 970  
 SARS-CoV-2 585 LDITPCSFCCGVSVITPTGNTSNQAVYIYQDWNCTEVPAVIAHDQLT---PTWVYVSTGNSVQTRAGCLIGAEHVN-NSYEDCIFIGAG  
 SARS-CoV 571 LDITPCAFGGVSVITPTGNTSNQAVYIYQDWNCTEVPAVIAHDQLT---PAWRIYTCNNGVFTQAOGLIGAEHV---TYSVCDIPIGAC  
 HKU1 669 YNIFPPYAGVRSAAFH---ONASSLALLYRNLKCSVLMNIS---LTTQ---PYEDSYLGCVNADNLTYDSSVSCALMRG  
 OC43 669 FMRCYSGVRSAAFH---ANSSPEALLFRNCKVNFNNSLROLQPI---NEDSYLGCVNADNLTYDSSVSCALMRG  
 MERS-CoV 649 YKLRCAVSVPSVVIYQD---KETKTHATLFGSVACEHISSTMSQYRSRSTRSMLKRDSTYGLPTPVGCVLGLVNSN-LFVEDCKLPLGS  
 229E 490 YSITCPNPQDQLVYQD---AVVGALEMSENFVTSYGSNVVEMP---KTFYVASN---CTYNTCD---AVLYTSSFG  
 NL63 673 FIVTCNPQDQVAVYQD---SIIGAMTAVNESRGQNLLNQLQF---NFFYVSN---GNGNCTT---AVMVTSSNG  
 980 990 1000 1010 1020 1030 1040 1050 1060  
 SARS-CoV-2 670 IGAQYQTOT-NSPRRARSVAQ-SI---I-AYTMSLGAENSV---AYSNNSIAIPLNTISVITVPLVPMSTKTSVDCNTMIGDOSTECS  
 SARS-CoV 655 ICASYHTVS-L---LRTSKQ-SI---V-AYTMSLGAQDSSI---AYSNNTIAIPLNTISVITVPLVPMSTKTSVDCNTMIGDOSTECS  
 HKU1 743 FQDVYNSPSSSSRRKRSRISASASYRFTVPEPPNVSF-VNDSDIESVGGYIIEQIPTNTIVGQEEFQIQTNSPKVITDCSLPVCSNYYAACH  
 OC43 746 FQDVYDK---NRSRGAITTYGTRNFPPEVNTVPEPPNVSF-VNDSDIESVGGYIIEQIPTNTIVGQEEFQIQTNSPKVITDCSLPVCSNYYAACH  
 MERS-CoV 735 LCALDPSTLTPRSVSPG-EMLRAS-APNFIQV-DL-NSSYFPLSNTLMSFNGTSEVYQIETQIQTICKVTDCKQVYVNGQFKQCE  
 229E 553 VCAUGDSIIA---VQPNRVSYD---SV-SAVIITANLISIISNTVSVQVYELQITSTPITVDCSTVYNGNQVRC  
 NL63 736 ICADGSLIIP---VPRFRNSN---GI-SAIITANLISIISNTVSVQVYELQITSTPITVDCSTVYNGNQVRC  
 1070 1080 1090 1100 1110 1120 1130 1140 1150  
 SARS-CoV-2 751 NLLQYGSFCQTLQNRALTGIAVEQVKQIYKTPPIKDPG---GF---FSQILB-DP---SKPSKREFIEDLLFNRVTLA  
 SARS-CoV 733 NLLQYGSFCQTLQNRALTGIAAEQDRNTRVEFVQKQIYKTPPIKDPG---GF---FSQILB-DP---LMLTKRPSKREFIEDLLFNRVTLA  
 HKU1 831 DILSYEFTGCDNISLTDLVEVNLDTTQLQVQNSLMNGVFTSLKLDGVNPNVFDINIFPSVLCGLGSCSKASSRAEIDLFLPKVLS  
 OC43 829 SOLVEYQGFCQDNNAIILTEVNLDTTQLQVQNSLMNGVFTSLKLDGVNPNVFDINIFPSVLCGLGSCSKASSRAEIDLFLPKVLS  
 MERS-CoV 819 QLLRVEYQGFCQSKINQHLGANRQODDSVRLNFAVSQKSSQSSPPIIPEGCF---GDFNLTLLVE-  
 229E 619 ELLQKQTSACKTIEDALRNSAMLESADMVSEMLTFTDKAFTLAVANSSF---GDNVLLSSVIPSRLRSGRSGVRAGSRAEIDLFLPKVLS  
 NL63 619 ELLQKQTSACKTIEDALRNSAMLESADMVSEMLTFTDKAFTLAVANSSF---GDNVLLSSVIPSRLRSGRSGVRAGSRAEIDLFLPKVLS  
 1160 1170 1180 1190 1200 1210 1220 1230 1240  
 SARS-CoV-2 830 DAGFIK-QGDCL---GDIARDLICQFKNGLTVPLPLTDEMIAQYI-SALLAGTITSWGTFGAGAALQIIPFQAMQYAFNGIGVQTNQV  
 SARS-CoV 812 DAGFMK-QGECI---GDIARDLICQFKNGLTVPLPLTDEMIAQYI-SALLAGTITSWGTFGAGAALQIIPFQAMQYAFNGIGVQTNQV  
 HKU1 919 DVGFEV-AYNNCT---GGESEIRDLCVSGYKGLVPLPLLSENOISCYTIAATASLSPFPTIA---AGFVYPLVYQVINGLGVMDV  
 OC43 918 DVGFEV-AYNNCT---GGESEIRDLCVSGYKGLVPLPLLSENOISCYTIAATASLSPFPTIA---AGFVYPLVYQVINGLGVMDV  
 MERS-CoV 902 DPGYMQ-GYDDCMQQGAPASDRLICQAGYVAGKVLPLLSENOISCYTIAATASLSPFPTIA---AGFVYPLVYQVINGLGVMDV  
 229E 702 GLGTWDAVYDQKST---KGLSLADIACAGYVAGKVLPLLSENOISCYTIAATASLSPFPTIA---ASIFSLAQSLNLYVALQTDV  
 NL63 885 GLGTWDAVYDQKST---KGLSLADIACAGYVAGKVLPLLSENOISCYTIAATASLSPFPTIA---ASIFSLAQSLNLYVALQTDV  
 1250 1260 1270 1280 1290 1300 1310 1320 1330  
 SARS-CoV-2 916 LYENOKILQANQFNAISLQKIGLQDLSSTA---SALGHLQDVNVQNAQALN1LKVQLSNNSFGALVSVLNDLILSLRQKVEAE  
 SARS-CoV 898 LYENOKILQANQFNAISLQKIGLQDLSSTA---TALGHLQDVNVQNAQALN1LKVQLSNNSFGALVSVLNDLILSLRQKVEAE  
 HKU1 1001 LNKNQKLIATAFNNALLSIQNGFSATN---SALARIOSVNVQNAQALN1LKVQLSNNSFGALVSVLNDLILSLRQKVEAE  
 OC43 1000 LSONQKLIANAFLNAYAIOEGFDATN---SALVIIQAVNVNAEALN1LKVQLSNNSFGALVSVLNDLILSLRQKVEAE  
 MERS-CoV 990 LSENKQKLIANKFQNLQAGMPTGFTT---EAFQVQDAVQVQNSLTSQRLNQFOATISSQIYALQDYLQV  
 229E 785 LONQKILQASFKNMTAIVDFTGVNDATQTSQALQTVATLNN1QDQDVNVQOGNSLNHTLSQRLNQFOATISSQIYALQDYLQV  
 NL63 968 LONQKILQASFKNMTAIVDFTGVNDATQTSQALQTVATLNN1QDQDVNVQOGNSLNHTLSQRLNQFOATISSQIYALQDYLQV  
 1340 1350 1360 1370 1380 1390 1400 1410 1420  
 SARS-CoV-2 991 VQIDRLITGRLQSLQTYVTOOLIRAAEIRASANLATAITMSECVLGDSKQWVDFCGKGYHLMFMSPPQAFHGVVFLHVTYVAQEKFTTAP  
 SARS-CoV 973 VQIDRLITGRLQSLQTYVTOOLIRAAEIRASANLATAITMSECVLGDSKQWVDFCGKGYHLMFMSPPQAFHGVVFLHVTYVAQEKFTTAP  
 HKU1 1076 VQIDRLITGRLTALNAYQSOQSLDISLVLKFGAALMENXNEVCKVQSKSPHIFNGCNHILSVLQNPYGLYHMFYKQISF1KPTVLS  
 OC43 1075 AQIDRLITGRLTALNAYQSOQSLDISLVLKFGAALMENXNEVCKVQSKSPHIFNGCNHILSVLQNPYGLYHMFYKQISF1KPTVLS  
 MERS-CoV 1065 AQIDRLITGRLTALNAYQSOQSLDISLVLKFGAALMENXNEVCKVQSKSPHIFNGCNHILSVLQNPYGLYHMFYKQISF1KPTVLS  
 229E 874 QVQDRLITGRLAALNAYQSOQVNLNQYKTFYQHATKTYEVTRASRQLOQWQYQKQHGFYQGNGTHIFLVSFLNVAPEGLVYHMBTLLPTQKVDEAWS  
 NL63 1057 QVQDRLITGRLAALNAYQSOQVNLNQYKTFYQHATKTYEVTRASRQLOQWQYQKQHGFYQGNGTHIFLVSFLNVAPEGLVYHMBTLLPTQKVDEAWS  
 1430 1440 1450 1460 1470 1480 1490 1500 1510  
 SARS-CoV-2 1080 AICHGKA---HFPRREGFVSN---GTHWVFTORNYEYQIQTITDNTFVSGNCWVYDDEPKFRSSRLSTQD  
 SARS-CoV 1062 AICHGKA---HFPRREGFVSN---GTHWVFTORNYEYQIQTITDNTFVSGNCWVYDDEPKFRSSRLSTQD  
 HKU1 1165 GLC1GSDV---GIAPKQGYFIKH---NDHNMFTGSSYQYEPISDSDVNFVPMNTCSNFTKAPLVY---LNHSVPLKLSDFESELISHWFKN  
 OC43 1164 GLC1GSDR---GIAPKQGYFVN---NNTKMTGSSYQYEPISDSDVNFVPMNTCSNFTKAPLVY---LNHSVPLKLSDFESELISHWFKN  
 MERS-CoV 1154 GLCDAANPNTCIAPIVNGYIIFKTNNTTRIVEDWSYTGSSFYAEPITSLNKTQVAPOTVQY-ISTHNLPPPLGNSTGIDEQDEDFK  
 229E 963 GLC1GSDV---GIYVLRQPNLALY---KEGNYRQTRIMRFEIPTIADFVQIYCNQTFVNISRE-  
 NL63 1146 GLC1GSDV---GIYVLRQPNLALY---SDNGVFRVTSRVMFOEPLRVPLSVDFQVQIYCNQTFVNISRE-  
 1520 1530 1540 1550 1560 1570 1580 1590 1600  
 SARS-CoV-2 1159 HTS-PDVLG-BISGINASVNNI---QKEIDRNEVARNLNLSEIDLQELQKGYIQLYKQWVYQWVYDDEPKFRSSRLSTQD  
 SARS-CoV 1161 HTS-PDVLG-BISGINASVNNI---QKEIDRNEVARNLNLSEIDLQELQKGYIQLYKQWVYQWVYDDEPKFRSSRLSTQD  
 HKU1 1245 QTS-IAPNLTLNHLHTNATFIDLE---YEMENLQIHSQISLNSLQYINLKDIDGYTQVWVYQWVYQWVYDDEPKFRSSRLSTQD  
 OC43 1244 QTS-VAPDLSL---D-INTVFLFIDLE---QVEMNRLQIHSQISLNSLQYINLKDIDGYTQVWVYQWVYQWVYDDEPKFRSSRLSTQD  
 MERS-CoV 1247 VST-SIPNFG-SLQ-TINTLFLD---TYENLMSLQOVKALNQYIISLDEKLQYNTYQWVYQWVYQWVYDDEPKFRSSRLSTQD  
 229E 1455 LPNTVTPDLD---VVEQYQNTQINLTSSEISTLENKSAEYLNTVQLQDNLNITSTVLWKLWNRVETYIKWVPPWVNLISLCSISVLFVMS  
 NL63 1228 LPKYVKBNF---DLTPEFLYVNLSSLEKOLEAKTASLQFTTVEQGLQDQI-  
 1610 1620 1630 1640 1650  
 SARS-CoV-2 1610 1620 1630 1640 1650  
 SARS-CoV 1610 1620 1630 1640 1650  
 HKU1 1610 1620 1630 1640 1650  
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 MERS-CoV 1610 1620 1630 1640 1650  
 229E 1610 1620 1630 1640 1650  
 NL63 1610 1620 1630 1640 1650



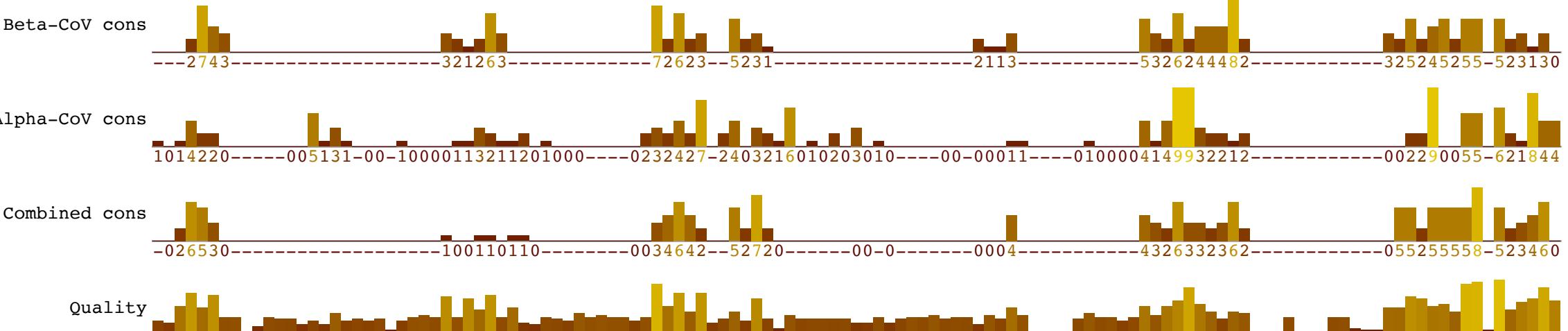


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 229E -A--LLHI--AGCQT-TNGTN-T--SHSVCNGCVGHSENVF-AVESGGYIPS---NFSNNWFLLNTNTSSVVDGVVRSFQPLLNCLWSVSGSQF--TTGF  
 NL63 L-SVK-----CYFNYSCVFSSVNVATVT--VNVTT-HNGRV-V--NYTVCDDCNGYTDNIF-SVQQDGRIPN----GFPNNWFLLTNGSTLVDCVSRLYQPLRLTCLWPVPGLKS--STGF  
 HKU8 V-SIR-----CPDKFSCSSQIVTKAIT--VNVTTFANGTI-D--KYAICDNCNGYPAHIF-PVSEGGLIPA---DFNFSNWFLLTNSSTIVDGRIVSEQPVLLMCLWAVPGLMS--TNSF  
*Ferret-CoV* NVTIKTQTN-----ANQIYWWFNPVRDLSYYVNWKTV--DSTI-VVSNCT--SDCSGYAANIF-AVETGGFIPS---SFSFNNWFVLTNSSTIVSGKFVSSQPLLVNCNLPVPSFGD--ETSI  
*CDPHE15* I-RFS-----CGHKDSCGHQVVYSTNVFNLNG--FI-T-NYQLPDSGEFSNDLF-TVGDDGSIPP---SFGFNNWFVLSNSSSISGTVVSNQPLRLTCLWPPIPSTG--ALAT  
 HKU10 V-NVN-----CAHKSSCQVYPVKDSVT--WVVNTTADGRV-K--DYTICGSCNGFPQHVF-PVSPGGLIIPP---DFSFDGWYVLTNSSTMVQGKFAVASQPLKLLCLWPVPALDS--SASV  
*BtRf-AlphaCoV* V-AVK-----CLVKETCAFFAAHKMVT--LNITTNGQGLI-K--NYSVCKDCNGFPQHVF-PVLDGGKIPG---DFSFDNWFLLTNSSTIVNGKIVDIQPLKLLCLWPVPALVA--NDNK  
*LuchengRn-CoV* SWADTH-----T-----SFEHGSVISYVQICKYPSNVVTI--TNNGCHTNMAG-PGATTCDVILSS-----PLECVLNRTY  
*Mink-CoV* NATISNTFA---SNGTTSNSWWFNPVYNLTYYRVNKT--DNTV-VVQNCT--NNCADYVNNIF-STESGGIIP-----GFSFNNWFVLTNDSTITDGRFLTRQPLLNCLWPVPSFSE--TSQN  
*Bat-CoV-1A* V-SFR-----CVYKETCAHQVITKPIT--AIVNTTADGLI-S--GYSVCCDDCQPKYVF-AVTEGGEVPP---NFRLDNWFYLTNSSTPISGMFTSVQPFNLSCVWTVPVLTN--NSLP  
*BtMr-AlphaCoV* A-TLR-----CANGYACAHQVYVNPVTT--VIVTTVAAG-I-Q--KYTLCDNCTGFPQHVF-ATMENGEIPP---SFNFANWFYLTNSSSPVSSRVVGLQPLLLTCLWPPIPALLG--TATD  
*BtNv-AlphaCoV* V-AFT-----CSKDDSCGHQIITKPIT--VNATTDDKGVI-T--SYTVCDCDGFPHHV-AVQEGGKIPG---SFDFTNWFYLTNTSSPFDGRFVFSNQPLQIQLWPIPALTS--TTGI  
*Porcine-EDV* V-ATR-----CYNRRSCAMQYVYTPTYYMLNVTSAGEDGI-Y--YEPCATCTGYAANVF-ATDSNCHIPE---GFSFNNWFLLSNDSTLLHGKVVSNQPLLVNCLLAIPKIYG--LGQF  
*BtCoV512* V-VSW-----CSAADSCAMQYINSTIYYNLNVTPGPGGI-T--YSVCTKHCTGLADNVF-STDQGHHIPP---IFPYNNWFLLNTSTLVQGVTRVQFLVNCVALPKLQG--LTTF  
 HKU2 WVHRP-----WGA-NANVTVRLCRWQKFSF-----NETAHFQOPAG-P-----SS-----AFCCLVNGSF  
*NL63-related* V-AIA-----CDNPTSCYHSVVTQLAT--VRVQTDKGLI-S--SFEPCQNCCEGFAENVF-AVEASGKIP-----DFSFNNWFLLTNSSSVVDGKVRSLQPLKVLCLRAPSLLS--TKEV  
*FCoV* NVTLLYSRSSTAT--WEYSAAYAYQGSNFTYYKLNNT--NGLK-TYELCEDYEHCTGYATNVF-APTSGGYIPD---GFSFNNWFLLTNSSTFVSGRFVTNQPLINCLWPVPSFGV--AAQE  
 TGE TVTFGDMRATTLEVAGTLVLDLWWFNPVYDVSYYRVNNK--NGTT-VVSNCT--DOCASYVANVF-TTOPGGFIPS---DFSFNNWFLLTNSTLVSGKLVTKOPLLVNCCLWPVPSFEE--AAST  
 OC43 VVQPERTIN-----ST-----QD-----GDNKLQGLLEV-SVCQYNMCEYPQT-----ICHPN-LG-NHRKELWHDGT-----VVSCLYKRNF  
*MERS-CoV* VLLPDG-C-----GTLLRAFYCI-----LEPRSG-----NHCPAGNSYTSFATYHTPATDCSDGNYNRNASLNSFKEYFN-----LRNCTFMYTY  
 HKU1 VVQPHN-----GVLEI-TACQYTMCEYPHT-----ICKSK--GSSRNESWHFDKS-----EPLCLFKKNF  
*Covid-19* LIVNNA-----TNVVIKV-CEFQFCNDPF---LGV-----YYHKNNKS--WMESEFRVYSS-----ANNCTFEYVS  
*Bovine-CoV* VVQPHTTI-----LGNKLQGFLEI-SVCQYTMCEYPNT-----ICNPN-LG-NQRVELWHDGT-----VVSCLYKRNF  
 HKU4 TIIPDG-C-----GTVLHAFYCI-----LKPRTV-----NRCPSTGTGVSYFIYETVHNDQCS-TINRNAALSFKSFFD-----LVNCTFFNSW  
 HKU5 VILPDG-C-----GTILHAFYCV-----LHPRTQ-----QNCAGETNFKSLSLWDTPASDCVSGSYNQEATLGAFKVYFD-----LINCTFRYNY  
 HKU9 LIMAPDNL-----GSW---SA-----VAPRNQNTNIYLLVCSNATLCINPG---FNR-----WGPAGSF-----APDALVDH-----SNSCF--VN  
*SARS-CoV* IIINNS-----TNVVIRA-CNFELCDNPF---FAV-----SKPMGT-----QTHTMIFDN-----AFNCTFEYIS  
 HKU24 VIEPSTQI-----VNGKLIGTLSA-SVCQYTMCEYPNT-----ACNPV-LG-NGRPSLWHASIG-----IVPCLYQRNF  
*Rat-CoV* VLEPYN-----GIIMA-SICQYТИCQLPHT-----YCKPN-TGGNTLIGFWHTDLR-----SPVCILTRNF  
*Bat-BCoV* AISLDG-----LYRRIEV-CNFQVCQNPV---MYY-----AKVSNPAQGNPLIYSQRAYTD-----YNSCVS-QTS  
*Hedgehog-CoV* VILPEK-C-----GTVITALYCV-----LEPKND-----TNCTGSSGYVSYLFENFSCDATT--DGNLKNSLQWFN-----IKDCLFEKNY  
 GCCDC1 VIIAPGEF-----GMW---GQ-----VNRPNNTFVHVVACSNATICAYPM---FNR-----WGPAGSIH-----AENSFVEH-----NSSCF--LN  
 HKU14 VVQPHTTI-----SDKVLLGLEI-SVCQYTMCEYPHT-----ICHPN-LG-NKRIELWHDFTD-----VVSCLYKRNF  
 MHV APRYWVTPLVKROYLFNFNOKGVITSAVDCASSYTSEIKKTOSMLPSTGVYELSGYTVOPGVVYRRVANLPACNIEEWLTARSPVSPLNWERKTFOCNFNLSSLLRYVOAESLFCNNIDASKVY



