logo

Name		Value
SARS-CoV-1	....ATATTAGGTTTTTACCTACC.CA....GGAAA..AGCCAACCAACCT.CGATC..TCTTGT.AGATCTGTTC....TCTAAAC.GAACTTTAAAATCTGT.GTAGCTGTCGCTCGGC.............	100
SARS-CoV-2	....ATTAAAGGTTTATACCTTCC.CA....GGTACAAACCAACCAACTTCGATC..TCTTGT.AGATCTGTTC....TCTAAAC.GAACTTTAAAATCTGT.GTGGCTGTCACTC.....	100
Hp-bCoV-2013	TTAAGCTTCGGCTTGTTGCATAGGACC....GGAAAGGTACTATCTACCCT...AAC..TCTTGT.AGTTAGACTC...TCTAAAC.GAACTTTAAAACTGGTTGTGTCCTTCA.....	100
A59	....GTATAAGAGTGATTGGCGTC....CGTACGTACCGTCTCAACTCTAAAAC..TCTTGT.AGTTTAAATC.TAATCTAA.....ACTTTATAAACGGC..ACTTCCTGCGTGTCCA.....	100
MHV-A59-C12	....TATAAGAGTGATTGGCGTC....CGTACGTACCGTCTCAACTCTAAAAC..TCTTGT.AGTTTAAATC.TAATCTAA.....ACTTTATAAACGGC..ACTTCCTGCGTGTCCAT.....	100
MHV-JHM	....TATAAGAGTGATTGGCGTC....CGTACGTACCGTCTCTACTCTAAAAC..TCTTGT.AGTTTAAATC.TAATCTAATCTAAACTTTATAAACGGC..ACTTCCTGCGTGTG.....	100
Rat-CoV-Parker	....................CGCTCTCAACTCTCC..TCTTGT.AGTTTAAATC.TAATCTAA.....ACTTTACAAACGGC..ACTTCCTGCGTGTCCATGCCCGTGGGCTTGGTCTTGTCATAGTG	100
OC43-ATCC-VR-759	....ATTGTGAGCGATTGCGTG....CGTGC.ATCCGCT..TCACTGATC..TCTTGTAGATCTTTGTAATCTAA.....ACTTTATAAAACATCCACTCCTG.TAATCTAT.....	100
Bovine-CoV	....GATTGCGAGCGATTGCGTG....CGTGC.ATCCGCT..TCACTGATC..TCTTGTAGATCTTTGTAATCTAA.....ACTTTATAAAACATCCACTCCTG.TAGTCTA.....	100
HKU14	....GATTCTGAGCGATTGCGTG....CGTGC.ATCC.GCC..TCAGTGAAC..TCTTGTAGTTCAATT.TAATCTAA.....ACTTTATAAAACACCTACTCCTG.TAATCTATG.....	100
HKU1	....GAGTTTGACCGATTGACGTT....CGTAC..CGTCTATGCAGCTTACGATC..TCTTGTCAGATCTCATT.AAATCTAA.....ACTTTTTAAACAAG...ATTCCTG.TTATCCATG.....	100
HKU24-R05005I	....GATAAAGTGTGAATCGCTTC....CGTAG..CATC.GCA..CCCTCGATC..TCTTGTTAGATCTAATC.TAATCTAA.....ACTTTATAAAACACTAGGTCCTGCTAGCCTAT.....	100
Beta-CoV-England-1	......ATTTAAGTGAAATAGCTTG....GCTATCTCACTTCCC....CTCGTC..TCTTGCAGAACTTTGAT....TTTAACG.AACTTAAATAAAGCCCTGTTGTTTAGCGTATTGTT.....	100
MERS-CoV	....GATTTAAGTGAAATAGCTTG....GCTATCCC..TCTCT....CTCGTC..TCTTGCAGAACTTTGTT....TTTAACG.AACTTAAATAATAGCGCCTGCTGGTTTGCGTGCTGCGA.....	100
HKU5-1	....GATTTAAGCAGAAATAGCCTG....GCTATCCC..TCTCT....CTCGTC..TCTTGCAGAACTTTGT.....TTTAACG.AACTTAAATAATAGCGCCTGCTGGTTTGCGTGCTGCGA.....	100
HKU4-1	....GATTTAACAGAAATAGCATA....GCTATCCCTCTCTCT....C.CGTTC..TCTTGTAGAACTCTTGT....TTTAACG.AACTTAAATAAAGCCCTTATACTTTGTCTAGGTAG.....	100
ErinaceusCoV-2012	....GAAATAAAAGTAATTGCGTGCG.TGCAATCAACTTTCC.CCCTCGATCG.TCTTGTACGATTCACTC....TTGAACG.AACTTAAATTGTGCATATTGTGTCAC.....	100
GCCDC1	....GATAAAAGTGATTAGCCGCG.TGCGTATCGCCGATCCTAAC.CCCGATCG.TCTTGTACGATTCACTC....TTGAACG.AACTTAAATTGTGCATATTGTGTCAC.....	100
HKU9-1	...TTTAAAATCTGTGTAGCCGTGTCACTTGGCTGCATGCCCAGTGCCACTTACGGCGTTATATCTTATAAACTTTTACTGTCGTGTGACAG..GACACGAGTAACTCGT.....	100
BM48-31	(see alignment)	100
consensus		100

X non conserved
X ≥50% conserved