Virus Report

(Comparisons)

26-01-2015 16:01:54

Segunences:

aowiuhf

TGGTATCAGAGCAAGGTTTTCTGTAAGTAATTTATGGCTTTCATGGGGTAAAACCCTTAGATAGGAGCCTGAGGGCTCTGCTATGTTCTATTTTGAGAAAATAAC TCTACTGAGAAATCATGAGTAAATGGACGTTGACACACTGACTTAAGGAAAAGAAGATGTGCCTAGGATGAAGTCGAGCACTGTTCTAGGCTGAAGGGTGAA AGAACCCCGGTGTGAGTTTCTTTACCTAGCAGAAGATATACTCTAGTATCCTCAAAACACCGTGAAGTATTTACAACCCTTTATGAACAAGTATTTTCTGGAGTT AGAGCTTAGTAAGAAGTAAGTGCATAAACTACCTCTGGAACTGGTTATAATCTTGAAATGAGCATATGTATCCCAAAGGTTACTCCTATCTGTCCGTGACCATAA GCATCTGCTGATAAGGTCTCTAATAAAGACCTTGCATTTAATTTACATATAAACACTTATCGTAGTGATCTTGGTTTTAAAGTCGCAATTCACTCTTTATCAAAAAA TCGTGAGCTGCTAATACAGAACAGGAAGCTTTTGGAAGAACAAAAGCAGCAGCTGTCTGAAATTAACAACCTATCTAAGGTTGTGCGTCTCCAACGAGCTGATT TAAAGGAGACCTTGAGAAGGACGGCCCTTGCGAAGGAGCTTCAGGCTCTTCGTAAAGACTATCTGGAAAGGCCCCCTCTTAGCAAGGACGACGTGGAAG AATTGGTTGTCCGCATCTCTGAACAACCAAAAATTTATTGAAAAAACAGACGGAAGCTCTTACTGAGGAGCTTACAAAGGAGGTTCAGGCACTGAAGCTAATCA TCCATAACTTCGAACAGAAGTTACTGGGATGAGCCTGGCCGGAAGCAAGGCCACAGCAATCTATCAAGAAGCCTTGCAATCTACTGCTGCTGGTTGGGAAGA CATCAACGCCAAGCTCAAAAGTATTCAAGATATACAAGGGAGTCACCCTCCAAAAGAGACCCCCTCCGGAGTTATCAAAGTTTTTTGAGGATCCCTACAGCATCC TCCGCAGACTATAAATGGCGTCAAGACCTAGAGTTTCTGGGAGCACTACCAGAACCATGATAGCAGAACCAGGGGTTCCCCTGGTGGATGATCAGATCCGGG AATACCGGAGTGCGGCCCGAGTCGCATACGAAGCACAAAGGATTGCCAGACGCACTGGCAATATTCTTGGAAGAATTGTTGGACGACAACCAAGGGAGCATA CCCTTGCGATGGTCGTGGACCCCAACAGTGAGCTGGAACGCTCACTGGCTCACCGGGCACGAACAATACCTGCTGAAGTCCTTTACATGACCCAGCGGGGTG AGCCTACAAATCGGGTCTATCGGAACAGAACTGAAGAAGGGTCCTAGTGACTCATGGTCAGCAAGATAGGACTTTCATCTTGCCAGAGTCATATGAAGAGCT CAGGGAGGCTGGATTTGAGTACATCCATCTAGGGGTACTCCAGGTGCGCATCCAAATTATGCACCGAACATATGATGGTACAATGGCATTAGTGGTCTTCCGT GACACCAGATGGACACAGGAGAACCACCAAGACAGATCCATTATAGCAACCATGGAGGCTGATCTCTCCCAAGGGCATCAGCTAATTTACGTCATCCCAGGATA TAATGATGACTATTCGGGATTTCTACCAGCACATTCAGATAAGCATCCTTACCAAAGGATATGAAGGCTTTCAAGGAGAAGCAAATCTCCTTATTACAAGAAGCT GCCGGTGCAGATTAAGCAACGTTCCTAACGTTGGCTTCCAATACAACATCCAGAACGTCGTGGAATTCTTAAAATCTAGGGGAGTAAAAGCCCTAAATGCGAC GAAATTAAGCACCAGGAGGTTCCAAGGAGGAGAATGGAACATCCGACCATCAGAAGTGGTGGTCCCTATGCAGCCAACAACTATGATAGTACGAGTTAATTAT GACTCGTCACGAAGCATCAGATTCGGGGGATTATGAAGCTAGCACATCATCATCAGCCCCGAGATATGAGCAAGACGGTGATGAAGATGAAGCACTTGGTGATA TCCATCAAGTAAATATGATCACCATCATAGAAGATGATGCAGAAGATGACTACCCACGGTTATCAGCTTTAGAGCGAATAATCGCTCCAGAAAGCATGGTGGGA GAGGAGGACACAATTGCTGAATTTTTAAGCAATCTGTCCTTAGATTCCTCCACTGATGAGGAATTCTACGACGCCGACAACTCATTATTCGAAGAAGAAGAATA TGATGGGGATTCCGAAGTTAGCACCCCAAGGAGCAAATACAATATATTTGCGCTAGAAGATGAGTACCCAAAACTTCAACAGCTGGAAAGCCTGGTACTCTCC ACAACAGAGTCTGCTATCAGTCGCTTTAGACCAGCAGACACAGATATGACTGGCGTAGGCCCTGGCTACGCACCAGCAACTGGAACAGCTGGCTATACTGGA GCCAGTTCATCTGATTTCCCTTACCCTAGAAGACCAAGAAGACGGGACAACAACTCGGAGTGGTTCAACCTACCCACCGCTAACGCGAGGCAAGCATCAATAT TCGTTATGCCTCAGGATTTTGACACAAAGGTCTTTGAA