390 400 410 420 430 440 450 460 470 480 490 500

*229E* VYFNGTG--RG-ACKGF-YS-NASSDVIRYNINFE--NLRRGIFTL-KTSYGAFFVYCTNNTL--VSGDAHIP--SGTVLGNFYCFVNNTI--G--NE--TTSAFGVGAL-PKTVRE  
*NL63* VYFNATG--SDVNCNGY-QH-NSVVDVMRYNLFNSANSLDNLKSGVIVF-KTLQYDVLFYCSNSSS--GVLDTTIP--FGPSSQPYCFINSTI--N--TT--HVSTFVGIL-PPTVRE  
*HKU8* VYFNGTA--PNKQCNGYATD-SAF-EALRFSLNFTDERVFA-GSGSVL-LVSLQYKFSCTNNSE--AVIDSGIP--FGNVVEPFYCFVSING--TSIFVGML-PAVLRE  
*Ferret-CoV* INFDTV-P---SQCNGA-TV-NGSFDVVRFLNFTANVASSSGTSFLALNTGGVVLYLSCFNETKTTNLYMSEGALP--FGTHEGALCYVSYNE--TSFKYIGVLPSSVKE  
*CDPHE15* IYFNGT--NGAQCNMF-DS-NAPFDAIRFLNGLSG--HNFVSGFVLHAANGATLGFSCTNSTDA--PY-LRQIP--FGIGDTPYYCYLNVT--D--IN--STMSFGAL-PLNLRE  
*HKU10* VYFNLSR--SSAQCNMATE-NGLADALRFSMNFSRNG-FS-GAHLSTL-KG-AIDYVFRCNTNTN--VTSFYSIP--FGYIKTVYRCFVSYNS--T--NG--TVSEFGVLPPLVKE  
*BtRf-AlphaCoV* IYFNITG--ANCNGFKDE-NAYADVLRFSLNFTNSQFN-GIHSVTV-NVVGDLKFSCSNTSTY--AESDEVLP--FGPLSSVYYCFVNGKL--G--ND--TFSRFGVIL-PPVVKE  
*LuchengRn-CoV* --SQQ-----Y-----AGVSYI-TWYNDHI--IA-----SI-----QGEVFTFDIG--EVLQW--SNFSACCGT  
*Mink-CoV* FCFNTEDF--AQCNMGY-RL-NSTADVIRFLNLFNTHETVAASGTTFFELNTTGGVVLHISCVNQSRTE-AVTSEGFLP--FGYHENSLYCYVSYNE--TLQKFLGVL-PPSVKE  
*Bat-CoV-1A* VYFNMSM--NENVTCGYSNFDVGVDAMRFSLNFTDQII--RKGVISV-FTSTNVNFSCTNNSVL--DG--DAVIP--FGDVAQTHYCFIKYFT--N--ITSGESVSQFVGIL-PPQVKE  
*BtMr-AlphaCoV* ITFDRNG--TSDVRCNGFASNE--TADAMRFSLNFTDAVFA-KEGVITL-KTLSNTFKFSCSNSSTY--QA--PYVIP--FGHIDQPYYCFTFYI--N--ETAGTTTSFVGML-PPVVRE  
*BtNv-AlphaCoV* IYFNSSRFTDPHQRCNGYNEV-GGLADHRLRAINVTDRGAF--QLGVISL-LAVHNAYNFSCSNISNF--VDATTIGVP--FGKTYQPYYCFCVTEGY--N--ISA--NRTFVGVM-PSDVRE  
*Porcine-EDV* FSFNHTMD--GVCNGA-AV-DRAPEALRFNINDTSVI--LAEGSIVLHTALGTNLFSVCSNSSDP--HLAIFAIP--LGATEVPYYCFLKVDT--Y--NS--TVYKFLAVL-PPTVRE  
*BtCoV512* LSFDSPLN-VPGFSCNGA-NG-SSSAEAFRFNVNDTKLF--VGAGAVTLNTVDGVNVSVICSNNAQ--PTRSN--NLQEDLPYYCFTNTSS--G--TN--HTVKFLSVF-PPIIRE  
*HKU2*--PSSQ-----HK-----GYMFGV-TWYNDFV--RI-----IF-----PPTVFELOLD--G-LQWEYVQFTGPVNA  
*NL63-related* ISLSGVN--AD-KCNGH-SI-NETAGALRFNLFNFTSNPINALASKIFI-SSSGNVTIFCSNSSDP--GSSADAFIA--MGSTSAAVYCFANSTF--G--NV--TSLDFIGVL-PVTVRE  
*FCoV* FCFEGAQF--SQCNGV-SL-NNTVDVIRFLNFTADQSGMGTAVFSLNTTGGVILEISCYSDTVSESSSYSGEIP--FGITDGPRYCYVLYNG--TALKYLGTL-PPSVKE  
*TGE* FCFEGAGF--DOCNGA-VL-NNTVDVIRFLNFTTNVOSKGATVFSLNTTGGVTLIEISCYT--VSDSSFFSYSGEIP--FGVTDGPRYCYVHYNG--TALKYLGTL-PPSVKE  
*OC43* -TYDVNA-----DYLYFHFYQ--EGGTFYA--YFTD-----TGVV-----TKFLFNVYLG--MALSHYYVM-PLTCNS  
*MERS-CoV* --NITE----D-----EILEWF--GITQT--AQGVHL--FSSRYVDL--YGGN--MFQFATLPVY--DTIKYYSIIPHSIRS  
*HKU1* -TYNVST-----DWLYFHFYQ--ERGTFYA--YAD-----SGMP--TTFLFLSYLG--TLLSHYYVL-PLTCNA  
*Covid-19* QPFLMDL--EGKQGNF--KNLREFVFKN--IDGYFKI--YSKHT--PINL-VRDLPQGFS--LEPLVLDLPIG--INITRFQTL-LALHRS  
*Bovine-CoV* -TYDVNA-----DYLYFHFYQ--EGGTFYA--YFTD-----TGVV-----TKFLFNVYLG--TVLSHYYVM-PLTCNS  
*HKU4* --DITA----D-----ETKEWF--GITQD--TQGVHL--YSSRKGD--YGGN--MFRFATLPVY--EGIKYYTVI-PRSR  
*HKU5* --TITE----D-----ENAEEWF--GITQD--TQGVHL--YSSRKENV--FRNN--MFHFATLPVY--QKILYYTVI-PRSR  
*HKU9* NTFSVNIS--T-----SRISLASF--KDGDLI--YHSGWL--PTSNFEHGFSRGSH--MTYFMSPVG--GNLPRAQOFF-QSIVRS  
*SARS-CoV* DAFSLDV--SEKSGNF--KHLREFVFKN--KDGFLYV--YKGYQ--PIDV-VRDLPSPGFNT--LKPIFKLPLG--INITNFRAI-LTAFS  
*HKU24* -TYNVA-----DNIYFHFYQ--DGGTFYA--YVG-----KSPI--TTLLFQVYLG--TVVTHYYVL-PLVCNA  
*Rat-CoV* -TFNVNA-----EWLYFHFYQ--QGGTFYA--YAD-----VSSA--TTFLFSSYIG--AVLTQYFVL-PYMCNP  
*Bat-BCoV* QSFNTSQ--GDKNGAFSK--DTITTYVFKY--QNGIFSV--FHGFH--NGTD--HVPI--PKPAFEFPLD--IPITHIFIIP-PTVEE  
*Hedgehog-CoV* --NVTD----D-----EREERWF--GIIQD--QQGVHL--YTSRKN-G--YSNN--MFLFATLPIY--DQILYYTVM-PRSI  
*GCCDC1* DTFEIPMG--T-----SRVNLAFR--QDGNL--YHTAWL--PTENYTL--SGDYP--LRYARSGVG--SNLPFAQFF-QSVARA  
*HKU14* -TYDVNA-----DFLYFHFYQ--EGGTFYA--YFTD-----TGTV-----TKFLFNVYLG--TVLSHYYVM-PLTCDS  
*MHV* GRCFGSISVDKFAVPRSROVDLOLGNNSGFLOTANYKIDTAATSCOLHYTLPKNNVTINNNPSSWNRRYGFNDAGVFGKNOHDVVYAOQCFTVRSSYCPAOPDIVSPCTTOTKPKSAFVNVDHCE



510 520 530 540 550 560 570 580 590 600 610 620 630

**229E** FVISRTGHFYINGYRYFSLGDVEAVNFNVTN---AA-TIVCTVALASYADVLVNVSQTAIANIIYCN-SVINRLRCDDQLSFDFDVPDGFYSTSPI--QPVELPMSIVSL-----

**NL63** IIVVARTGQFYINGFKYFDLGFIIEAVNFNVTT---ASATDFWTVAFATFVDVLVNVSATNIQNLLYCD-SPFEKLQCEHLQFGLQDGFYSANFL--DDNVLVPETYVAL-----

**HKU8** IVITTRYGSIYLNGFSIFQGPPIQGVLFNVTN---RGATDLWTVALSNTTEVLAEVQSTAIAKALLYCD-DPLSQLKCQQLQFSLPDGFYATASL--FQHELPRTFVTL-----

**Ferret-CoV** IAVSKWGFVYINGYNYQTFFPIDSVAFNLTT---GNSGAFWTIAAYTFTDVLEVSQDQIKSVTYCN-SHINDIKCSQMSENLPDGFYPVSQH--SLPNVNKTTFVTL-----

**CDPHE15** TIVIASNGDVMNGYRYFAAGDLSSVVDLPSQ---QVFGSTFWTIAYTFTVLEVDGTSINRMLYCD-NPLNRVKCSHTQFDLVDGFYPLTDVLD--AVKPFVFVTL-----

**HKU10** IVVSRYGTIYLNGVRIFKLPPLEGGVVFNVTS---DVGSDFWTVAFQAQNLEVNAATNIVNLHYCD-NTLNKVKCQQLSFQLPDGFYPTTNL--VDSVVPKTYVAL-----

**BtRf-AlphaCoV** IVISRYGSFYMNGVKLFEVYPVESVVFNVTS---SVGSDFWTVAYAQNTEVLLLEVNVSTDIKDILYCD-TPANKLKCQQLSFSLLEDGFYPAALV--TGVEVPRTYVAL-----

**LuchengRn-CoV** -----GNKCGFSYATTLSEWLVRTDNDGTVIDYVICDCTDFESQLKCKNMVFELETPAVYSGSAVE---LQSAIYYV--SNELPDCDFSFADMFM-----

**Mink-CoV** IAISKDGGFYINGYNYQTFFPIDCISFNLT---GNTGAFWTIAAYTSDVMVDVENTAIKRVVYCN-SHIDDIRNCNQLTNPNLPDGFYPVSPK-AIGFVNKTTFVTL-----

**Bat-CoV-1A** FVIMRNQDFHLLNGYRIFSVDRVESAIFNIST---NDGRDFWTVAFANNAEVLTEINATSIQNLLYCN-NPVNSIKCQQLRFNLDLGFYSHTFD-SADNLPTTIVRL-----

**BtMr-AlphaCoV** FVITKTGNVYLNGYRIFTVDDVVSVNFNISS---TDHRDFWTVAFKNTEVMDIEDTYIKQOLLYCN-TPLNVVKCQQLKFVLLDDGFYSYSSP-VDEVLPRTTIVRL-----

**BtNv-AlphaCoV** IVVSRYGSVYINGYKIFNVGELYGVVLNFSS---LTGSDFWTVAYANEVNVLVDIEETYITGILYCD-TPLNRLKCQQQRFFMDDGFYSAI-D-LAPPVQTIPLL-----

**Porcine-EDV** IVITKYGDVYVNGFGYLHLGLDAVTINFHTGHTDDDVSQFWTIASTNFVDALIEVQGTSIQRILYCD-DPVSQLKCSQVAFDLDGFYPISSRNLLSHEQPISFVTL-----

**BtCoV512** FVITKYGNVYVNGYIYLRTTRPLTVAVHLASS-HSQDVAGFWTIAATNFTDVLVEVNNTGIQRLLYCD-TPENSVKCSQSFELLEDGFYSMTADNVYAVTKPHTFVTL-----

**HKU2** -----GRMTKFNVVTIEISSVLLTDQSGAVTRYSYCADGFVNGLQCKLRLFDIPPGVYNSNEVE--YPVALYTV--VHNMSVCQRPESY-----

**NL63-related** FVFAATGQIYINGYNFFSTFPICISFNLT---DNVTDFWTVAYTQFVDTLVAVNNNTLHQEVLYCD-DVIHKLKCSQSFQDLPDGFYASALV-RDERLDKTFVTL-----

**FCoV** IAISKWGHFYINGYNNFFSTFPICISFNLT---GVSGAFWTIAAYTSDVQVNTAIKNTVYCN-SHINNIKCSQTLANLNNGFYPVASS-EVGFVNKSVLL-----

**TGE** IAISKWGHFYINGYNNFFSTFPICISFNLT---GDSDFWTIAAYTSDVQVNTAIKNTVYCN-SHVNNIKCSQITANLNNGFYPVSSS-EVGLVNKSVLL-----

**OC43** -----KLTLEYWVTPPLTSRQYLLAFNQDGIIFNAVDCMSDFMSEIKCKTQSIAAPTGVYELNGYT--VQPIADVYR-RKPNLPNCNIEAWLNDK-----

**MERS-CoV** I---Q---SDRKA-WAAFYVYKQLQPLTFLDFSVDGYIRRAIDCGFNDLSQLHCSYESFDVESGVYSVSSFE-AKPGSGVVE-QAEGVECDFSPLLSG-----

**HKU1** I-----SSNTDNETLQYWVTPLSKRQYLLKDNRGVITNAVDCSSSSFFSEIQCQKTKSLLPTGVYDLSGFT-VKPVATVHR-RIPDLPDCDIDKWLNFF-----

**Covid-19** YLTPG-----DSSSGWTAGAAAYVGQLQPRTFLLKYNENGTTDAVDCALDPLSETKCTLKSFTEVKGIQTSNFR-VQPTESIV-RFPNITNLCPFGEVFNAT-----

**Bovine-CoV** -----ALTLEYWVTPPLTSRQYLLAFNQDGVIFNAVDCKSDFMSEIKCKTLSIAPSTGVYELNGYT-VQPIADVYR-RIPNLPDCNIEAWLNDK-----

**HKU4** K---A---NKREA-WAAFYVYKHLQHLYLDFSVDGYIRRAIDCGHDDLSQLHCSYTSFEVDTGVYSVSSYE-ASATGTFIE-QPNATECDFSPMLTG-----

**HKU5** P---F---NDRKA-WAAFYIYKLHPLTYLLNFDEVEGYITKAVDCGYDDLAQLQCSYESFEVETGVYSVSSFE-ASPRGEFIE-QATTQECDFTPMLTG-----

**HKU9** NAI-D---KGDMGCTNFDVNLHVAHLINRDLLVSYFNNGSVANAADCADSAAEELYCVTGSFDPPTGVYPLSRYR-AQVAGFVRVTQRGSYCTPPYSVLQ-----

**SARS-CoV** -----PAQDIWGTSAAAAYFVGYLKPTTFMLKYDENGTITDAVDCSQNPLAELKCSVKSFIDEKGIIQTSNFR-VVPSGDVV-RFPNITNLCPFGEVFNAT-----

**HKU24** -----RQTYEYWVTPLIKREYLLFVGNGVITNAVDCASDHMSEIQCMTQNIKPVTGVYELTGYT-VQPIADVYR-RIPNLPDCIEEQWLNDP-----

**Rat-CoV** T-----TSG-VFSPQYWVTPPLVKRQYLFNFQNKGIIITSAVDCASSYTSEIKCKTQSMPNTGVYDLSGYT-VQPVGLVYR-RVRNLPDCKIEEWLAA-----

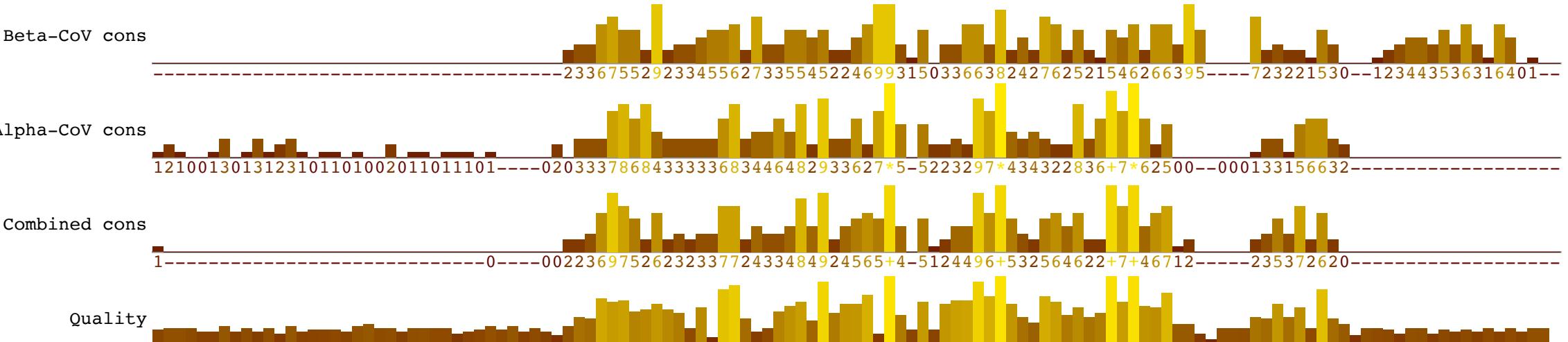
**Bat-BCoV** -----P---AQGQGYTRGNYVYVYQAQLKKSTFMFGFDSNGLITRYIDCAAGPKDELMCSQGFSNITPGVYTTTNYR-ATPKQHVVITSADVSAECPFQSLINVTE-----

**Hedgehog-CoV** SN---Y---ASYHA-FSAFYIYKLHINYMVDVDVNGYITRAVDCGYNDYTQLOCSYQFDMDSGVYASAYFN-ARSRGYYYE-AAELTECDLDVLFKN-----

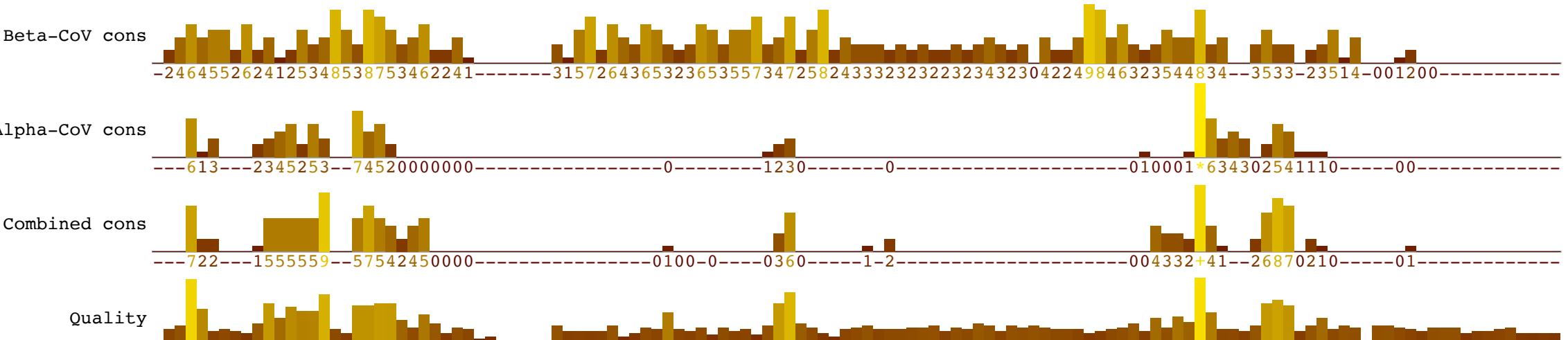
**GCCDC1** R-----DSACAFHNLYIAPVQPKELLVKYNDQGMPVEIADCSDADSTQELYCITGFTTPSVGVYQLSRYR-AEAKKLVQVTQQEDSCAIPYTTIL-----

