

Virus Report

(Comparisons)

25-01-2015 14:55:49

Sequences:

HI-V:

TGGTATCAGAGCAAGGTTTTCTGTAAGTAATTTATGGCTTTCATGGGGTAAACCCTTAGATAGGAGCCTGAGGGCTCTGCTATGTTCTATTTTGAGAAAATAAC
TGTGTAAGTAGTAATGCATGATAAATCGAATAAGTTCTTTACGGATGGTCCTAAATTTATGAAATCATTCTTTTACTATCTGAGTGAGACCCTGTGTTTCCAAGT
TCTACTGAGAAATCATGAGTAAATGGACGTTGACACACTGACTTAAGGAAAAGAAAGATGTGCCTAGGATGAAGTCGAGCACTGTTCTAGGCTGAAGGGTGAA
AAGAAAAATGAGTTGAGTTGGGAGTAAGACTCTGAGAGGCTTGAGGCAAAAGAACCGGAACCTTACCTAGCAAAGTAAGACCCAAGGGAGGGCAGTAGCTGG
AGAACCCCGGTGTGAGTTTCTTTACCTAGCAGAAGATATACTCTAGTATCCTCAAAACACCGTGAAGTATTTACAACCCCTTTATGAACAAGTATTTTCTGGAGTT
AGATGTCAGCCACCAAAGATCATGATCTATCAGAAACAACTTGAGTCATACAGAGAAGAACTCAGATGTAACCTGAATGACTAATGATCAGGGACCTCTATTAA
AGAGCTTAGTAAGAAGTAAGTGCATAAACTACCTCTGGAACCTGTTATAATCTTGAAATGAGCATATGTATCCCAAAGGTTACTCCTATCTGTCCGTGACCATAA
GAAACCTTCTTTAAAGAAAACTCTGGATTCTACTGCGATCTTGAATTTGAAAAGTACAAAACTCTCATTCCAAATCTGTTTCAGATCTTTCTTATTTAGACTTA
GCATCTGCTGATAAGGTCTCTAATAAAGACCTTGCAATTTAATTTACATATAAACACTTATCGTAGTGATCTTGGTTTTAAAGTCGCAATTCACCTCTTTATCAAAAAA
TCGTGAGCTGCTAATACAGAACAGGAAGCTTTTGAAGAACAAGGAGCAGCTGTCTGAAATTAACAACCTATCTAAGGTTGTGCGTCTCCAACGAGCTGATT
TAAAGGAGACCTTGAGAAGACAGGACGCTCCTTGCGAAGGAGCTTCAGGCTCTTCGTAAGACTATCTGGAAAGGCGCCCTCTAGCAAGGAGGACGTGGAAG
AATTGGTTGTCCGCATCTCTGAACAACCAAAATTTATTGAAAAACAGACGGAAGCTCTTACTGAGGAGCTTACAAAGGAGGTTTCAGGCACTGAAGCTAATCA
TCCATAACTTCGAACAGAAGTTACTGGGATGAGCCTGGCCGGAAGCAAGGCCACAGCAATCTATCAAGAAGCCTTGCAATCTACTGCTGCTGGTTGGGAAGA
CACCGGCATCGGATTTACTGATAAAGGAGTTATCAGTACCACTACCCTATCAAGGCAGCTAAATTCGCTGTTATTTCTGGTGATCAAAACAAATCAGCAGATCA
CCACCATCTCGGAAGAACTAACCAACTTCACAACCGGGTAAAAACCTTGAAAGGAAGGACAGGGGTATCTGCAAGCCCCCTGTATAAATCGGAGATTGAATC
CATCAACGCCAAGCTCAAAAGTATTCAAGATATACAAGGGAGTCACCCCTCCAAAAGAGACCCCTCCGGAGTTATCAAAAGTTTTTGAGGATCCCTACAGCATCC
TCCGCAGACTATAAATGGCGTCAAGACCTAGAGTTTCTGGGAGCACTACCAGAACCATGATAGCAGAACCAGGGGTTCCCTGGTGGATGATCAGATCCGGG
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CAGGGAGGCTGGATTTGAGTACATCCATCTAGGGGTACTCCAGGTGCGCATCCAAATATGCAACCGAACATATGATGGTACAATGGCATTAGTGGTCTTCCGT
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TGATGGGGATTCCGAAGTTAGCACCCCAAGGAGCAATACATATATTGCGCTAGAAGATGAGTACCCAAAACCTTCAACAGCTGGAAGGCCTGGTACTCTCC
ACAACAGAGTCTGCTATCAGTCGCTTTAGACCAGCAGACAGATATGACTGGCGTAGGCCCTGGCTACGCACCAGCAACTGGAACAGCTGGCTATACTGGA
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TCGTTATGCCTCAGGATTTTGACACAAGGTCTTTGAA

Test:

TGGTATCAGAGCAAGGTTTTCTGTAAGTAATTTATGGCTTTCATGGGGTAAACCCTTAGATAGGAGCCTGAGGGCTCTGCTATGTTCTATTTTGAGAAAATAAC
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GCCAGTTCATCTGATTTCCTTACCCTAGAAGACCAAGAAAAGTGGGACAACAACCTCGGAGTGTTCAACCTACCCACCGCTAACGCGAGGCAAGCATCAATAT
TCGTTATGCCTCAGGATTTTGACACAAAGGCTCTTTGAA

Secondary Structure:

HI-V:

[illegible]

Test:

[illegible]

Percentual similarity of domains: 0%