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TGGTATCAGAGCAAGGTTTTCTGTAAGTAATTTATGGCTTTCATGGGGTAAAACCCTTAGATAGGAGCCTGAGGGCTCTGCTATGTTCTATTTTGAGAAAATAAC TCTACTGAGAAATCATGAGTAAATGGACGTTGACACACTGACTTAAGGAAAAGAAGATGTGCCTAGGATGAAGTCGAGCACTGTTCTAGGCTGAAGGGTGAA AGAACCCCGGTGTGAGTTTCTTTACCTAGCAGAAGATATACTCTAGTATCCTCAAAACACCCGTGAAGTATTTACAACCCTTTATGAACAAGTATTTTCTGGAGTT AGATGTCAGCCACCAAAGATCATGATCTATCACAAACCAAACCTTGAGTCATACAGAGAAGAACTCAGGTGTAACTGAATGACTAATGATCAGGGACCTCTATTAA AGAGCTTAGTAAGAAGTAAGTGCATAAACTACCTCTGGAACTGGTTATAATCTTGAAATGAGCATATGTATCCCAAAGGTTACTCCTATCTGTCCGTGACCATAA GCATCTGCTGATAAGGTCTCTAATAAAGACCTTGCATTTAATTTACATATAAACACTTATCGTAGTGATCTTGGTTTTAAAGTCGCAATTCACTCTTTATCAAAAAA TCGTGAGCTGCTAATACAGAACAGGAAGCTTTTGGAAGAACAAAGCAGCAGCTGTCTGAAATTAACAACCTATCTAAGGTTGTGCGTCTCCAACGAGCTGATT TAAAGGAGACCTTGAGAAGGACGTCCTTGCGAAGGAGCTCTCAGGCTCTTCGTAAAGACTATCTGGAAAGGCCCCCTCTTAGCAAGGACGACGTGGAAG AATTGGTTGTCCGCATCTCTGAACAACCAAAAATTTATTGAAAAAACAGACGGAAGCTCTTACTGAGGAGGCTTACAAAGGAGGTTCAGGCCACTGAAGCTAATCA TCCATAACTTCGAACAGAAGTTACTGGGATGAGCCTGGCCGGAAGCAAGGCCACAGCAATCTATCAAGAAGCCTTGCAATCTACTGCTGCTGGTTGGGAAGA CATCAACGCCAAGCTCAAAAGTATTCAAGATATACAAGGGAGTCACCCTCCAAAAGGAGACCCCTCCGGAGTTATCAAAGTTTTTGAGGATCCCTACAGCATCC TCCGCAGACTATAAATGGCGTCAAGACCTAGAGTTTCTGGGAGCACTACCAGAACCATGATAGCAGAACCAGGGGTTCCCCTGGTGGATGATCAGATCCGGG AATACCGGAGTGCGGCCCGAGTCGCATACGAAGCACAAAGGATTGCCAGACGCACTGGCAATATTCTTGGAAGAATTGTTGGACGACAACCAAGGGAGCATA CCCTTGCGATGGTCGTGGACCCCAACAGTGAGCTGGACGCTCACTGGCTCACCGGGCACGAACAATACCTGCTGAAGTCCTTTACATGACCCAGCGGGGTG AGCCTACAAATCGGGTCTATCGGAACAGAACTGAAGAAAGGATGCTAGTGACTCATGGTCAGCAAGATAGGACTTTCATCTTGCCAGAGTCATATGAAGAGCT CAGGGAGGCTGGATTTGAGTACATCCATCTAGGGGTACTCCAGGTGCGCATCCAAATTATGCACCGAACATATGATGGTACAATGGCATTAGTGGTCTTCCGT GACACCAGATGGACACAGGAGAACCACCAAGACAGATCCATTATAGCAACCATGGAGGCTGATCTCTCCCAAGGGCATCAGCTAATTTACGTCATCCCAGGATA