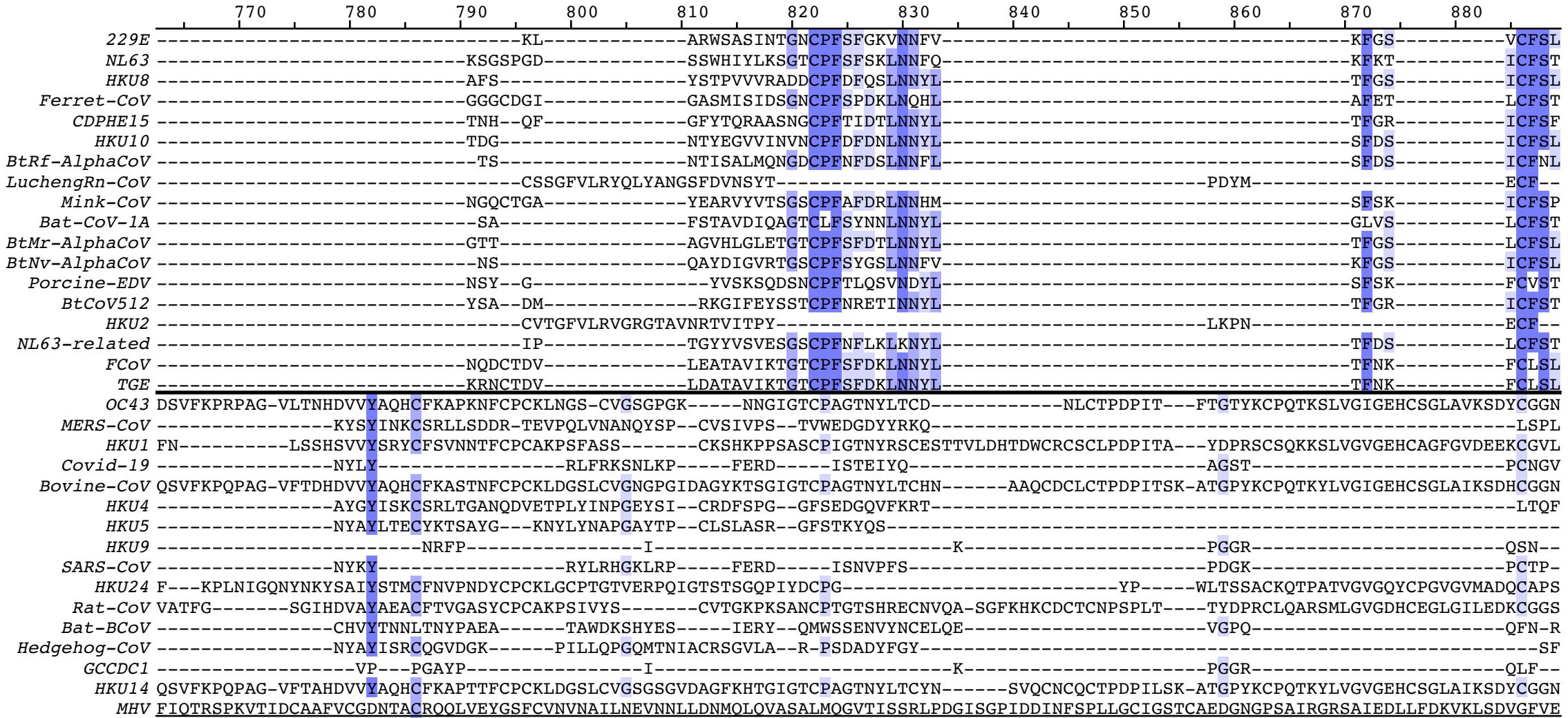
**HKU14** -----TLSLEYWVTPPLTPRQYLLAFNQDGIIFNAADCMSDFMSEIKCKTQSIAAPSTGVYELNGYT-VQPIADVYR-RIPNLPDCNIEAWLNDK-----

**MHV** GLGVLEDNCGNADPHKGICICANNFIGWSHDTCLVNDRCOIFANILLNGINSGTTCTSDLOLPNTEVVTGICVKYDLYGITGOGVFKEVKADYNSWOTLLYDVNGNLNGFRDLTTNKTYTIRSCYS-----



640 650 660 670 680 690 700 710 720 730 740 750 760  
 229E ---PVY---HKHTFIV--LHVKFEHQQRGP-----GKCYNCRPSVI---NITLA---NFn-----ETKGPLCVDTSHFTQFVD---NV---  
 NL63 ---PIY---YOHTDIN--FTAT--ASFG-----GSCYVCKPHQV---NISLN---G-----NTSVCVRTSHFSIRIYIY---NR---V---  
 HKU8 ---PRH---FTHSWIN--LRIKWKN-----GVCYNCPPASSW-IDFVTFSNNGT-ENV-----LPERTLCVNNTQFTTNLTL---IE---E---  
*Ferret-CoV* ---PAN---FEHTSIN--VTGNVLLIYYA-----HPVF-R-SGNVTLH-----P-----QGTQTICVNNTQFTFNFDSD---QCHQLA---  
*CDPHE15* ---PTF---ADHSFVY--FNFSLMFDDLN-----E-----D-----FRLQSFLN-T-I-----NGQLSYCVQSRQFTSGSV---RT---N---  
 HKU10 ---PYH---ATHTYIE--LNITVNY-----VRD-----E-----IAI-----T-----GGGELETCVNNTQFTTLFST---NA---A---  
*BtRf-AlphaCoV* ---PYH---ATHMFVN--LTvhLGN-----SVDADA-----Y-----VQI-----N-----GVNDTVCVNNTQFSTHFQQ---LD---  
*LuchengRn-CoV* DGTGNFEGLRRHVFNSNCVNYTAWFACAD-----DY-SCIIFNA-----IFAEVR-YKLSQPDG-----LVN-----PFIKCNGLDLYSITKG---  
*Mink-CoV* ---PAN---FDHVYVN--ITGNVRLKIQR-G-----RPYN-N-GGNATLS-----F-----Q-----GCIDVSQFTVNVNH---TCTLID---A---  
*Bat-CoV-1A* ---PKY---VTHSYIN--VTVGVNFEENG-----LTNYL-----VDFAPDGTSIGK-----DNITTVCVDTTTFTTRLNV---LA---F---  
*BtMr-AlphaCoV* ---PRL---MTHNFLN--FTIFVSFYFDDDQARPDGFFYECATCAPKYYK-----LAFVDDWTSAVS-----SNVTSICVNYASFTTRLFT---FY---A---  
*BtNv-AlphaCoV* ---PEY---TGLTNIT--LDVNVTFNPP-----SCIQCAPTVNS-----ILLN-----G-----ESGGSVCVSTNRFSVDFKL---TV---  
*Porcine-EDV* ---PSF---NDHSFVN--ITVSAAFFGGLS-----S-----A-----NLVASDT-T-I-----NGFSSFCVDTRQFTITLFY---NV---T---  
*BtCoV512* ---PTF---NDHGfvN--VTVGGNFDSYY-----P-----P-----KFTANGT-L-V-----NNGTVVCVTSNQFTLRHDF---MV---G---  
 HKU2 -CGSNYCPFKRUVFNSNCVNVNTSWTSGLL-----RDY-QHLVLVPNG-----KFN-----PFTECNGLN--RIVDD---  
*NL63-related* ---PTH---SDHSNVT--LYVSFNTYSSS-----TC-TTKPDHVT-SCQYNVT-----IVG-----ENDGPVCVKSKQFTPLQTS---S---  
*FCoV* ---PSF---FTYTAVN--ITIDLGMKLSGYG-----QPIAST-LSNITLP-----MQD-----NNTDVYCIRSNQFSVYVHS---TCKSSLWDNIF---  
 TGE ---PSF---YTHTIVN--ITIGLGMKRSGYG-----QPIAST-LSNITLP-----MOD-----HNTDVYCIRSDOFSVYVHS---TCKSALWDNIF---  
 OC43 -SVPSPLNWERKTFNSNCNFNMSSLLMSFIQ-----ADSFTCNNIDAABIYGMCFSSITIDKFAIPNGRKVDSLQNLGYLQSFNYRIDTTATSCQL--YYNLPAANVS-VSRFPNSTWNKRGFIE---  
*MERS-CoV* -TPPQVNFKRLVFTNCNLYNLTKLLSLFS-----VNDFTCSQISPAIASNCYSSLILDYFSYPLSMKSDSLVSAGAPISQFNYKQSFNSPTCLI--LATVPHNLTT---ITKPL---  
 HKU1 -NVPSPLNWERKIFNSCNFNLSTLLRLVH-----TDSFSCNNFDESKIYGSCKSIVLDKFAIPNSRRSDLQLGSSGFLOSSNYKIDTTSSSCQL--YYSLPAINVT-INNYPSSWNRRYGFNN---  
*Covid-19* -RFASVYAWNKRISNCVADYSVLYNSAS-----FSTFKCYGVSPPTKLNDLCFTNVYADSFVIRGDEVRQIAPGQTGKIADYNYKLPDDFTGCVI--AWNS-NNLDS-KVGGNY---  
*Bovine-CoV* -SVPSPLNWERKTFNSNCNFNMSSLLMSFIQ-----AYSFTCNNIDAABIYGMCFSSITIDKFAIPNGRKVDSLQNLGYLQSFNYRIDTTATSCQL--YYNLPAANVS-VSRFPNSTWNRRGFTE---  
 HKU4 -VAPQVNFKRLVFSNCNLYNLTKLLSLFA-----VDEFSCNGISPDSIARGCYSTLTVDYFAYPLSMKSYIRPGSAGNIPLYNYKQSFANPTCRV--MASVLANV-T---ITKPH---  
 HKU5 -TPPIYNFKRLVFTNCNLYNLTKLLSLFQ-----VSEFSCHQVSPSSLATGCYSSLTVVDYFAYSTDMSYYLQPGSAGIAVQFNYKQDFSNPTCRV--LATVPQNLT---ITKPS---  
 HKU9 -DPPQPVWRYMLYDCVFDFTVVVDSL-----THOLQCYGVSPRRLASMCYGSVTLDVMRINEETHNNLFNRVPDTFSLYNVALPDNFYGCYH--AFYL-NSTAP-YAVA---  
*SARS-CoV* -KFPSVYAWERKKISNCVADYSVLYNSTF-----FSTFKCYGVSATKLNDLCFSNVYADSFVVKGDDVRQIAPGQTGVIADYNYKLPDDFMGCYH--AWNT-RNIDA-TSTGNY---  
 HKU24 -QVPSPLNWERKTFNSNCNFNMSSLLSKVR-----ATSFSCNNIDAASKIYDMCGSITIDKFAIPNSRKVDSLQFGSSGYIIONNYRLQSATSCQL--YYGIPANNVT-VTKKNPSGWNNRYGFVE---  
*Rat-CoV* -TVPSPLNWERKTFQNCNFNLSSLLRFVQ-----AESLSCSNIDAASKVYGMCFGSISIDKFAIPNSRRVDSLQLGKSSLQSFNYKISTRATSCQL--YYSLAQNNVT-VINHNPSWNRRYGFND---  
*Bat-BCoV* ATIPSPAFWRRHYVRNCNLYDISVFTDNAD-----VYSLQCYGVAPSSIADMCEEAHIDYMKISEKDIIFSFKPGSGAGDFAKNYKLPDFMGCTV--VFTN-QELTC-NATSGQL---  
*Hedgehog-CoV* -DAPIIANYSRVFTNCNLYNLTKLLSLVQ-----VDEFVCHKTTPEALATGCYSSLTVDFALPFMSKSTLAIGSAAISMFNYNQDYSNPTCRI--HAAVTANVSTALNFTANA---  
 GCCDC1 -EPSPAAWRATISNCTDFESLLRTL-----TYNLKYGYGISPARSTMCYAGVTLIDFKLNTTHLSNMLGSVPDAVISIYNALPSNFYGVH--AYHL-NSTTP-YAVA---  
 HKU14 -SVPSPLNWERKTFNSNCNFNMSSLLMSFIQ-----ADSFTCNNIDAABIYGMCFSSITIDKFAIPNGRKVDSLQNLGYLQSFNYKIDTSATSATSCQL--YYNLPAANVS-VSRFPNSTWNRRGFTE---  
 MHV GRVSAAFHKDAPEPALLYRNINC SYVFSNNISREENPLNYFDSYLGCVNADNRTDEALPNCDLRLMGAGLCSVYDKSRRRAHRSVSTGYRLTTFEPYTPMLVNDSVOSVDGLYEMOIPNFTIGHHEE





Beta-CoV cons



Alpha-CoV cons



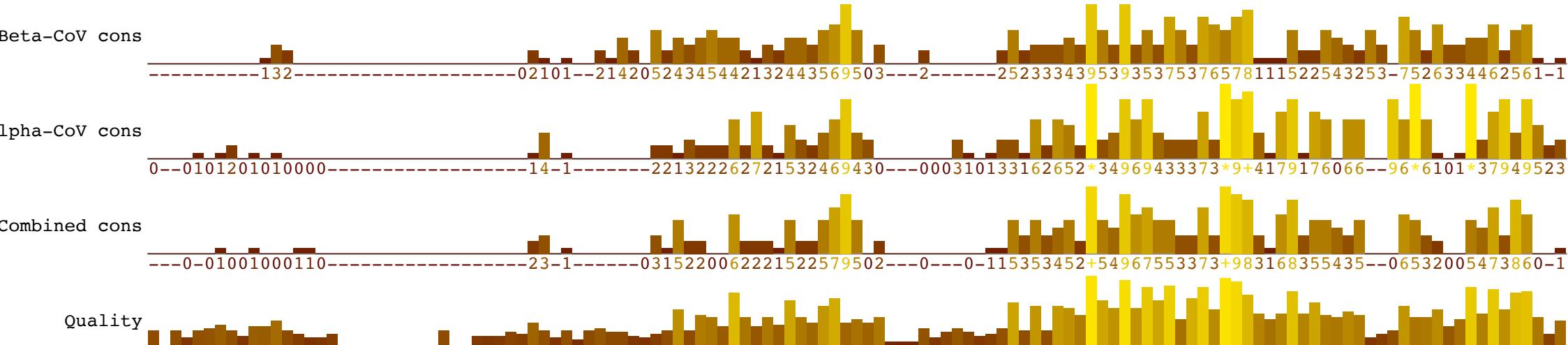
Combined cons



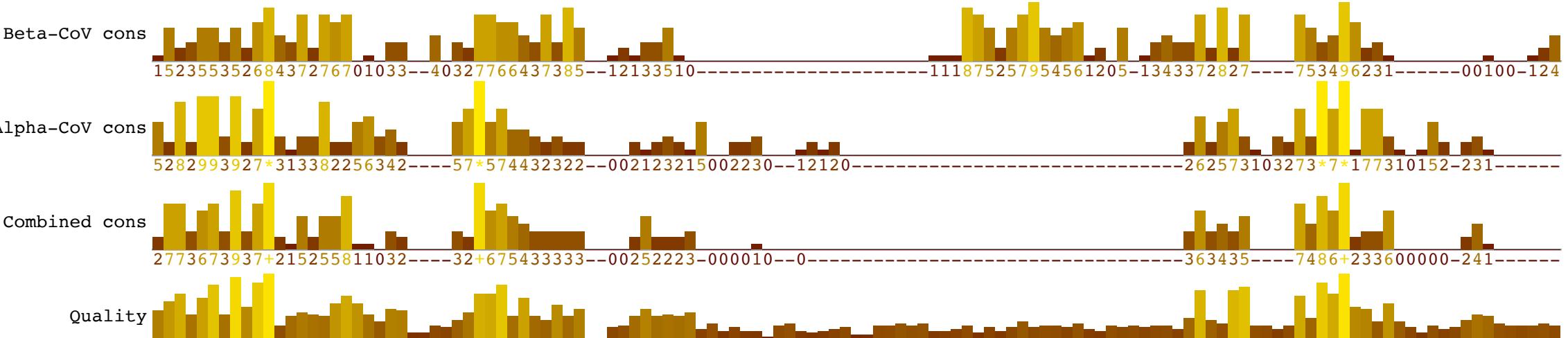
Quality



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 NL63 V-EVPGSCNFPLEATW-----HY-T-----SYTIVGALYVTWSEGSNTGVPY---PVSGIREFSNLVLNNCTKYNIYDVGTGIIRSSNQSLAGG--ITYVSNSGNLLGFKNV  
 HKU8 NGTIKGCTLGIYKRA-----SS-Q-----YIPIWNVWVAYTSGDNILGVRE---PNVGVRDQSVVHQNVCTSNTIFCHSGRGIIRPANISYIAG--VYYTAASQOLLGFKNV  
*Ferret-CoV* L-PTGSDCAFSLTVQN-----RYFT-----RVFAHLFVTYKYGLDHLGQVOT---PDVGVKDLSVVYQNVCTEYNNIYGHAGVGIIRSTNQTLLGG--LYYTSLSGDLLGFKNV  
*CDPHE15* G-ESGAGCGVDMVYES-----QYNM-----FKVTTIFVSYSEGDIAGMPK---PSEGIRDVSKIHLDCVCSTYSIYGHGTGDIIRLTNDTLLGG--LYYTS...  
*HKU10* K-PIAGGCTMRLVKQI-----LV-S-----YFDIGVLYVSYTPGSKITGVLQ---PTGGVYDPSIVHMGVCTDYYVYGLTGRGVISVSNNTTYIAG--LYYTS...  
*BtRf-AlphaCoV* Q-PVGSSCTISIMKNW-----MG-F-----KAPWKSVYVTFKRGNRITGVKN---ASTGIFDPSVMQLNECTDYTIYGVSGRGIIRKANSSYISG--LYYTS...  
*LuchengRn-CoV* -----GYFALY-----NGYVIYN--AKFVSKGLTVCVVQ-----PVEPELDVCKSYTIDGVTFOQILRNTTHQIDSFHNL...  
*Mink-CoV* V-PLGDDCRLDLTVQT-----RYFT-----GVFAHVYVSYKFGLDIVGLPR---ADAGLKDL...  
*Bat-CoV-1A* K-PNGGCMMSSVARG-----YFGE-----FQKGVLYV...  
*BtMr-AlphaCoV* V-ANGGCTMNIVTQG-----PYGL-----PHIAVLYVSYTEGDNIIGVPLSNIPPLGVSDMSQVYLDTCTTYIYGMTGRGVITRSNNNTFITG--LYYTS...  
*BtNv-AlphaCoV* V-DNGGCPMPINAIN-----YIKI-----EYPIGVLYVTHSPGEAITGVPKGFIKRLGFLDASILH...  
*Porcine-EDV* S-LLAGACTIDLFGYP-----AFGS-----GVKL-TSLYFQFTKGELITGTPK---PLEGITDVSFM...  
*BtCov512* S-PADGACELKYYVWN-----TIGA-----VSHLAGTLYVQHTKGDIITGTPK---PLOGLN...  
*HKU2* -----GWSWND-YQDSIYDW...  
*NL63-related* K-QLPGGC...  
*FCoV* S-PVGANCKFDVAART-----RTNE-----QVVRSLYVVIYEEGDNIVGVPS---DNSGLHDLSVLHLD...  
*TGE* S-PVGANCKFDVAART-----RTNE-----QVVRSLYVVIYEEGDNIVGVPS---DNSGVHDL...  
*OC43* -----SCTCRPQ-----AFLGWSADS-CLQGD...  
*MERS-CoV* ---EGGGWLVASG-----S-----TVA-----MTEQLQMGFGITVQYGTDTNSVCPKL---EFAND-TKIAS...  
*HKU1* ---D-GSYNV...  
*Covid-19* ---EGFNCYFPLQSYG-----F-----QPTNGVYQ...  
*Bovine-CoV* -----PCTCQPQ-----AFLGWSVDS-CLQGD...  
*HKU4* ---EGGG...  
*HKU5* ---HSDGE...  
*HKU9* -----SAFIDTV-----I-----NAAHY-SPFSYVYGLAVITLKPAAGSKLVCP-V---AN-----DTVVITDRCVQY...  
*SARS-CoV* ---PALNCYWPLNDY...  
*HKU24* ---MPGNSILTCS...  
*Rat-CoV* ---N-ICNCSAD-----AFVGWAKDS-CLSNDRCHIF...  
*Bat-BCoV* ---MPRNCYVRSNN...  
*Hedgehog-CoV* ---QGRNYYLGRK-----S-----YKPKT-DEGDVQM...  
*GCCDC1* -----NSFV...  
*HKU14* -----PCTCQPQ-----AFLGWSVDS-CLQGD...  
*MHV* AYNNCTGGOEVRD...  
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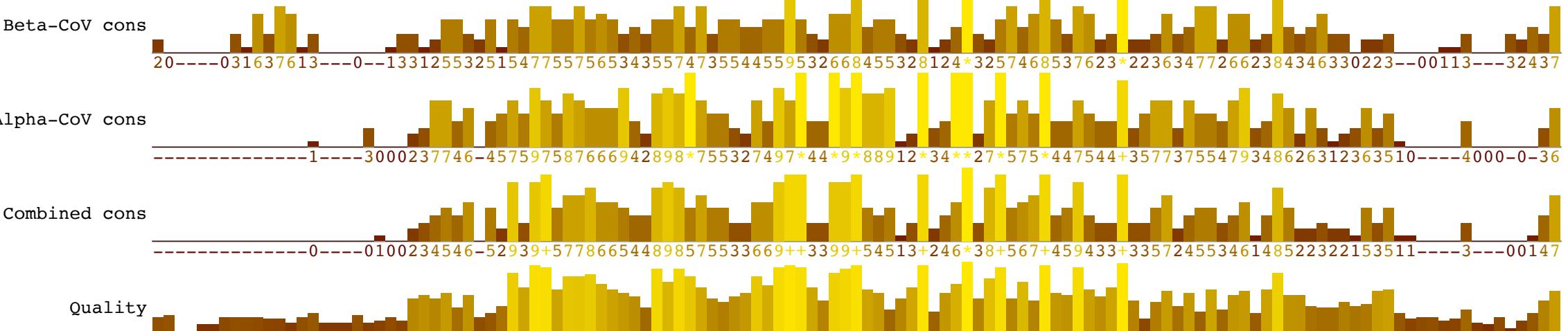


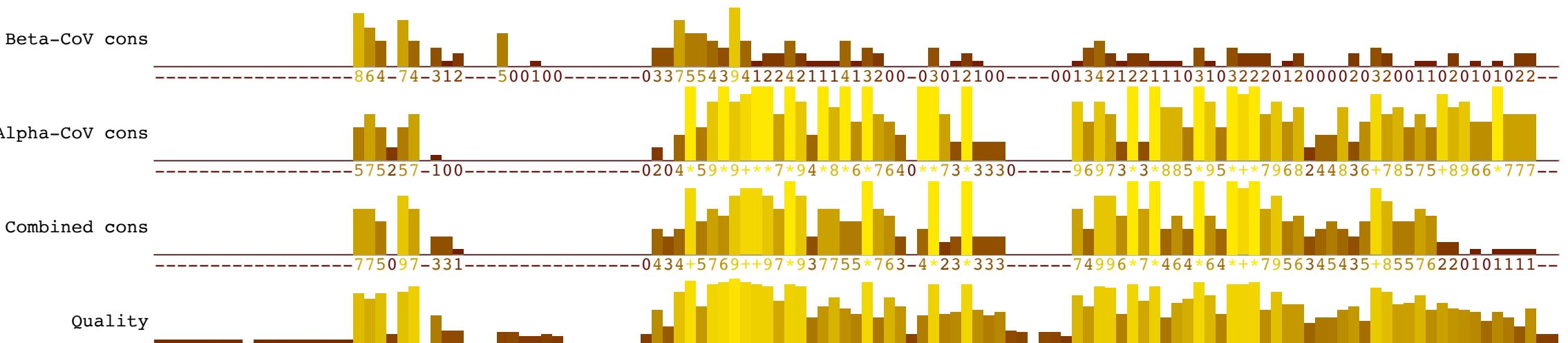
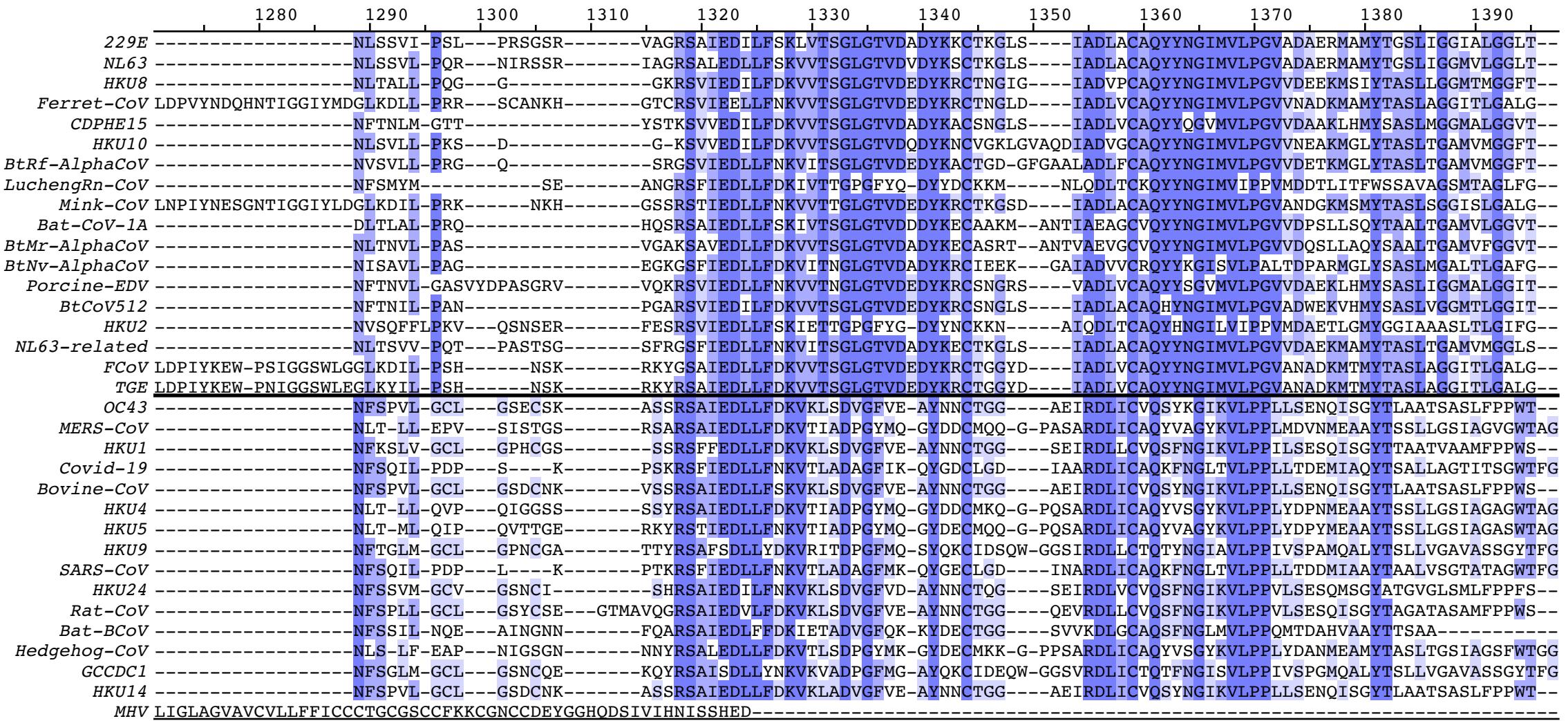
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 NL63 STGNIFIVTPCNQPDQVAVYQQS--IIGAMTAVNESR--YGLQNLQLPNFFYYV--SNGGN--NCTTAVMTYSSNFGICADGSLIPVR-PRN--  
 HKU8 TTGEVFSVTPCNPSSQAVVVKDR--LGVMSSTSTVS--IPFNNNTIPTPSFYH--SNATS--SCDDPSVYSSIGICDDGGITFVN-STR--  
*Ferret-CoV* TTGEVYSIVPCQLSAQAAVINGK--IVGAUTSVQSPI--LDLPHHVTPQFYHHSIYNYSATPTSYRTNNGFDKYL--VNCTPIITYSNMGVCENGALVFIN-ITQ--  
*CDPHE15* TTGEIYSVTPCSLTQQVAVVSD--IVGVVSSSANVDQFSFTYTTVTEQFYMM--TDGNS--SCSEPVLTYSLLGVCKDGAIRLQ-PRE--  
*HKU10* TTGVLYAVEPCDTSSQAVVGDA--IIGAMTSSANVT--FGVNTTVLPSPFYH--TNFGF--GCQTPVLEYGSMGLCADGSVVDVT-LRQ--  
*BtRf-AlphaCoV* TTGEVFSVTPCQLTMQAAVNDD--IVGVVSSNTSG--INFTHITTKTFFYH--TNVQQ--NCSEPVLTYANIGVCSDGSITEVT-VRK--  
*LuchengRn-CoV* IRGVVYAVESCRNFYYS-VFKTL--SAIGLYLSGATCD--STDVTTFMT--KA--RA--TTFVDSSLGCFIDVSV-TDGNYTDCLNPI---GNGFCVDVNVTGQPVVG--  
*Mink-CoV* TTGTVYSITPCELSAQAVIDGS--IVGAITSVNSEL--LGLKNHIVTPYFYYSVYNYTADNN--SRSMQKYD--VNCTPVISYSNMGVCANGALVFIN-VTH--  
*Bat-CoV-1A* TTGEIFSISPCQLTTQVAAVSDK--IVGVAATKEVT--LPFDTHVSLGSFYHYKNSNAE--LCKNPSLMYGGLGVCEDGRILVNIS-RSE--  
*BtMr-AlphaCoV* TTGVVNVYVPCQLSSQAVASIDA--IVGMASSTPNVS--IDFNVTVVADNFYYL--SNSAQ--PCDQPVLYAGIGICSDGSITNST-ARR--  
*BtNv-AlphaCoV* TTGDIYSVIPCQTSMOYAVIADN--IVGVISADTATG--ITFNHTIATPMFYYS--TNTDR--NCTEPVLTYATMGICADGAIGYVQ-PRV--  
*Porcine-EDV* TSGAVYSVTPCSFSEQAAYVNDD--IVGVISSLSNST--FNNTRELPGFFYH--SNDGS--NCTEPVLVYSSNIGVCKSGSIGYVP-SQY--  
*BtCoV512* TTGQIYSVTPCQLVQQAFVEDR--IVGVISSANNTG--FFNSTRTFPGFYH--SNDTT--NCTSPRLVYSSNIGVCTSAGIQLLS-PKA--  
*HKU2* ILGKVYEVAPCFEASYDVLFRSS--SSFGLLYRSFDCN--QLRISASRF--AE--RLLPSHNGTATALGCLFNATYAPNDTMVNCTNPL---GDGFCADLL--SNVVVRR--  
*NL63-related* TDSTVYTVPACATATQLAQVKQK--VLGAITAVKNDS--FGFNSTLTLPLFYH--SNGKV--NCTEPILVYSSIGICPDGTMIQIK-PVE--  
*FCoV* SDGVIYSVTPCDVSAQAVIDGA--IVGAMTSINSEL--LGLTHWTTTPNFFYYSIYNTSERTR--GTAIDSND--VDCEPVITYSNIGVCKNGALVFIN-VTH--  
*TGE* SDGVIYSVTPCDVSAQAVIDGT--IVGAITSINSEL--LGLTHWTTTPNFFYYSIYNTNDRTR--GTAIDSND--VDCEPVITYSNIGVCKNGAFVFIN-VTH--  
*OC43* ITNRTFMIRSCYSGRVSAAFHAN--SSEPALLFRNIKCN--YVFNNSLT--RQLQPINYFDSYLGCVVNAYNSTAISVQTCDLTV---GSGYCVDYS---KNRRSRGAIT  
*MERS-CoV* DDGNYYCLRACSVVPVSVIYDKE--TKTHATLFGSVACE--HISSTMSQYSRSTRSMLK--RRDSTYGPLQTPGVCLGLVNSS-LFVEDCKLPL--GQLCALPDTPSTL-TPRSVRSP  
*HKU1* VTNKTYNIFPCYAGRVSAAFHQN--ASSLALLYRNLKCS--YVLNNIS--LTTQPYFDSYLGCVFNADNLTDYSSVSCALRM--GSGFCVDYNSPSSSSRRKRSIS  
*Covid-19* QTLEILDITPCSFGGVSVITPGNTNSQAVLYQDVNCT--EVPAIHADQLPTWRVYSTG---SNVFQTRAGCLIGAEH-V-NNSYECDIPI--GAGICASYQTQTNSPRRAR-SVA  
*Bovine-CoV* LTNRTFMIRSCYSGRVSAAFHAN--SSEPALLFRNIKCN--YVFNNTLS--RQLQPINYFDSYLGCVVNADNSTSSVQTCDLTV---GSGYCVDYS---TKRRSRRSIT  
*HKU4* DDGNYYCVRPCSVVPVSVIYDKS--TNLHATLFGSVACE--HVTTMMSQFSRILTQSNLR--RRD-SNIPLOTAVGCVIGLSNNS-LVVSACKLPL--GQLCAVPPVSTFR-SY---SA  
*HKU5* DNGNYYCVRPCSVVPVSVIYDKA--SNSHATLFGSVACS--HVTTMMSQFSRMTKTNLL--ART-TPGPLQTTVGCAMGFI NSS-MVVEDCQLPL--GQLCAPIPPTSSR-VRRATSGAS  
*HKU9* INSTTYSIMPCTVTPVSV--GYHPNFERALLFNGLSCS--QRSRAVTE-PVSLWASATA--QDAFDTPSGCVNVELRNTTIVNTCAMPI--GNSLCFINGSIATA-NAD--SLP  
*SARS-CoV* KTSEILDISPCAFCGGVSVITPGTNASSEVAVLYQDVNCT--DVSTAIAHADQLTPAWRIYSTG---NNVFQTOAGCLIGAEH-V-DTSYECDIPI--GAGICASYHTVSL---LRSTS  
*HKU24* LTNRTYMIROSCYSGRVSAAYHSD--TDEPALLYRNLKCS--YVFNNNIS--ISSRSVIRYFDSYLGCVVNADDDIAEAVGSCNLTV--GSGYCVDYS---STWRAKRDLN  
*Rat-CoV* VTNKTYLLRSCYSGRVSAAYHSD--APEPALLYRNLKCD--YVFNNNIS--REETPLNYFDNYLGCVNADNSTEHVDACDLRM--GSGLCVNYS---TAHRARRSVS  
*Bat-BCoV* YNNTIYGILPCAQAQVSIVTAGSNHD--NVAVVYNGLCN--AFQSKMQG-VDNTEWSVYTNF---SSPIDTPVGCLTGAIQAO-ENSTEQFAL--GLDSCVNYTADTRL-RT-AR-AAD  
*Hedgehog-CoV* KNGTYYCMAPCVSVPVSVIYDKS--SDVHATLYSSVECN--HIKSVATVFSRQTESKLR--S-SDNGLLQTAVGCVIGFRNTS-DTVEDCTLPL--GQLCAKPPSFSSR-STN---VN  
*GCCDC1* AGDKVYSITPCVSVPVISV--GYDPGHQRALVFNGLDCS--ARANAVSM-PASEYWTAAAST--TARGSEPVLDTPSGCVVNVNNTCTHTVSVCEMPI--GNSLCLVSNFTCSD-VATASLSPN  
*HKU14* LTNRTFMIRSCYSGRVSAAFHAN--SSEPALLFRNIKCN--YVFNNFTA--RQLQPINYFDSYLGCVVNADNSTASTVQTCDLTV---GSGYCVDYS---TQLRSRAIT  
*MHV* SNRFGAISASLOEILTRLEAVEAKAOIDRLINGRLTALNAYISKOLSDSTLIKVSAAQAIKEVNECVKSOTTRINFCGNGNHILSLVONAPYGLYFIHF SYVPISFTTANVSPGLCISGDRGLAPKA

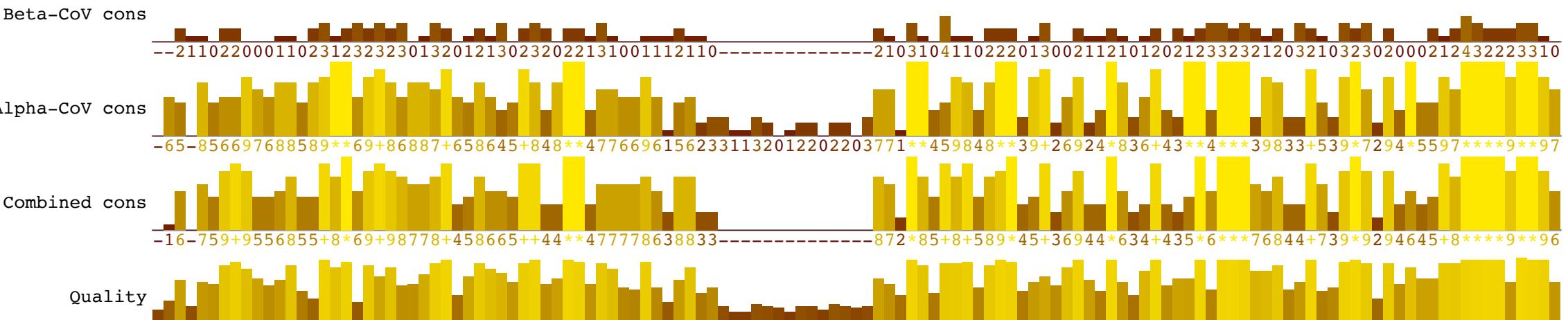
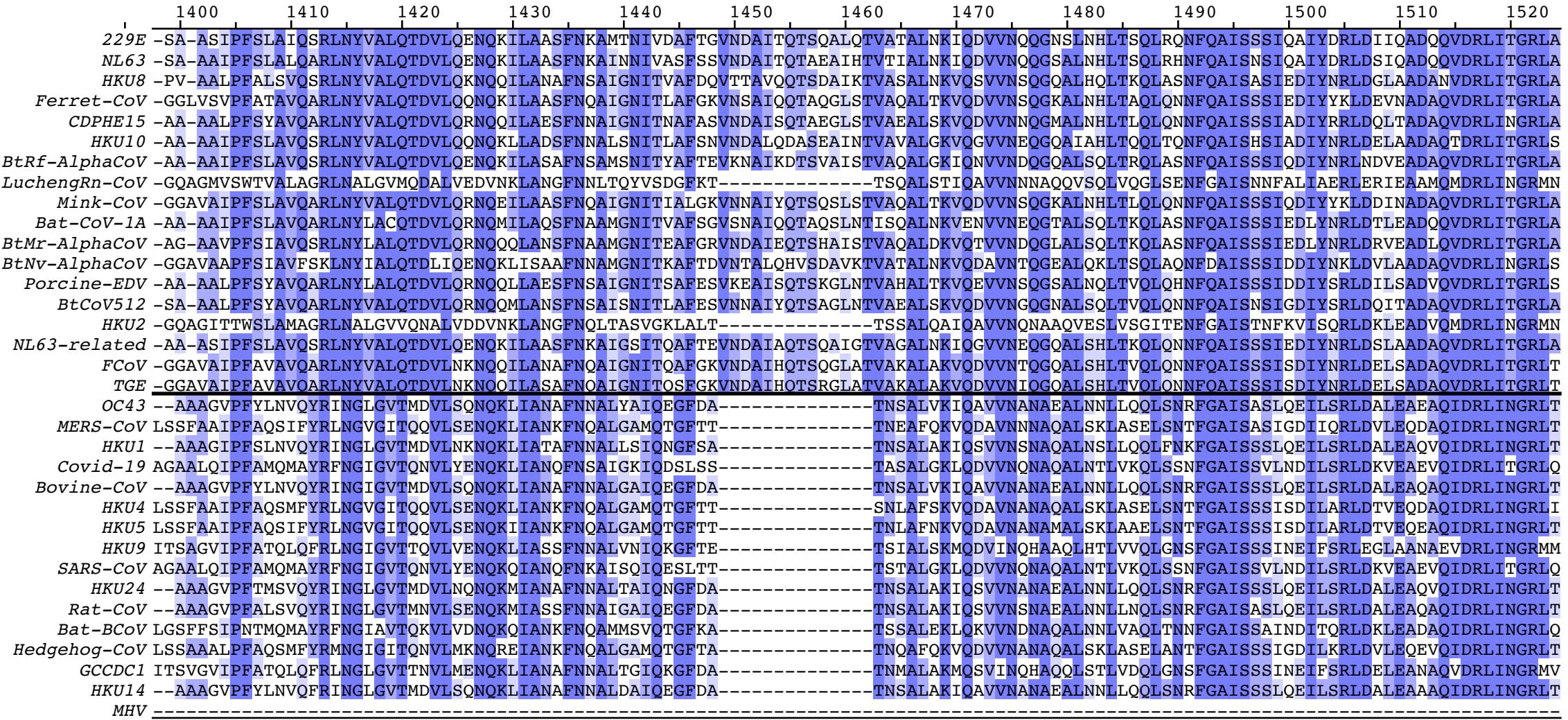