TAATGATGACTATTCGGGATTTCTACCAGCACATTCAGATAAGCATCCTTACCAAAGGATATGAAGGCTTTCAAGGAGAAGCAAATCTCCTTATTACAAGAAGCT
GCCGGTGCAGATTAAGCAACGTTCCTAACGTTGGCTTCCAATACAACATCCAGAACGTCGTGGAATTCTTAAAATCTAGGGGAGTAAAAGCCCTAAATGCGAC
GAAATTAAGCACCAGGAGGTTCCAAGGAGGAGAACATCCGACCATCAGAAGTGGTGGTCCCTATGCAGCCAACAACTATGATAGTACGAGTTAATTAT
GACTCGTCACGAAGCATCAGATTCGGGGATTATGAAGCTAGCACCATCATCATCAGCCCCGAGATATGAGCAAGACGGTGATGAAGATGAAGCACTTGGTGATA
TCCATCAAGTAAATATGATCACCATCATAGAAGATGATGCAGAAGATGACTACCCACGGTTATCAGCTTTAGAGCGAATAATCGCTCCAGAAAGCATTGGTGGGA
GAGGAGGACACAATTGCTGAATTTTTAAGCAATCTGTCCTTAGATTCCTCCACTGATGAGGAATTCTACCGACGCCGACAACTCATTATTCGAAGAAGAAGAATA
TGATGGGGATTCCGAAGTTAGCACCCCAAGGAGCAAATACAATATATTTTGCGCTAGAAGATGAGTACCCAAAACTTCAACAGCTGGAAAGCCTGGTACTCTCC
ACAACAGAGTCTGCTATCAGTCGCTTTAGACCAGCAGAAACACAACAACACTCGGAGTGGTTCAACCACCACCACCAACACTGGAACACCAGCAACCTCAATAT
TCGTTATGCCTCAGGATTTTGACACAAAGGTCTTTGAA
TCGTTATGCCTCAGGATTTTGACACAAAGGTCTTTGAA

Secondary Structure:

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Energies of secondary structure: aowiuhf: -2272.4

aowiuhf: -2272.4

GC content: aowiuhf: 44.0923076923 aowiuhf: 44.0923076923

Number of Proteins: aowiuhf: 3

aowiuhf: 3

Percentual similarity of domains: 100%

Common domains	Position in aowiuhf	Position in aowiuhf	Length in aowiuhf	Length in aowiuhf	ĺ
DUF1319	768:898	768:898	130	130	ĺ
zf-CCHC	2444:2461	2444:2461	17	17	ĺ
RVP	2771:2882	2771:2882	111	111	ĺ
RVT_1	3030:3190	3030:3190	160	160	ĺ
MP	1806:1927	1806:1927	121	121	ĺ