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229E -----V-SYDSVSAIVT-ANLSIPSNWTSVQVEYLQITSTPIVVDCSTYVCNGNVRCEVLLKQYTSACKTIEDALRNSAMLESADVSEMLTFDKKAFTLAN---VSSF-G-DY  
 NL63 -----S-SDNGISAIIT-ANLSIPSNWTSVQVEYLQITSTPIVVDCATYVCNGNPRCKNLLKQYTSACKTIEDALRNSAHLETNDVSSMLTFDSNAFLSAN---VTSF-G-DY  
 HKU8 -----V-RGEPDPAISM-GNISVPSNFTVSQIYQVEYLQMSIRPVSIDCAMYVCNGNPHCTRLLQOYISACRTIEEALQLSARLESFEVNSMLTVSETALDLVN---ISTF-GGDY  
*Ferret-CoV* -----S-EN-PVQPIST-GNVTIPSNFTISVQVEYLQMSSEAVSIDCAQYVCNGNPRCNRLLAQYISACHAIEQALITSTRLEALESMISISDNALALAT---VETFNSSDY  
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 HKU10 -----Y-QPAPVSPPIAT-TNISIPLNFTISIQTTEYIQMYSQSVSVDSCMYVCNGNVRCLRLLTQYATACRNIESALQLSARLESLEVSSVSVSENALKLAN---ITHF-D-SY  
*BtRf-AlphaCoV* -----T-EPEPVAPIIT-GNLSIPSNWTSVQVEYLQMYNKPVSVDCQTYVCNGNPRCLKLLTQYATACRTIEEALQLSARLESVEVANMAVSDEAVQLAN---VSYF-D-TY  
*LuchengRn-CoV* -----IFIQTH-DDTYARPILTAQQIELPIDHYVSVKEQFIQTSTPKFDVDCERYICDVSSDCRELLVKYGGYCSKILADIKSSSIQLDYQILGLYKTLAVDFKV---PDI---DFGDF  
*Mink-CoV* -----T-NG-DVQPIST-GNVTIPSNFTISVQMEYVQVSTELVSIDCARYVCNGNARCGRLLSQYVSACQTIEQALAMGSRLESMELESMSISENALKLAS---VEEFNSSAM  
*Bat-CoV-1A* -----D-T-FVPSAVIS-GNITIPANFSFVVPQPEYIQIMTKPVSVDCSVYVCNGNPRCLQLLTQYASVCRTEEPQLQNLARLEALELTSMIIIVQDKTLKLGV---ASNFN-DTF  
*BtMr-AlphaCoV* -----A-AADPVPVVIS-GNISVPTNFTFSVQVEYIQMLKPVTVDCSVYVCNGNPRCLQLLAQYASACRTIEQALQLSARLESVEVNSMISISQEALALGV---IDNFK-HDF  
*BtNv-AlphaCoV* -----V-STTPATPITT-GNLTIPVNFTVSQIAEYVQVSLRAVVDCATYVCNGNVRCLQLLRQYVTACTSVENALALNARLESQEVAADMALAVDYAAVRSSELN-TPQFE-NGF  
*Porcine-EDV* -----G-QVK-IAPVT-GNISIPTNFSMSIRTEYQLQYNTPVSVDCATYVCNGNSRCKQLLTQYTAACKTIESALQLSARLESVEVNSMLTISEEEALQLAT---ISSFNGDGY  
*BtCov512* -----A-QPO-VQPMFO-GNISIPTNFTMSVRTEYIQLFNPKPVSVDCAMYVCNGNDRCKQLLSQYTSACKNIESALQLSARLESMEVNSMLTVSDEALKLAT---ISOFPGGY  
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*NL63-related* -----T-TPQVVAPIVT-ANISIPLNFTTSVQVEYLQLSSRPVSVDCATYVCNGNPRCLTLLTQYSSACKTAAEALQLTARLEASEVNSMIKSLPTAIDNANKLGVSTYQ-G-GF  
*FCoV* -----S-DG-DVQPIST-GNVTIPTNFTISVQVEYMQVYTPVSIDCARYVCNGNPRCNKLLTQYVSACQTIEQALAMGARLENMEVDSMLFVSENALKLAS---VEAFNSTEN  
 TGE -----S-DG-DVQPIST-GNVTIPTNFTISVQVEYIQVYTPVSIDCARYVCNGNPRCNKLLTQYVSACOTIEQALAMGARLENMEVDSMLFVSENALKLAS---VEAFNSSET  
 OC43 TG---YRFTNFEPPFTVNSVNDSEPVGGLYEIQIPSEFTIGNMVEFIQTSSPKVTIDCAAFVCGDYAACSQLVEYGSFCNDINAILTEVNELLDTTQLQVANSLMNGVTLSTKLKGVNFNVDDI  
*MERS-CoV* GE--MRLASIAFNHP---I-QVDQLNSSYFKLISIPTNFSFGVTQYEIQTTIQLKVTVDCKQYVCNGFKCEQLLREYGQFCSKINQALHGANLRQDDSVRNLFAVKSSQSS---PIIPG---FGGDF  
 HKU1 AS---YRFVTPEPNVSFVNDSIESVGGLYEIKIPTNFTIVGQEEFIQTNSPKVTIDCSLFVCSNYAACHDLLSEYGTFCNDINSILDEVNGLLDTTQLHVADTLMQGVTLSSNLNTLHFVDVNI  
*Covid-19* SQ---SIIAYTMS---LGAENSVAYSNNIAIPTNFTISVTEILPVSMTKTSVDCTMYICGDSTECNSNLLQYGSFCQTLNRALTGIAVEQDKNTQEVAQVKQIYKTP---PIK---DFGGF  
*Bovine-CoV* TG---YRFTNFEPPFTVNSVNDSEPVGGLYEIQIPSEFTIGNMEEFIQTSSPKVTIDCSAFVCGDYAACSQLVEYGSFCNDINAILTEVNELLDTTQLQVANSLMNGVTLSTKLKGVNFNVDDI  
 HKU4 SQ--FQLAVLNFTSP---I-VVTPINSSGFTAAIPTNFSFSVTQYEIETSIQKVTVDCKQYVCNGFTRCEKLLVEYGQFCSKINQALHGANLRQDESVYSLYSNIKTSTQ---TLEYG---LNGDF  
 HKU5 DV--FQIATLNFNTSP---L-TLAPINSTGFVVAVPNTFTFGVQEFIEETTIQKITVDCCKQYVCNGFKCEDLLKEYGQFCSKINQALHGANLRQDESIANLFSSIKTQNTQ---PLQAG---LNGDF  
 HKU9 R---LQLVNYDPL---YDNSTATPMTPVYWKVPTNFTLSATEEYIQTAPKITIDCARYLCGDSRCLNVLLHYGTCFCNDINKALSRVSTILDSALLSLVKELSI-NTRDEVTTFS---FDGDF  
*SARS-CoV* QK---SIVAYTMS---LGADSSIAYSNNIAIPTNFSISITTEVMPVSMAKTSVDCNMYICGDSTECANLLQYGSFCQTLNRALSGIAAEQDRNTREVFAQVKQOMYKTP---TLK---YFGGF  
 HKU24 TG---YRFTNFEPPFTVNLDSVESVGGLYEIQIPTEFTIGNLEEFVQTTSPKVTIDCAAFVCGDYAACREQLVEYGSFCNDINTILNEVNSMIDTSQYQLASTLMNGVTLSSRLKGIFNQDDI  
*Rat-CoV* TG---YKLTTFEPPFTVSIVNDSVESVGGLYEMQIPTNFTIASQQEFIQTRSPKVTIDCAAFVCGDYTVCRQQLVDYGSFCNDINAILGEVNNLIDTMQLQVASALIQGVTLNSRLADGISQIDD  
*Bat-BCoV* RALTFNYMYFAQELP---SQSEQSVAFDNSTLQIPIINFETIGIEYETIPVTPMPKVTVDCAQYVCGDNSECRTLLVQYQGTFCEITNDALRGVALQQDVNQKDFVSSVRRLAKVSSPMNLG---NLNGF  
*Hedgehog-CoV* NT--FALVHMLFQNP---L-KVSVLNSTEFQVSIPQNFSFGITEEFIETSIQKVTVDCKQYVCNGFERCEQLLVQYQGQFCSKINQALHGVNLRQDDATKSLFEDIKVQPSV--PLMTS---LTGDF  
 GCCDC1 L----LSLVVYDPT---D-AGLKVLTPVYWVSIPTNFTLAATTEYIQTAPKVNVDCKVYLCGDSERCIDVLSQYGAFCEDVNKALADVSAIIDSSMVTMSELTA-GVMWSETPQA---NVGSY  
 HKU14 TG---YRFTNFEPPFTVNAVNDSLQPVGGLYEIQIPSEFTIGNMEEFIQTSAPKVTIDCAAFVCGDYAACSQLVEYGSFCNDINAILTEVNELLDTTQLQVANSLMNGVTLSTKLKGVNFNVDDI  
 MHV **GYFVODDGEWKGFTGSSYYYPEPITDKNSVIMSSCAVNYTKAPEVFLNTSIPNPPDFKEELDKWFKNOTSIAPDLSLDFEKLNVTLLDTYEMNRIODAIKKLNESYINLKEVGTYEMYVKWPWYVWL**







1530 1540 1550 1560 1570 1580 1590 1600 1610 1620 1630 1640 165

229E ALNVFVSHTLTKYTEVRASROLAQQKVNECVKSQSQRGYFCGN-GTHIFSLVNAAPEGLVFLHTVLLPTQYKDVEAWSGLCV--DGINGYVLRQPNLALYKEG-----NYYRITSRIMFEPRIP

NL63 ALNAFVSQVLNKYTEVRGSRRLAQQKINECVKSQSQRGYFCGN-GTHIFSLVNAAPEGLVFLHTVLLPTQYKDVEAWSGLCV--DGIGYGVLRQPNLVLYSND-----GVFRVTSRVMFQPRLP

HKU8 ALNAFVTQTLTKYTEVRASRLLAQEKINECVKSQSTRYGFCGN-GTHLFSIPNAAPEGIMLFHTVLVPTEYVSVTAWSGYCH--NG-VGYAVKDVGNQLFQFN-----NTFYITPRNMYPQPRTP

*Ferret-CoV* SLNAFVTQTLTNQAQIRASRQLSKEKINECVRSSRFGFCGN-GTHLFSLANAAPRGVMLFHTVLLPTSYKTVTAWSGVCAISNNKTMGFIVKDVDSTLTFKNH-----DDKFYLTPRTMYEPRVA

CDPHE15 ALNAFVSQTLTKYQSVQASRSLAKQKINECVLSQSPPRYGFCDGGGRHVTQAAPOQILFLHTVLRPTGSVNVTAAAGICV--DGA-GYALNQPGLVLIYQD-----GTYLITPRVMFEPQRP

HKU10 ALNAFVTQTLTKYTEVRASRQLAQEKINECVKSQSRYGFCGN-GTHLFSIVNNAPNGLMFHTVLVPTVYVTVSAWAGVCF--NNRA-YILRDVESVLF-NN-----NGYYVTNRKMYEPRVP

BtRf-AlphaCoV ALNAFVTQTLTKYTEVRASRQLAQEKINECVKSQSRYGFCGN-GTHLFSISNAAPQGIIFFHTVLLPTEYQTVTAWSGVCV--NNHYGLVLRNVDALYKRN-----DSYFITSRDMYEPRVP

LuchengRn-CoV ILQNFTVNTYKLKISELKSOQALAQSLINECVYAQSSRNGFCGD-GLHLFSLMQRAPDGIMFFHYTLKPNNTIIVETTPGLCLSND---VCIAPKDGFLFVRLSTA-----RDTDWHFTTRNRSPEPI

Mink-CoV ALNAFVTQTLTRQAEVRASRQLAKQKVNECVRSQSSRFGFCGN-GTHLFSLANAAPNGMVFHTVLVPTAYQTVTAWSGICASDGDRFTGLIVKDVDSTLFRNY-----NNSFYLTPTRTMYQPRIA

Bat-CoV-1A ALNAFVTQQLRYSEVRASRQLAQEKINECVKSQSRYGFCGN-GTHVFSVANAAPDGIMFLHANLVPТАFIEVSAGAGVC--DGR-ALVLRGRDEVLFQKP-----DTDQYLITPRLLFEPV

BtMr-AlphaCoV ALNAFVAQQLTKYTDVRASRQLAQDKINECVKSQSFRYGFCGN-GTHVFSVVNAAPDGMMFFHSVLLPTAYMEVAAFSGLCV--EGN-GYVLRDTGNVLFER-----NGQYLITARKMFEPV

BtNv-AlphaCoV ALSTFVSAQLVKYSEVKASRNLLAMQKVNECVKSQSRLGFCGN-GTHLFSMVTGAPDGLMFLHTVLLPTEYKEVAAWAGLCV--GGK-AFVLRDVQLLLFIR-----IIQYLVTSRNMYQPRVP

*Porcine-EDV* ALNAFVAQTLTKYTEVQASRKLAQQKVNECVKSQSQRGYFCGGDGEHIFSLVQAAPQGLLFLHTVLVPGDFVNVLIAAGLCV--NGETIALTREPGLVLFTHE-LQTYTATEYFVSSRRMFEPRKP

BtCov512 ALNAFVAQSLTKYAEVQASRTLAKQKVNECVKSQSPPRYGFCDGEGEHIFSLTQAAPOGLMFLHTVLVPGNFINVATVGLCV--DETIAMTLRQSGFVLFVQN-----GNYLVSPRKMFEPRRP

HKU2 VLQLFVTNTYKLKIAELRNTHRYVQSLINECVYAQSLRNGFCQ-GLHVLSLMQAPSGIMFFHYSLIPNNNTIVKTTPLCESDELGSKCIVAKDGLVLS--A-----NLSYWQWSPRNLKYKPENL

NL63-related ALNSFVTQTLTRQTVRASRELAMQKINECVKSQSDRYGFCGN-GTHLFSIANAAPEGLLFLHTVLVPPDDVTVTAWSGICF--GGDKGFIILRDFQTLIKYD-----NKYKVTSRKMFQPRNA

FCoV ALNAFVSQTLTRQAEVRASRQLAKDKVNECVRSQSRQYGFCGN-GTHLFSLANAAPNGMIFFHTVLLPTAYETVTAWSGICASDGDRFTGLVVKDVDQLTLFRNL-----DDKFYLTPRTMYQPRVA

TGE ALNAFVSOTLTRQAEVRASROLAKDKVNECVRSOSORFGFCGN-GTHLFSLANAAPNGMIFFHTVLLPTAYETVTAWSGICASDGDRFTGLVVKDVDVOLTLFRNL-----DDKFYLTPRTMYQPRVA

OC43 ALNAYVSQQLSDSTLVKFSAAQAMEKVNECVKSQSRRINFCCN-GNHIIISLVQNAPYGLYFIHFVSYVPTKVTARVSPGLCIA----DRGIAPKSGYFVN-----NTWMYTGSYYYYPEPI

MERS-CoV TLNAFVAQQLVRSESAALSAQLAKDKVNECVKAQSKRSGFCQ-GTHIVSFVVNAPNGLYFMHGYYPSNHIEVVSAYGLCDAAN-PTNCIAPVNGYFIKTNNTR-----IVDEWSYTGSSFYAPEPI

HKU1 ALNAYVSQQLSDISLVKFGAALAMEKVNECVKSQSRRINFCCN-GNHILSLVQNAPYGLLFMHFSYKPISFKTVLVSPGLCISG-----DVGIAPKQGYFIKHN-----DHWMFTGSSYYYYPEPI

Covid-19 SIQTYVTQQLIRAAEIRASANLAATKMSECVLGQSKRVDGCK-GYHLMSPQASPHGVVFLHTVYVPAQEKNFTTAPAIICHG-----KAHPREGVVFVSN-----THWFVTOQRNFYEPQII

Bovine-CoV ALNAYVSQQLSDSTLVKFSAAQAMEKVNECVKSQSRRINFCCN-GNHIIISLVQNAPYGLYFIHFVSYVPTKVTAKVSPGLCIA----DRGIAPKSGYFVN-----NTWMFTGSGYYYYPEPI

HKU4 SLNAFVSQQLVRSETAARSAQLASDKVNECVKSQSRRNGFCGS-GTHIVSFVVNAPNGFYFHGVYVPTNTVNTAAYGLCN----PPLCIAPIDGYF1TNQTTT-----SVDTEWYTGSFYKPEPI

HKU5 SLNAFVAQQLVRTEAAARSAQLAQDKVNECVKSQSRRNGFCGT-GTHIVSFAINAPNGLYFFHGVYQPTSHVNATAAYGLCNEN-PQKCIAPIDGYFVNLQTTSTVADSDQOWYYTGSSFFHPEPI

HKU9 VLNTYVTQLLIQASEAKAONALAAQKISECVKAQSLRNDFCGN-GTHVLSIPQOLAPNGVLFIHAYPTPEYAFVQTSAGLCHNG-----TGYAPRQGMFVLPNNT-----NMWHFTTMQFYNPVNI

SARS-CoV SLQTYVTQQLIRAAEIRASANLAATKMSECVLGQSKRVDGCK-GYHLMSPQAAAPHGVVFLHTVYVPSQERNFTTAPAIICHG-----KAYFPREGVVFVNG-----TSWFITQRNFFSPQII

HKU24 ALNAYVSQQLSDITLVKFSASQAIKEVNECVKSQSRTNFCCN-GNHILSLVQNAPYGLYFIHFVSYQPTKTTAYVSPGLCLAG-----DVGVAPKSGYF1KVN-----DKWMFTGSGAYYHPEPI

Rat-CoV ALNAYVSKHLSDMTLIKVSAQAIKEVNECVKSQSRRINFCCN-GNHILSLVQNAPYGLYFIHFVSYVPTSTTVNVSPGLCISG-----DRGLAPKAGYFVQDH-----GEWKFTGSYYYYPEI

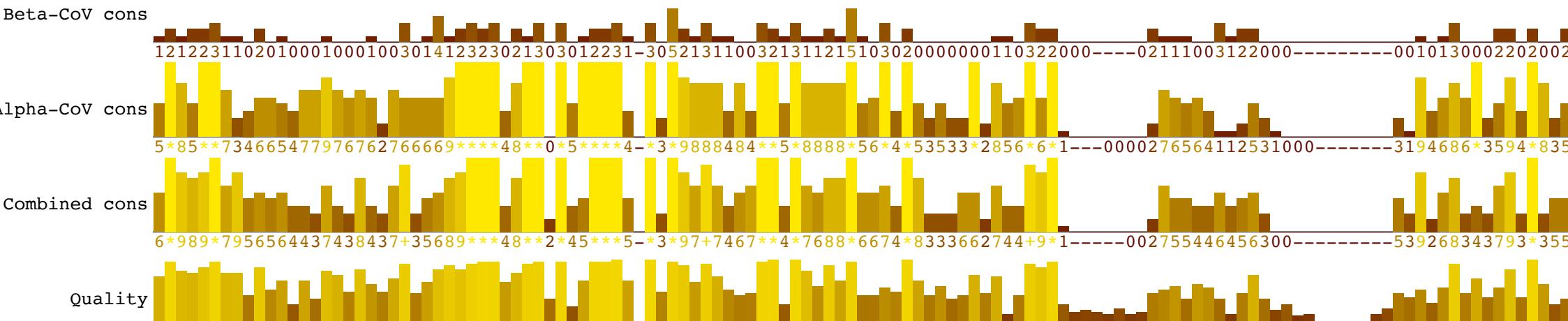
Bat-BCoV VLQFTVTQQLIRASEIRASAKLAAQKMSECVQGQSQRLDFCGR-GMHLMSFPQAAPYGMVFLHVLYKPTDYINVTTVPAICSN-----VAYFPVDPGVFVYYN-----NSLMFTKRNYFEPEII

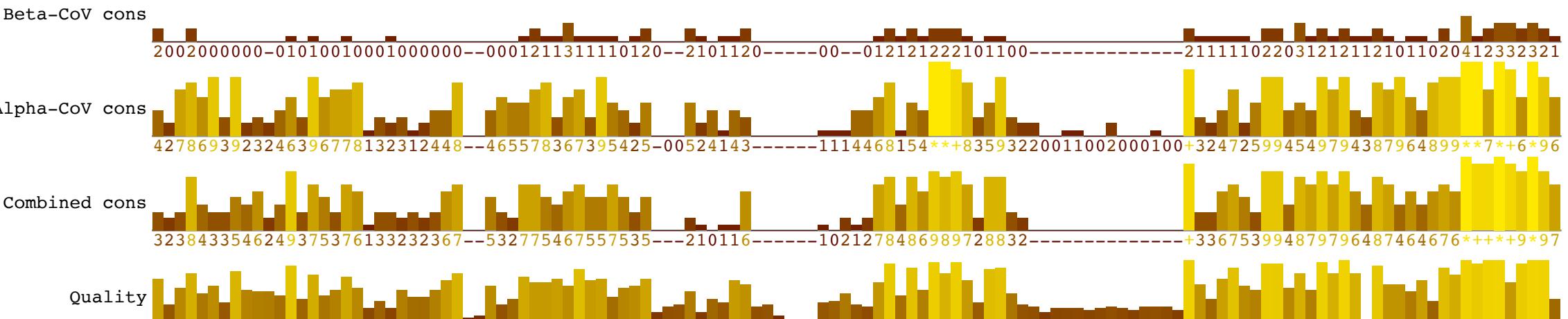
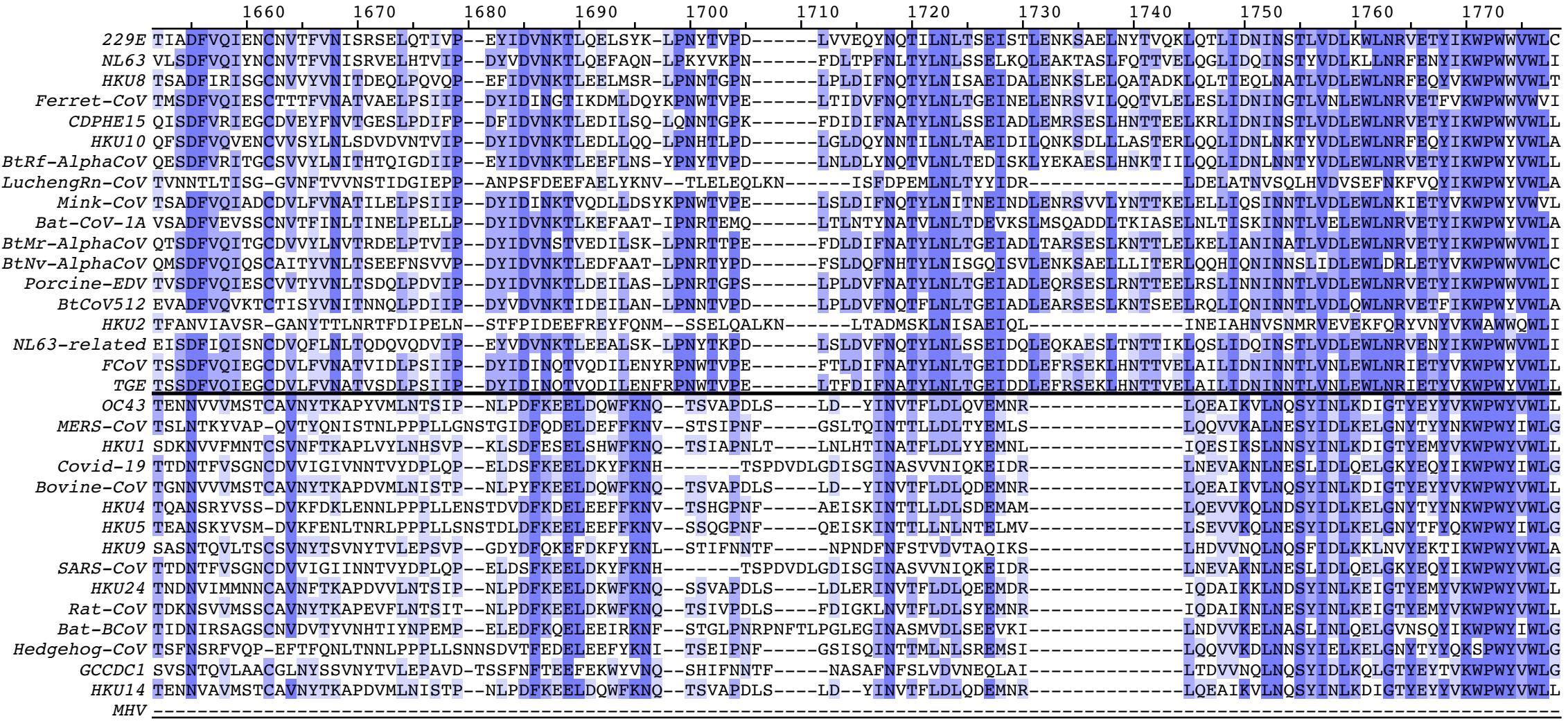
*Hedgehog-CoV* SLNAFVSQQLVRSEAAARSSQLAKEKINECVKAQSTRSGFCQ-GTHIVSFVINAPNGFYIHGVYHPQDYVNQTAAYGLCDSS--SSKCIAAKNGYFVKNDSD-----SDSWSYTGSSFYQPEPI

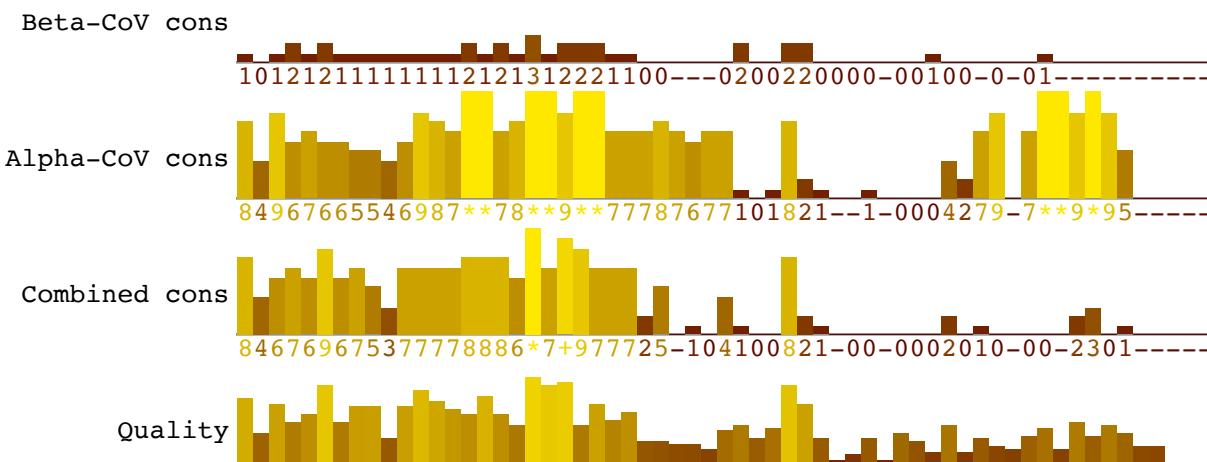
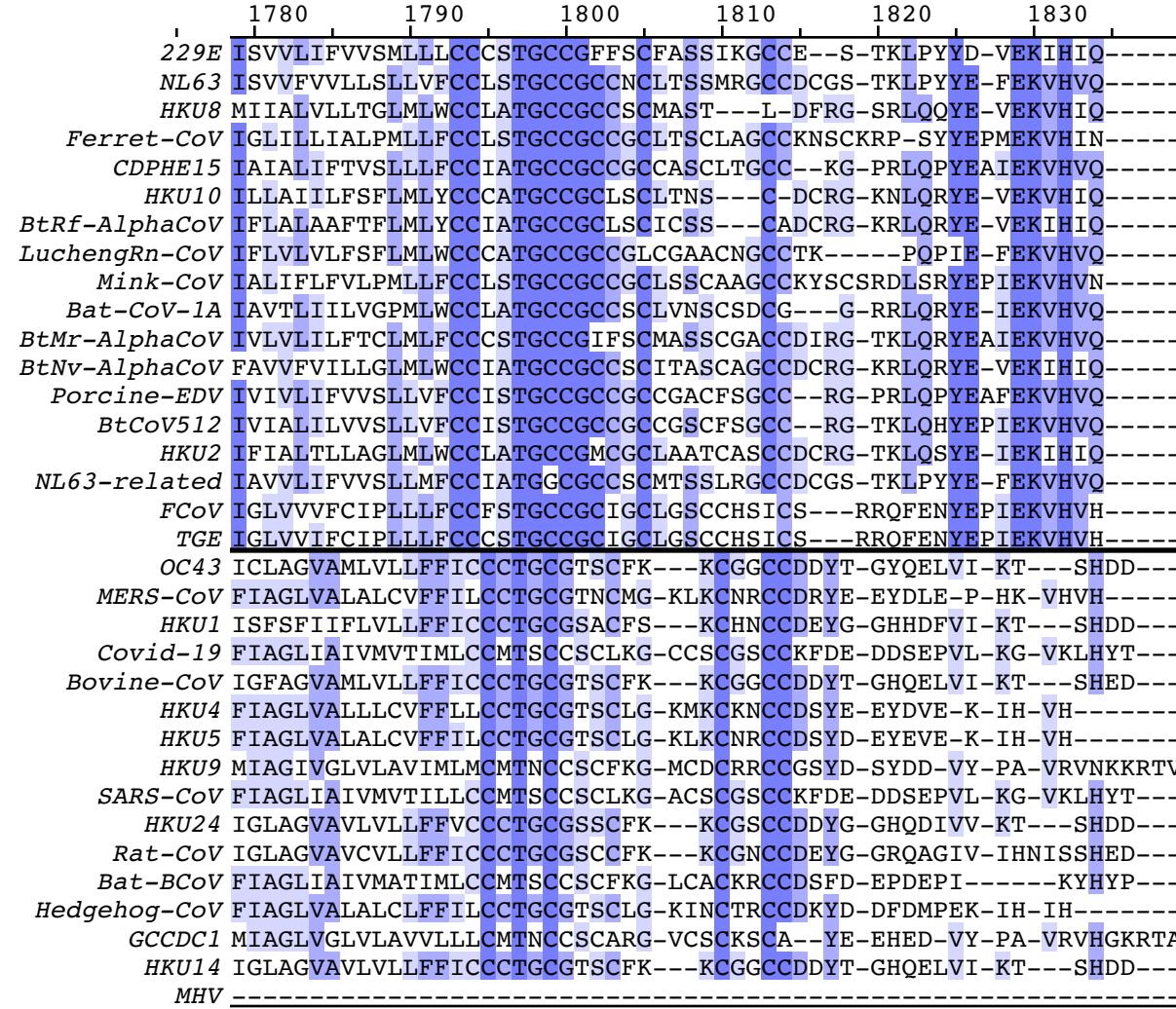
GCCDC1 VLNTYVTQLLIRASEVKAQALASQKISECVKAQSQRNDFCCN-GTHVLSIPQMAPNGVLFHYSYQPTAYNLVRTAAGLCFND-----TGYAPLGGFLVLPNNT-----DRWLFTKMSFYDPVNI

HKU14 ALNAYVSQQLSDSTLVKFSAAQAMEKVNECVKSQSRRINFCCN-GNHIIISLVQNAPYGLYFIHFVSYVPTKVTAKVSPGLCIA----DRGIAPKSGYFVN-----NTWMFTGSGYYYYPEPI

MHV -----



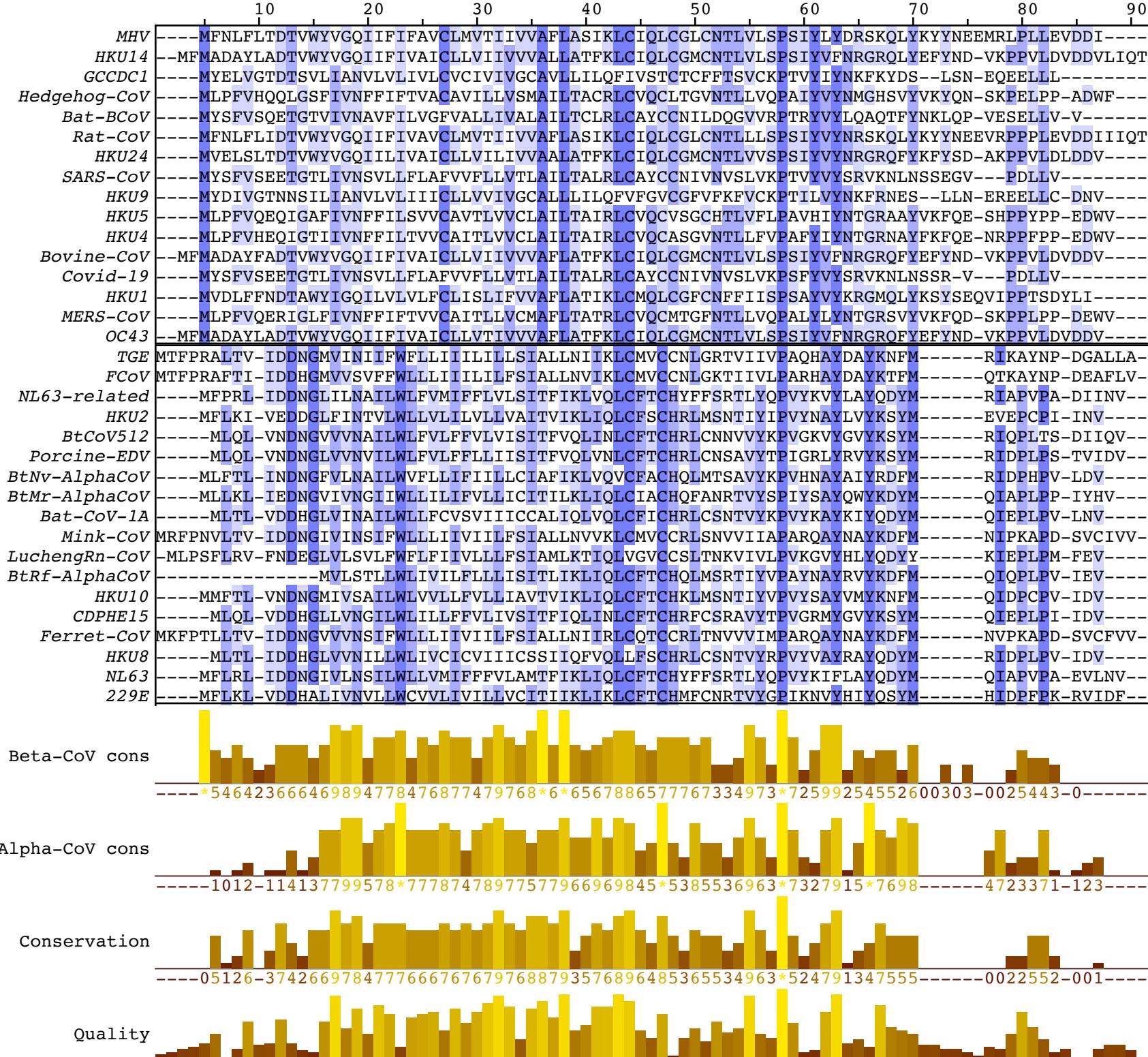




## Envelope (E) protein

10 20 30 40 50 60 70 80

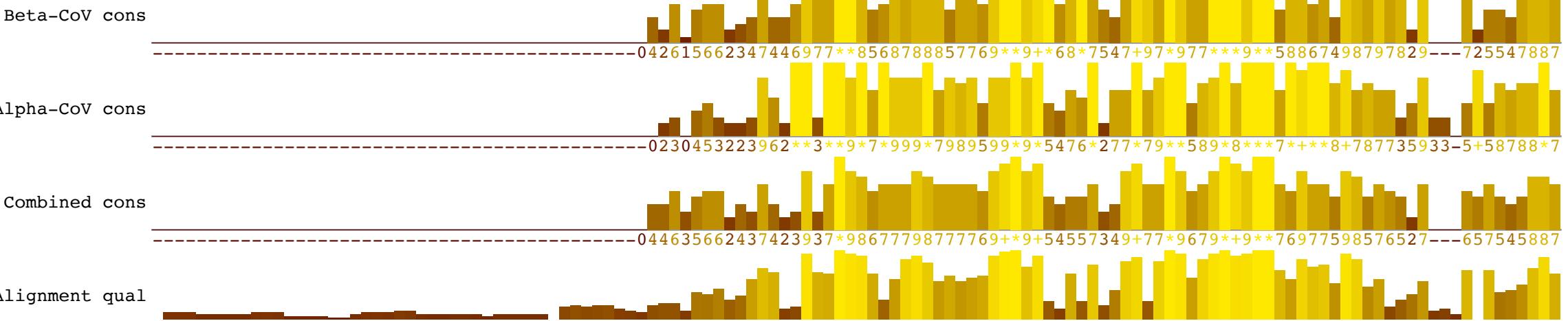
SARS-CoV-2	1 --MYSFVSEETGTLIVNSVLLFLAFVVFLLVTLAILTALRLCAYCCNIVNVSLVKPSFYVYYSRVKNLNSS--R-VPD----LLV-
SARS-CoV	1 --MYSFVSEETGTLIVNSVLLFLAFVVFLLVTLAILTALRLCAYCCNIVNVSLVKPTVYVYYSRVKNLNSS--EGVPD----LLV-
HKU1	1 --MVDLFFNDTAWYIGQILVLVLFLCLISLIFVVAFLATIKLCMQLCGFCNFFIISPSAYVYKRGMQLYKSYSSEQVIPPSTDYLI-
OC43	1 MFMADAYLADTVWYVGQIIIFIVAIICLLVTIVVVVAFLATFKLCIQLCGMCNTLVLSPSIYVFNRGRQFYEFYNDVKP-PVLDVDDV
MERS-CoV	1 --MLPFVQERIGLFIVNFFIFTVVCAITLLVCMAFLTATRLCVQCMTGFNTLLVQPALYLYNTGRSVYVKFQDSKPPLPPDEWV-
229E	1 --MFLKLV-DDHALIVNVLLWCVVLIVILLVCITIIIKLIKLCFTCHMFCNRTVYGPIKNVYHI----YQSYM HIDPFP-KRVIDF
NL63	1 --MFLRLI-DDNGIVLNSILWLLVMIFFFVLAMTFIKLIQLCFTCHYFFSRTLQYPVYKIFLA---YQDYMQIA PVP-AEV LNV

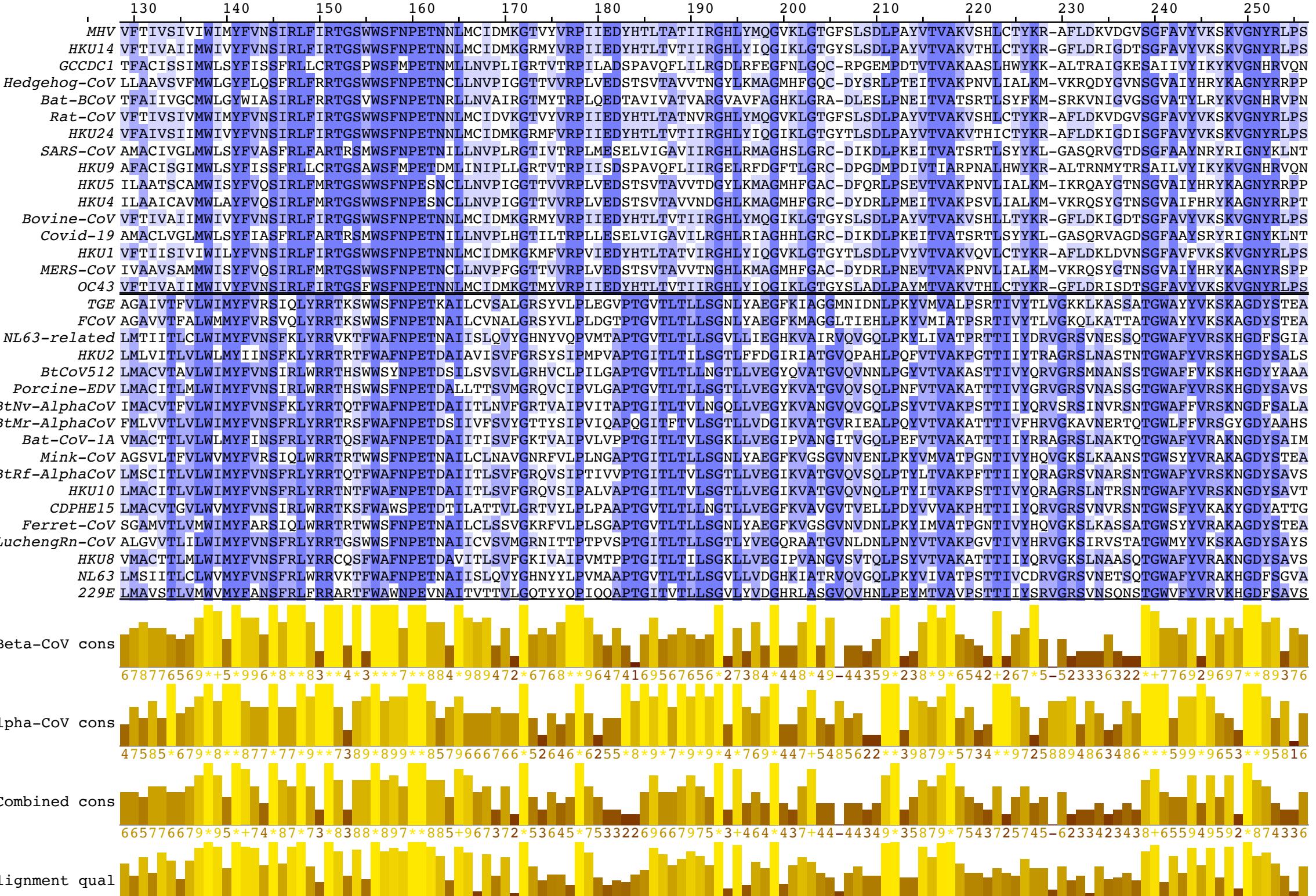


## Membrane (M) Protein

	10	20	30	40	50	60	70	80	90
<i>SARS-CoV-2</i>	1	-----MADSNGTITVEELKKLLEQWNLVIGFLFLTWICLLQFAYANRNRLFYI	I	KLIFLWLLWPVTLACFVLAAV	--YRINWITGGIAIAM				
<i>SARS-CoV</i>	1	-----MADNGTITVEELKQLLEQWNLVIGFLFLAWIMLLQFAYSNRNRLFYI	I	KLVFLWLLWPVTLACFVLAAV	--YRINWVTGGIAIAM				
<i>HKU1</i>	1	---MNKSFLPQFTSDQAVTFLKEWNFSLGVILLFITIILQFGYTSRSRMFVYLIKMI	I	LWLMWPLTITLTIFNCF	--YALNNNAFLAFSIVF				
<i>OC43</i>	1	MSSKTPAPVYIWTADEAIKFLKEWNFSLGIIILLFITIILQFGYTSRSRMFVYVIKMI	I	LWLMWPLTIILTIFNCV	--YALNNVYLGLSIVF				
<i>MERS-CoV</i>	1	-----MSNMTQLTEAQIIAIKDWNFIAWSLIFLLITIVLQYGPSRSMTVYVF	KMFVWLWLPSSMALSIFS	AV--YPIDLASQIISGIV					
<i>229E</i>	1	-----MSND-NCTGDIVTHLKNWNFGWNVIL	TIFIVILQFGHYKYSRLFYGLKMLVWLWLPVLALSIFDTWANWDSNWAFVA	FSFLM					
<i>NL63</i>	1	-----MSNSSVPLLEVYVHLRNWNFSWNL	ILTLFIVVLQYGHYKYSRLLYGLKMSVLWCLWPLVLALSIFDCFVN	FNFNVWDWF	FGFSILM				
	100	110	120	130	140	150	160	170	180
<i>SARS-CoV-2</i>	85	ACLVGLMWLSYFIASFRLFARTRS	MWSNPETNILLNVPLHGTILTRP	PLESELVIGAVILRGHLRIAGHHLGRC-DIKDLPKEITVATSR					
<i>SARS-CoV</i>	84	ACIVGLMWLSYFVASFRLFARTRS	MWSNPETNILLNVPLRG	TIVTRP	LMESELVIGAVIIRGHLRMAGHSLGRC-DIKDLPKEITVATSR				
<i>HKU1</i>	86	TIISIVIWIYL	YFVN	SIRLFIRTGS	WWSFNPETNNL	MCIDMKGKMFVRPVIEDYHTLT	ATVIRGHL	YIQGVKLGTGYTLS	DLPVYVTVAKVQ
<i>OC43</i>	90	TIVAIIMWIVYFVN	SIRLFIRTGS	FWSFNPETNNL	MCIDMKGTMV	YRPIIEDYHTLT	VTIIRGHL	YIQGIKLGTG	YSLADLPAYMTVAKV
<i>MERS-CoV</i>	84	AAVSAMMWISYFVQ	SIRLFMRTGS	WWSFNPETNC	LLNVPGGGTTVVRPLVED	STSVTAVVTNGH	LKMAGMHFGAC-DYDRLPNE	VTVAKPN	
<i>229E</i>	84	AVSTLVMWVMYFANSF	RRLFARRTF	WA	NPEVNAITVTTVL	GQTYQP	IQQAPTGITV	LLSGVLYVDGH	RLASGVQVHLP
<i>NL63</i>	85	SIITLCLWV	WV	MYFVN	SFR	LWR	RVKTF	WAFNPETNAIISLQVY	GHNYYL
	190	200	210	220	230				
<i>SARS-CoV-2</i>	175	TLSYYKLG-ASQRVAGD	SGFAA	YSRYRIGNYKLNTDH	SSSSDNIALLVQ	--			
<i>SARS-CoV</i>	174	TLSYYKLG-ASQRVGTD	SGFAA	YNRYRIGNYKLNTDH	AGSNDNIALLVQ	--			
<i>HKU1</i>	177	VLCTYKRA-FLDKLDV	NSGF	AVFVKSKVGN	YRLPSS-KPSGMDT	ALLRA	--		
<i>OC43</i>	181	HLCTYKRG-FLD	RISDTSGF	AVVVKSKVGN	YRLPSTQKGSGMDT	ALLRNNI			
<i>MERS-CoV</i>	174	VLIALKMV-KRQSYGT	NSGV	AIYHRYKAGNYRSPP	ITA-DIEL	ALLRA	--		
<i>229E</i>	175	TTI	IYSRVGR	SVNSQN	STGWVYVRV	KHGDFSAVSSPM	SNMTENER	LLHFF	
<i>NL63</i>	176	TTIVCDR	VGR	SVNETSQTG	WAFYVRA	KHGDFSGV	ASQEGVL	SEREKLLH	LI

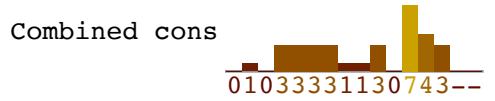
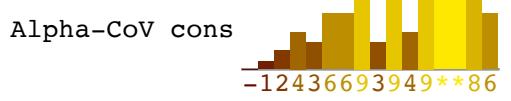
	10	20	30	40	50	60	70	80	90	100	110	120	
<i>MHV</i>				-M-S-T-T-Q-A-P-E-P-V-Y-Q-W-T-A-D-E-A-V-Q-F-L-K-E-W-N-F-S-L-G-I-I-L-L-F-I-T-T-I-L-Q-F-G-Y-T-S-R-S-M-F-I-Y-V-V-K-M-I-I-L-W-L-M-W-P-L-T-I-V-L-C-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>HKU14</i>				-M-S-S-T-T-P-A-P-V-Y-I-W-T-A-D-E-A-I-K-F-L-K-E-W-N-F-S-L-G-I-I-L-L-F-I-T-T-I-L-Q-F-G-Y-T-S-R-S-M-F-V-Y-V-I-K-M-I-I-L-W-L-M-W-P-L-T-I-I-L-T-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>GCCDC1</i>				-M-S-G-G-N-G-T-I-P-R-S-E-V-M-A-A-L-R-D-W-N-F-A-I-S-V-V-L-L-F-I-T-A-L-L-Q-W-G-Y-P-T-R-C-K-P-L-W-V-I-K-M-C-I-L-W-L-L-W-P-V-S-I-A-A-V-F-A-A-L-Y-P-I--								NDVAFGFAI	
<i>Hedgehog-CoV</i>				-M-S-N-D-T-L-S-Q-E-E-V-I-A-V-I-K-D-W-N-F-A-W-S-I-V-F-L-I-I-T-A-V-L-Q-Y-G-Y-P-S-R-S-M-T-A-Y-V-V-K-M-F-L-W-L-L-W-P-C-S-I-A-L-A-V-F-S-A-V-Y-P-I--								NLVSQIISG	
<i>Bat-BCoV</i>				-M-S-G-D-S-N-N-T-I-T-Q-E-E-L-F-K-V-V-K-E-W-N-F-A-L-G-I-M-L-F-I-F-I-T-L-L-Q-Y-A-Y-A-T-R-S-L-G-L-Y-I-L-K-L-V-L-W-L-M-W-P-L-T-F-A-C-F-V-L-A-A-I-Y-H-V--								NNVFMGFAI	
<i>Rat-CoV</i>				-M-S-S-T-T-S-A-P-Q-T-V-Y-Q-W-T-A-D-V-A-V-R-F-L-K-E-W-N-F-L-L-G-I-I-L-L-F-I-T-T-I-I-L-Q-F-G-Y-T-S-R-S-M-F-I-Y-V-V-K-M-I-I-L-W-L-M-W-P-L-T-I-V-L-C-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>HKU24</i>				-M-S-G-Q-A-T-T-P-E-P-I-Y-S-W-T-A-D-E-A-I-R-F-L-K-E-W-N-F-S-L-G-I-I-L-L-V-I-T-T-I-I-L-Q-F-G-Y-T-S-R-S-M-F-V-Y-V-I-K-M-I-I-L-W-L-M-W-P-L-T-I-I-L-V-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>SARS-CoV</i>				-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-L-F-L-A-W-I-M-L-L-Q-F-A-Y-S-N-R-N-F-L-Y-I-I-K-L-V-L-F-W-L-L-W-P-V-T-L-A-C-F-V-L-A-A-V-Y-R-I--								NWVTGGIAI	
<i>HKU9</i>				-M-S-N-N-C-T-N-T-V-P-R-P-E-V-I-A-A-L-K-D-W-N-F-A-V-S-V-I-L-L-F-I-T-V-L-L-Q-W-G-Y-P-S-R-C-K-P-I-W-V-I-K-M-F-L-I-L-W-L-L-W-P-L-S-I-A-A-V-F-A-A-I-H-P-I--								NSVAFGFAI	
<i>HKU5</i>				-M-A-S-S-N-V-T-L-S-N-D-E-V-L-R-L-V-K-D-W-N-F-T-W-S-V-V-F-L-L-I-T-I-V-L-Q-Y-G-Y-P-S-R-S-M-F-V-Y-V-I-K-M-F-V-L-W-L-W-P-A-S-M-A-L-S-I-F-C-A-V-Y-P-I--								DLASQIISG	
<i>HKU4</i>				-M-S-S-N-G-S-L-T-K-D-E-V-V-N-I-I-K-D-W-N-F-S-W-S-I-I-F-L-L-I-T-I-V-L-Q-Y-G-Y-P-S-R-S-M-M-V-Y-V-F-K-M-F-I-L-W-L-L-W-P-A-S-M-A-L-S-I-F-S-A-I-Y-P-I--								SLSSQIISG	
<i>Bovine-CoV</i>				-M-S-S-V-T-T-P-A-P-V-Y-T-W-T-A-D-E-A-I-K-F-L-K-E-W-N-F-S-L-G-I-I-L-L-F-I-T-V-I-L-Q-F-G-Y-T-S-R-S-M-F-V-Y-V-I-K-M-V-I-L-W-L-M-W-P-L-T-I-I-L-T-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>Covid-19</i>				-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-L-F-L-T-W-I-C-L-L-Q-F-A-Y-A-N-R-N-F-L-Y-I-I-K-L-I-F-L-W-L-L-W-P-V-T-L-A-C-F-V-L-A-A-V-Y-R-I--								NWITGGIAI	
<i>HKU1</i>				-M-M-N-K-S-F-L-P-Q-F-T-S-Q-A-V-T-F-L-K-E-W-N-F-S-L-G-V-I-L-L-F-I-T-T-I-I-L-Q-F-G-Y-T-S-R-S-M-F-V-Y-L-I-K-M-I-I-L-W-L-M-W-P-L-T-I-T-L-T-I-F-N-C-F-Y-A-L--								NNAFLAFSI	
<i>MERS-CoV</i>				-M-S-N-M-T-Q-L-T-E-A-Q-I-I-A-I-K-D-W-N-F-A-W-S-L-I-F-L-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-S-M-A-L-S-I-F-S-A-V-Y-P-I--								DLASQIISG	
<i>OC43</i>				-M-S-S-K-T-T-P-A-P-V-Y-I-W-T-A-D-E-A-I-K-F-L-K-E-W-N-F-S-L-G-I-I-L-L-F-I-T-T-I-I-L-Q-F-G-Y-T-S-R-S-M-F-V-Y-V-I-K-M-I-I-L-W-L-M-W-P-L-T-I-I-L-T-I-F-N-C-V-Y-A-L--								NNVYLGFSI	
<i>TGE</i>	-M-K-I-L-L-I-L-A-C-V-I-	-A-C-A-C-G-E-R-Y-C-A-M-K-S-D-T-D-L-S-C-R-N-S-T-A-S-D-C-E-S-C-F-N-G-G-D-L-I-W-H-L-A-N-W-N-F-S-W-S-I-I-L-I-V-F-I-T-V-L-Q-Y-G-R-P-Q-F-S-W-F-V-Y-G-I-K-M-L-I-M-W-L-L-W-P-V-L-A-L-T-I-F-N-A-Y-Q-V-S-R-Y-V-M-F-G-F-S-I											
<i>FCoV</i>	-F-K-Y-I-L-L-I-L-A-C-I-I-	-A-C-V-Y-G-E-R-Y-C-A-M-Q-D-S-G-L-Q-C-I-N-G-T-N-S-R-C-Q-T-C-F-E-R-G-D-L-I-W-H-L-A-N-W-N-F-T-W-S-V-V-F-L-L-I-T-I-V-L-Q-Y-G-Y-P-S-R-S-M-F-V-Y-V-I-K-M-F-V-L-W-L-W-P-A-S-M-L-S-I-F-D-C-W-A-N-F-G-K-D-D-W-V-F-F-A-F-S-L											
<i>NL63-related</i>				-M-S-T-T-M-S-N-E-S-I-P-L-T-E-V-Y-E-H-L-R-N-W-N-F-S-W-S-V-I-L-I-V-F-I-T-V-L-Q-Y-G-R-P-Q-F-S-W-L-V-Y-G-I-K-M-L-I-M-W-L-L-W-P-V-M-L-A-L-S-I-F-D-C-W-A-N-F-G-K-D-D-W-V-F-F-A-F-S-L									
<i>HKU2</i>				-M-S-S-N-S-T-E-S-V-P-V-T-E-V-L-E-H-L-R-N-W-N-F-S-W-S-N-I-I-L-T-V-F-I-A-V-L-Q-Y-G-N-M-K-Y-S-F-F-L-Y-G-V-K-M-L-I-M-W-L-L-W-P-L-V-L-A-L-S-I-F-N-A-Y-A-D-F-G-V-N-N-W-W-F-F-S-F-S-I									
<i>BtCoV512</i>				-M-S-S-N-Q-S-V-P-V-E-E-V-I-K-H-L-R-N-W-N-F-S-W-S-N-I-I-L-T-I-L-L-V-L-L-Q-Y-G-H-Y-K-Y-S-R-V-L-Y-G-L-K-M-A-I-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-A-W-S-F-N-V-N-N-W-V-F-F-A-F-S-I									
<i>Porcine-EDV</i>				-M-S-N-G-S-I-P-V-D-E-V-I-E-H-L-R-N-W-N-F-T-W-N-I-I-L-T-I-L-L-V-L-L-Q-Y-G-H-Y-K-Y-S-V-F-L-Y-G-V-K-M-A-I-L-W-L-L-W-P-L-V-L-A-L-S-I-F-L-D-A-W-S-F-Q-V-N-N-W-V-F-F-A-F-S-I									
<i>BtNv-AlphaCoV</i>				-M-S-S-S-N-Q-T-I-P-V-T-E-V-V-E-H-L-R-N-W-N-F-A-W-N-V-I-L-T-V-L-L-V-L-L-Q-Y-G-N-F-K-Y-S-K-V-L-Y-G-L-K-M-I-V-L-W-V-L-W-P-T-V-L-A-L-S-I-F-N-A-W-A-E-F-N-V-N-N-W-V-M-F-S-F-S-I									
<i>BtMr-AlphaCoV</i>				-M-S-D-N-S-T-V-P-V-A-D-V-I-E-H-L-R-N-W-S-F-A-W-N-I-L-L-T-L-L-I-A-L-L-Q-Y-G-N-F-K-Y-S-K-L-L-Y-G-L-K-M-A-L-W-L-L-W-P-C-V-I-A-L-S-I-T-N-A-W-A-D-F-N-V-N-N-W-A-Y-F-G-I-S-I									
<i>Bat-CoV-1A</i>	-M-I-N-F-I-V-L-C-C-A-F-I-G-V-I--Y-A-D-N-G-D-	-N-T-T-A-V-T-P-T-L-P-P-Q-Q-I-P-L-K-E-V-M-E-H-L-R-N-W-N-F-S-W-N-I-V-L-T-V-F-L-I-V-M-Q-Y-G-H-F-K-Y-S-T-I-L-Y-G-L-K-M-L-I-V-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-A-W-S-T-F-G-Q-K-W-V-M-F-A-F-S-I											
<i>Mink-CoV</i>	-M-E-V-L-L-V-A-A-T-A-V-L-S-V-I-P-T-S-H-A-T-N-L-Y-C-L-M-N-T-T-S-V-A-N-N-T-C-Q-N-T-N-G-I-C-D-T-C-I-N-E-G-N-L-I-W-H-L-M-N-W-N-F-S-W-S-V-L-L-I-L-F-I-T-I-L-Q-Y-G-R-P-Q-V-S-W-F-I-Y-G-I-K-M-L-I-M-W-L-L-W-P-L-V-L-A-L-T-I-F-N-A-Y-E-S-Y-T-N-S-R-Y-V-M-F-G-F-S-V												
<i>BtRf-AlphaCoV</i>				-M-S-N-N-S-T-V-P-V-E-V-V-E-H-L-R-N-W-N-F-S-W-N-V-V-L-T-V-F-L-L-V-L-L-Q-Y-G-Q-Y-K-Y-S-K-L-L-Y-G-L-K-M-L-I-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-S-W-A-S-F-G-V-N-N-W-V-F-F-A-F-S-I									
<i>HKU10</i>				-M-S-N-G-D-N-S-T-I-P-T-D-V-V-I-Q-H-L-R-N-W-N-F-S-W-N-V-I-L-T-I-F-L-V-L-L-Q-Y-G-H-Y-K-Y-S-A-V-A-L-Y-I-L-K-M-T-I-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-S-W-S-S-F-G-N-N-W-T-M-F-A-F-S-I									
<i>CDPHE15</i>				-M-A-N-G-S-V-S-E-E-I-I-H-H-L-R-N-W-N-F-G-W-N-I-I-L-T-I-F-I-V-V-L-Q-Y-G-Q-Y-K-Y-S-A-F-L-Y-G-L-K-M-L-V-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-A-Y-A-S-F-N-V-N-N-W-V-F-F-A-F-S-I									
<i>Ferret-CoV</i>	-M-I-E-I-V-L-L-V-C-S-A-I-A-A-M-V-P-T-T-A-S-G-N-R-Y-C-L-V-N-N-T-E-G-A-T-Q-N-C-S-	-I-N-D-C-I-D-E-G-S-L-I-W-H-L-M-N-W-S-F-S-W-S-V-V-L-I-V-F-I-A-I-L-Q-Y-G-R-P-Q-F-S-W-F-M-Y-G-V-K-M-L-I-M-W-L-L-W-P-L-V-L-A-L-T-I-F-N-A-Y-E-F-K-S-S-R-Y-V-M-F-G-F-S-V											
<i>LuchengRn-CoV</i>				-M-V-L-F-C-E-F-N-T-S-S-A-D-G-C-S-A-C-V-A-Q-G-C-T-G-A-G-V-T-S-A-I-T-W-H-L-L-N-W-N-F-S-W-S-V-I-L-T-V-F-V-S-V-L-Q-Y-G-N-L-K-Y-S-M-I-L-Y-I-L-K-M-I-I-M-W-L-L-W-P-I-V-I-A-L-T-I-F-D-A-I-E-T-F-K-Q-E-R-Y-V-M-F-G-F-S-I									
<i>HKU8</i>	-M-I-F-L-I-L-G-I-I-C-A-C-S-A-N-G-D-	-N-T-T-A-P-P-T-V-P-Q-S-I-P-V-E-Q-V-V-E-H-L-R-N-W-N-F-S-W-N-V-I-L-T-V-F-L-I-V-L-Q-Y-G-N-F-K-Y-S-K-L-L-Y-G-L-K-M-V-I-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-A-W-S-F-G-V-N-W-T-F-F-A-F-S-I											
<i>NL63</i>				-M-S-N-S-V-P-L-L-E-V-Y-V-H-L-R-N-W-N-F-S-W-N-L-I-L-T-L-F-I-V-V-L-Q-Y-G-H-Y-K-Y-S-R-L-L-Y-G-L-K-M-S-V-L-W-C-I-L-W-P-L-V-L-A-L-S-I-F-D-C-F-V-N-F-N-V-D-W-V-F-F-G-F-S-I									
<i>229E</i>				-M-S-N-D-N-C-T-G-D-I-V-T-H-L-K-N-W-N-F-G-W-N-V-I-L-T-I-F-I-V-I-L-Q-F-G-H-Y-K-Y-S-R-L-F-Y-G-L-K-M-L-V-L-W-L-L-W-P-L-V-L-A-L-S-I-F-D-T-W-A-N-W-D-S-N-W-A-F-V-A-F-S-F-I									





260                    270

<i>MHV</i>	NK-PSGADTALLRT--
<i>HKU14</i>	TQKGTGMDTALLRNNI
<i>GCCDC1</i>	AREDGDTIAM-FIA--
<i>Hedgehog-CoV</i>	VVVDEEL-AL-LRA--
<i>Bat-BCoV</i>	ARASEDQEDLLVVS--
<i>Rat-CoV</i>	NK-PSGADTALLRI--
<i>HKU24</i>	SQKGNGMDTALLRNNV
<i>SARS-CoV</i>	DHAGSNDNIALLVQ--
<i>HKU9</i>	TTEDGDRLAM-FVA--
<i>HKU5</i>	IIQDQEL-AL-LRA--
<i>HKU4</i>	IIQDEEL-AL-LRA--
<i>Bovine-CoV</i>	TQKGSGLDTALLRNNI
<i>Covid-19</i>	DHSSSSDNIALLVQ--
<i>HKU1</i>	SK-PSGMDTALLRA--
<i>MERS-CoV</i>	ITADIEL-AL-LRA--
<i>OC43</i>	<u>TOKGSGMDTALLRNNI</u>
<i>TGE</i>	-RTDNLSEQEKLHHLV
<i>FCoV</i>	-RTDNLSEHEKLHHLV
<i>NL63-related</i>	SQENVLSEREKLLHLV
<i>HKU2</i>	NSSDNLTENDRLLHLV
<i>BtCoV512</i>	NPTEVVTDSEKILHLV
<i>Porcine-EDV</i>	NPSAVLTDSEKVHLV
<i>BtNv-AlphaCoV</i>	TDNETMTERERLLLHLV
<i>BtMr-AlphaCoV</i>	TGTGQMTESEKLLHIA
<i>Bat-CoV-1A</i>	TMADSFSENERLLLHLV
<i>Mink-CoV</i>	-RNDNLSEHEKLHHLV
<i>BtRf-AlphaCoV</i>	SHSDSMTEDERVHLHV
<i>HKU10</i>	SSADSLTEDEKLLHLV
<i>CDPHE15</i>	HSNDKVPDSEVVLLHLV
<i>Ferret-CoV</i>	-RHDLNLEHEKLHHLV
<i>LuchengRn-CoV</i>	-SGGSLSDQEQLLHLV
<i>HKU8</i>	NPSDSYTDTDRLLLHLV
<i>NL63</i>	SQEGVL SEREKLLHLI
<i>229E</i>	<u>SPMSNM TENERLLLHFF</u>



# Nucleocapsid (N) protein

10 20 30 40 50 60 70 80 90

SARS-CoV-2 1 MSD-NGPQ-NQ-----RNAPRITFGGPDSTGSNQNGERSGARSK-----QRRPQGLPNNTA**SWFTALTQHGK**-EDLKFPRGQGV  
SARS-CoV 1 MSD-NGPQSNQ-----RSAPRITFGGPTDSTDNNQNQNGGRNGARPK-----QRRPQGLPNNTA**SWFTALTQHGK**-EELRFPRGQGV  
HKU1 1 MSYTPGHYAGSRSSSGNRSGILKKTSWADQSERNYQTF---NRGRKTQPKFTVST--QPOQGNTIPHYSWFSGITQFQKGRDFKFSDGQGV  
OC43 1 **MSFTPQKQSSSRASSGNRSGNG**-ILKWADQSDQFRNVQ---TRGRRAQPKQTATSQOPSGGNVPYYSWFSGITQFQKGKEFEFVEGQGV  
MERS-CoV 1 -----MASP-----AAPRAVSFADNNNDITNTN-----LSRGRGR-----NPKPRAAPNNTVSWYTGLTQHGK-VPLTFPPGQGV  
229E 1 -----MATVKWADASEPQ-----RGRQ-----GRIPYSLYSPLLVDS-EQPWKVIPRNLV  
NL63 1 -----MASVNWADDRAA-----RK-----KFPPPSFYMPPLLVSSEDKAPYRVIPRNLV

100 110 120 130 140 150 160 170 180

SARS-CoV-2 73 PINTNSSPDDQI**G**YRRATRR-IRGGD**G**KMKDLSPRWFYLLGTGP**E**AGLPYGANKD**G**IIVVATEGALNTPKD**H**IGTRNPANNAIAVLQLP  
SARS-CoV 74 PINTNSGPDDQI**G**YRRATRR-VRGGDG**G**KMKELSPRWFYLLGTGP**E**ASLPYGANKEGIVVVATEGALNTPKD**H**IGTRNPNNNAATVQLLP  
HKU1 86 PIAFGVPPSEAKGYWYRH**R**HSRRSF**K**TAD**Q**QQ**K**OLL**P**RWFYLLGTGP**E**ANASYGESLEG**V**FVANHQADTSTPSDVSSRDPTTQEAIPTRF**P**  
OC43 87 PIAPGVPA**T**EAKGYWYRH**R**NSRF**K**TAD**Q**NQR**Q**LL**P**RWFYLLGTGP**E**AKD**Q**Y**G**TID**D**G**V**W**A**SNQ**Q**AD**V**NT**P**AD**I**VD**R**DP**S**DEAIPTRF**P**  
MERS-CoV 64 ELNANSTPAQNAGYWR**R**QDR**K**-INT**G**NG-**I**K**Q**LAPRWY**F**YY**T**TGTGP**E**AA**L**PFRAV**K**D**G**IV**V**V**H**EDGAT**D**APS-TF**G**TRNPNN**D**SAIV**T**QFA  
229E 45 PVNKKD-KNKLIGYWNV**Q**KR--FRTRKG**K**RVD**L**SP**K**LHFYLLGTGP**H**KDAFK**R**ERVE**G**V**V**V**A**VG**A**KT**E**PT-GY**G**V**R**R**K**N**S**EP**E**IP**H**-F-  
NL63 43 PI**G**KGN-K**D**E**Q**I**G**Y**W**N**V**Q**E**R-WRM**R**R**C**Q**R**V**D**IPPKV**H**FYLLGTGP**H**K**D**LK**F**Q**R**SD**G**V**V**W**V**A**K**E**G**AK**T**V**N**T-S**L**G**N**R**K**R**N**Q**K**P**L**E**P**K-F-

190 200 210 220 230 240 250 260 270

SARS-CoV-2 163 QGTTLPKG**F**YAE**G**SRGG**S**QASS**R**SS--RSRNSSRN**N**STPGSSRG-TSPARMA-----GNGGDAALALLLDRLNQLES**K**-MSG**K**G--Q-Q  
SARS-CoV 164 QGTTLPKG**F**YAE**G**SRGG**S**QASS**R**SS--RSRGNSRN**N**STPGSSRG-NSPARMA-----SGGETALALLLDRLNQLES**K**-VSG**K**G--Q-Q  
HKU1 177 PGT**I**LP**Q**GY**V**EG**S**GR**S**ASN-SRP-----GSRSQS**R**GPNNR**S**LS**R**S**N**SNFR**R**HS-----DSIV**K**P**D**MADE**I**ANL-VLA**K**LG**K**D-S  
OC43 178 PGT**V**LP**Q**GY**V**IE**G**SG**R**SAPN-SRS-----TSRT**S**RASSAG**S**RS**R**AN**S**GN**R**TP-----TSG**V**TP**D**MAD**Q**I**A**SL-VLA**K**LG**K**D-A  
MERS-CoV 152 PGT**K**LP**K**N**F**H**I**E**G**GG**N**S**Q**SS**R**ASS--LSRNSSR**S**SS**S**Q**G**SR**S**G-NSTRGTS--PGPS**G**IGAV**G**DD**L**Y**L**LD**L**LN**R**Q**A**LF-ES**G**K**V**--K-Q  
229E 130 -NQ**K**LP**N**G**V**T**V**A**E**EP**D**SR-----APS**R**S**Q**SR**S**Q**R**SR**V**ES**K**QS-RNPSSDRNH**N**SQ**D**IM**K**A**V**A**A**ALK**S**LG**F**D**K**P**Q**E**K**D**K**SA**K**T**G**TP**K**P  
NL63 128 -SIAL**P**PELS**V**VE**F**ED**R**SN**N**NS**R**ASS**R**S**S**TR**N**NN**S**RD**S**SR**S**TS**R**Q-Q**S**RT**R**SD**S**N**Q**SS**D**LVA**A**VT**L**KN**I**LG**F**D**N**Q**S**K**P**SSSS-GT**S**TP

280 290 300 310 320 330 340 350 360

SARS-CoV-2 241 Q**O**Q**G**OT-V**T**K-----K**S**AA-E**A**SK**K**P**R**Q**K**R**T**T**A**K-Y**N**V**T**Q**A**F**G**R**R**P**E**Q**T**Q**G**N**F**G**D**Q**E**L**I**R**Q**G**T**D**Y**K**H**W**P**Q**I**A**Q**F**A**P**S**A**S**  
SARS-CoV 242 Q**O**Q**G**OT-V**T**K-----K**S**AA-E**A**SK**K**P**R**Q**K**R**T**T**A**K-Y**N**V**T**Q**A**F**G**R**R**P**E**Q**T**Q**G**N**F**G**D**Q**D**L**I**R**Q**G**T**D**Y**K**H**W**P**Q**I**A**Q**F**A**P**S**A**S**  
HKU1 248 -K**P**Q**Q**-V**T**K-----Q**N**AK**E**IR**H**K**I**L**T**K**P**Q**K**R**T**P**N**K-H-C**N**V**Q**C**F**G**K**R**G**P**S**Q--N**F**G**N**A**E**M**L**K**L**G**T**D**P**Q**F**P**I**L**A**E**L**A**P**T**P**  
OC43 249 T**K**P**Q**Q-V**T**K-----H**T**A**K**E**V**R**Q**K**I**L**N**K**P**Q**K**R**T**P**N**K-Q-C**T**V**Q**C**F**G**K**R**G**P**N**Q--N**F**G**G**E**M**L**K**L**G**T**S**D**P**Q**F**P**I**L**A**E**L**A**P**T**A**  
MERS-CoV 233 S**Q**P**K**V-I**T**K-----K**D**AA-A**A**AK**N**K**M**R**H**K**R**T**S**T**K**S-F**N**M**V**Q**A**F**G**L**R**G**P**G**D**L**Q**G**N**F**G**D**I**L**Q**L**N**K**L**G**T**E**D**P**R**W**P**Q**I**A**E**L**A**P**T**  
229E 214 S**R**N**O**SP**A**SS**Q**S**A**AK**I**L**A**R**S**Q**S**SET**K**EQ**Q**K**H**E**M**Q**K**P**R**W**K**R**Q**P**N**D**V**T**S**N**V**T**O**C**F**G**P**R**L**D**H**-N**F**G**S**A**G**V**V**A**N**C**V**K**A**G**Y**P**Q**F**A**E**L**V**P**  
NL63 214 K**K**PN**K**P**L**-----S**Q**P**R**AD**K**P**SQL**K**K**P**R**W**K**V**P**T**R**E--E**N**V**I**Q**C**FG**P**R**D**F**N**H--N**M**G**D**S**D**L**V**Q**N**C**V**DA**K**G**F**P**Q**L**A**E**L**I**P**N**Q**A

370 380 390 400 410 420 430 440 450

SARS-CoV-2 314 F**F**G**M**S**R**I**G**M**E**V**T**P-----SGT**W**L**T**Y**T**G**A**I**K**L**D**D**K**D**P**F**N**K**D**Q**V**I**I**L**N**K**H**I**D**A**Y**K**T**F**P**P**T**E**P**K**D**KK**K**K**A**D-E**T**Q**A**L**P**Q**R**Q-----  
SARS-CoV 315 F**F**G**M**S**R**I**G**M**E**V**T**P-----SGT**W**L**T**Y**H**G**A**I**K**L**D**D**K**D**P**Q**F**K**D**N**V**I*I*L**N**K**H**I**D**A**Y**K**T**F**P**P**T**E**P**K**D**KK**K**K**T**D-E**A**Q**P**L**P**Q**R**Q-----  
HKU1 321 F**FFF**GS**K**L**D**L**V**K**R**D--SEAD**S**P**V**K**D**V**F**E**L**H**Y**S**G**S**I**R**F**D**S**T**L**P**F**E**T**I**M**K**V****L**E**E**N**L**N**A****Y**V**N**S**Q**-----N**T**D**S**D**S****L****S****S****K**P**Q**R**K**R**V**K**Q**  
OC43 323 F**FFF**GS**R**L**E**L**A**K**V**Q**N**L**S**GN**P**D**E**P**Q**K**D**V**Y**E**L**R**Y**N**G**A**I**R**F**D**S**T**L**S**G**F**E**T**I**M**K**V**N**E**N**L**N**A**Y**QQ**Q**D**G**-----M**M**N**M****S**P**K**P**Q**R**Q**R**G**H**K**  
MERS-CoV 306 F**M**G**M**S**Q**F**K**L**T**H**Q**N-NDD**H**G**N**P**V**F**L**R**Y**S**G**A**I**K**L**D**P**K**N**P**N**K**W**L**E**Q**N**I**D**A**Y**K**T**F**P**K**E**K**K**Q**K**A**P**K**E****S**T**D****Q****M****S****E****P****P****K**E**Q**R**V****G**  
229E 302 M**L**F**D**SH**I**V**S**K**E**G-----N**T**V**V**L**T**F**T**T**R**V**T**P**K**D**H**E**H**L**G**K**F**L**E**E-L**N**A**F**T**R**E**M**QQ**Q**P**L**N**N**P-----S**A**L**E**F**N**-----  
NL63 287 I**F**F**D**SE**V**S**T**D**E**V**G**-----D**N**V**Q**I**T**Y**T**K**M**V**A**K**D**N**K**N**L**P**K**I**E**Q-I**S**A**F**T**K**P**S****S**I**K**E**M**Q**S**Q-S**S**H**V**A**Q**-----

460 470 480 490 500

SARS-CoV-2 387 -----K**K**Q**Q**T-----V**T**L**L**P**A**AD**L**DD**F**S**K**Q**L**Q**Q**S**M**S**S**A**D****S**T**Q**A-----  
SARS-CoV 388 -----K**K**Q**P**T-----V**T**L**L**P**A**AD**M**DD**F**S**R**Q**L**Q**N**S**M**S**G**A**S**A**D****S**T**Q**A-----  
HKU1 400 L**P**EQ**F**D**S**L**N**L**S**A-G**T**Q**H**I**S**N**D**F**T**P**E**D**H**S**L**L**A**T**L**DD**P**V**E**D**S**V**A**-  
OC43 402 G**Q**G**E**D**N**I**S**V**A**V**P**K**S**R**V**Q**Q**N**K**S**R**E**L**T**A**E**D****I**S**L**KK**M**D**E**P**T**E**D****T**  
MERS-CoV 392 I**T**Q**R**T**R**T**R**P**S**-----V**Q**P**G**P**M**I**D**V**N**T-----  
229E 363 -----P-S**Q**T**S**P-----A**T**V**E**P**V**R**D**E**V**S**I**E**T**D**I**I**D****E**V**N**-----  
NL63 348 -----N**T**V**L**N**A**I-----P**E**S**K**P**L**A**D**D**S**A**I**I**E**I**V**N**E**V**L**H-----

10 20 30 40 50 60 70 80 90 100 110 120

*MHV* MSFVPGQENAGGRSSSVN RAGNGILKTTW-----ADQT-----ERGPNNQNRGR-RNQPKQTATT-QPNNSGVVPHYSWFSGITQFQKGKEFQFAEGQGVPIAPGVATEAKGYWYRHNR  
*HKU14* MSFTPQKQP-SSRASSGNRAGNGILK---W-----ADQS-----EQSRNIQTRGR-RAQPKQTQVTSQQPTGGNVVPPYSWFSGITQFQKGKEFEAEGQGVPIAPGVATEAKGYWYRHRSRR  
*GCCDC1* MSGR-----NT-PKGRQ-----TP-KVSFKQESDSDSE-ADRHTTRNGARPKT--NNGP-ATIKPEK-----PRAAPMQNVSWFAPLVQTGK-SDLKFAHGEQGPVQGDGTYQHGYWLRTQ-R  
*Hedgehog-CoV* MAT-----PQ-----QPRAVTF-----ADNN-----GNQQN-----GNN-RGRPRQPK-----PRPAPNVSWSWYTGITQHGK-QPLAFPAQGVPLNANSTPKQNAGYWRROD-R  
*Bat-BCoV* MAA-----E-----GQ-RVTFLTQGSQNQNN-----NNQEN-----QDGR-GARPKTRK-----PRPAPQNNVSWFTPLTQHGK-QALTFNPQGVPLNANDDPSRIGYWRRI-R  
*Rat-CoV* MSFVPGQENAGSRSSSGNRAGNGILKTTW-----ADQT-----ERGQNNQNRGR-RNQPKQTATT-QPNTGSVVPHYSWFSGITQFQKGKEFQFAGQQGVPIANGIPSSQEKGWYRHNR  
*HKU24* MSHTPGKAG-S-RSSSGNR-NGGILK---W-----ADQT-----EP-NNVNSRGR-RAQPKQTANT-QASGGNVVPPYSWFSGITQFQKGQEFQFEGQGVPLTNGVPPSEEKGWYRHNR  
*SARS-CoV* MSDN-----GPQSNQRS-----AP-RITF-GGPTDST-----DNNQN-----GGRN-GARPKQRR-----PQGLPNNTASWFTALTQHGK-EELRFPRQGVPIINTNSGPDDQIGYRRAT-R  
*HKU9* MSGR-----NRSRSGTP-----SP-KVTFKQESDGSDSEERRNGNRNGARPKN-----NNSR-GSAPKPEK-----PKAAPPQNVSWFAPLVQTGK-AELRFPRGEQGPVQGDSTYEHGYWLRTQ-R  
*HKU5* MAT-----PA-----PPRAVVF-----ADNN-----ETPTN-----SQR-SGR-PRTK-----PRPAPNTTVSWFTGLTQHGK-QPLAFPPQGVPLNANSTPAQNAGYWRROD-R  
*HKU4* MAT-----PA-----APRTISF-----ADNN-----DNQPN-----QQO-RGRGRNP-----PRPAPNNTVSWYTGLTQHGK-NPLAFPPQGVPLNANSTTAQNAGYWRROD-R  
*Bovine-CoV* MSFTPQKQS-SSRASSGNRSGNGILK---W-----ADQS-----DQSRNVQTRGR-RAQPKQTATSQQPSGGNVVPPYSWFSGITQFQKGKEFEAEGQGVPIAPGVATEAKGYWYRHNR  
*Covid-19* MSDN-----GPQ-NQRN-----AP-RITF-GGPTDST-----GSNQN-----GERS-GARSKQRR-----PQGLPNNTASWFTALTQHGK-EDLKFPQGVPIINTNSPDDQIGYRRAT-R  
*HKU1* MSYTPGHYA-GSRSSSGNR-SGILKKT SW-----ADQS-----ERNYQTFNRGR-KTOPKFTVSTQ-----PQGNTIPHYSWFSGITQFQKGRDFKFSDQGVPIAFGVPPSEAKGYWYRHRSRR  
*MERS-CoV* MAS-----PA-----APRAVSF-----ADNN-----DITNT-----NLS-RGRGRNP-----PRAAPNNTVSWYTGLTQHGK-VPLTFFQGVPLNANSTPAQNAGYWRROD-R  
*OC43* MSFTPQKOS-SSRASSGNRSGNGILK---W-----ADQS-----DOFRNVOTRGR-RAOPKOTATSOOPSGGNVVPYYSWFSGITOFQKGKEFEFVEGOGVPIAPGVATEAKGYWYRHNR  
*TGE* MAN-----QGQRVSW-----GDES-----T-K-----TRGRSNSRGRKN-----NNIPLSFFNPITLQQGSKFWNLCPDFVPKGIGN-RDQQIGYWNRQ-T  
*FCoV* MAT-----QGQRVN -----GDEP-----S-K-----RRGRSNSRGRKN-----NDIPLSFYNPITLEQGSKFWNLCPDLVPKGIGN-KDQQIGYWNRQ-I  
*NL63-related* -----MATVN -----AD-----D-KRGK-----R-----KYPPPSFYLPPLVVKSDQPYKVI PRNLVPKGKGN-KDQQIGYWNVQ-R  
*HKU2* -----MATVN -----GDAV-----EQAE-----SRG-----R-----KRIPLSLFAPLRVTDGKNFNWNVMPRNGVPTGRGN-PDQQIGYWEQK  
*BtCoV512* -----MASVKF-----QP-----RGR-----SK-----GRVPLSLFAPLRVTDEKPLYKVLPPNAVPQGMGG-KDQQIGYWEQ-Q  
*Porcine-EDV* -----MASVSF-----QD-----RGR-----KRVPLSLYAPLRVTNDKPLSKVLANNAVPTNKGN-KDQQIGYWNNEQ-I  
*BtNv-AlphaCoV* MSS-----SKGNVGF-----DN-----A-ARGR-----S-----GRVPFSFYMPVINNSSQPFYKVMQNAVPKGQGN-KDQQIGYWNNEQ-V  
*BtMr-AlphaCoV* -----MSSVKF-----EA-----SGR-----T-----GRTPLSFFAPLTVSSDKFVNVLPKNAVPTGKGN-AKQQVGYWNNEQ-P  
*Bat-CoV-1A* -----MASVQF-----EDT-----N-SRGR-----S-----GRIPLSYYAPLRVADDKFLKSMPGNGVPKGGMG-KAEQIGYWT EQR  
*Mink-CoV* MAG-----NGNRVN -----GDEP-----SPSQ-----NRGRSRSRSRKN-----ENIPLSFFNPITHGNKKFWNVAPKDLVPKGKGN-RDQQVGYWNRQ-E  
*LuchengRn-CoV* -----MSSNVSW-----ADQV-----ETVN-----RRQRSRSRGRSO-----NRTNASNPLSWFTSIIDESNGFISLMPHSGVPTGMGT-AAQQCGYWYRAP-T  
*BtRf-AlphaCoV* -----MASVSF-----QE-----E-KRGR-----S-----GKVPLSYYNPIVSGDKFWKVMPPNAVPKGKGN-KDQQIGYWNNEQ-P  
*HKU10* MAS-----GGNNVSF-----AN-----K-PRGR-----S-----GKVPLSYYSPVMIDGDQFWKVMPTNAVPTGMGD-KNQQIGYWNNEQ-P  
*CDPHE15* -----MASVKF-----AK-----NNR-----R-----GRMEYSYFAPLIINSDQPIWKVLPPNAVPVGKGG-KDEQIGYWNNEQ-P  
*Ferret-CoV* MAG-----NGQRVN -----GDEP-----VPSQ-----KRGRSRSRSRRN-----ADIPLSYFNPITHEGKKSFWTVAPKDFVPIKGKGN-KDQQVGYWNRQ-Q  
*HKU8* -----MASVSF-----SD-----E-PRGR-----S-----GRVPLSLFAPLRTTDGSLFFRAMPQNGVPKGFGN-KDQQVGYWNNEQ-V  
*NL63* -----MASVNW-----AD-----D-RAAR-----K-----KFPPPSFYMPLLVSSDKAPYRVI PRNLVPIGKGN-KDEQIGYWNVQ-E  
*229E* -----MATVKW-----ADAS-----E-P-QRGR-----O-----GRIPYSYSPLLVDSEQ-PWKVIPRNLLVPVNKKD-KNKLIGYWNVOK--

