Virus Report

18-01-2015 14:49:59

Secondary structure results

Virus name: HI-V

Sequence:

TGĠTATCAGAGCAAGGTTAAACATGGTCATGTCCGGCTAAAAACCTAGTGTTAGGTTCGG GGAAAGAGGTCTACTGTGTTTCAAATATCCATTTGTTATTATCCAAACAGTGTCTGTTC CTTGTGTTTAGGATGGAAACCTGAGGTTGCGAACTACGTGTTCAAAGGAGTGATCATTTC AAAAGGAATGTCATTCTAACTATCGTATGCTCTAGACTTGCCCTGTTATAGCAACAGAGGA TCCTGTCTATCAGAAAATAGTTTGAACAATTCCGATGATCTCTCATACTTTAGGTATAACAA CGCCACGTATAAGGAGCTAATTCTTTTGTAAAAGACGTAACCCAGAAGAGAGTACCCAGA GAGGGGAAGACTTGGGGAGAAGGTGAACAGAAGAGAGAGCTTACTGTACCTGGTACTGT TATAGGAAGATTTTGTTATGTCCAGTCGCTGGGAGAATAGTATACAAGAGTGGTATGAGA AGTCACACACAGCAAACCTTGAGTACCTTGACCTAGCATCTACTAGTAAAGTGACCAACA ACCAGTTAGCACATAACCTTGCAGTAACCTTTGATAGAGTAAATTTAGGTAACCGAGTTTT TATTAAAAACCTGAAGCAGATTCAAGAGTCTATTCTTGAATTAAACACCCGTGTTGATACTA TAGAAGTAGCTCTAAGGAGGTTAACCAAGACCTTCAGAGAAAGCAAACCACTTACCGAAA GTGAAGTCCGTAAGTTGGTTGAAGAAATTGCTCAACAACCTAAGATCGTCGAGAAACAAG CACTCGAGATCTCTCAACAACTTGAGCAGAAACTAGAGAAGGTGGAAAAGCTCTTGCACA AACTTGACCAGTGGGTTGGTCAATGACTGATAGCCCAGCCTATCAAGAAGCTTTAAAGGA AGCAGAAAAGGTTGACCCACCAGCAGTTGGGATAACAACCTCCACCGGAGTAACAGCGG TTCAAGGGTTTAAAACCGTTATTAAGCAGAACAACGTTCAGATCTGCTTACTCGCTGTCAT AGGAAGTTGCCATTCCAGAAGACTTAATCACAAAACTCCAAGGATTATCTATTCAGGAGAA AGGAGAGGCAAAAGTCACAAGAAAACCCGAGCCAAAAGGAACGCTGTTTGGATTCAAAG ATCCTTACAAGATCCTAGCAGCAGAAAAGGCTAAGATCACACTTAAGCCTGTGAAAGAGT AGAAAGATGAGTCGAGCACGACCGCAACACCCAGTTCCTAGTGTGACTACAACCACTAGT GAGCAAAACAGGGAAGGACCTCTATACGAGGATCAGATCAGAGACTACCGAAGAGACA AAGAAGGATCTTTAACCTTCGAAGAAGAGCCAGAAGGTTAAGAAGATCAATGATGGGGTC TAGATATCAAGAGACCCTAGAACAAGAAATAGATCCACAGACAACACTGAGGTTGTCCAT GCAAGAACGAGCAAGACTAGTACCAGCTGAGGTACTGTACAGATCCAGGCGAGACACTG AAGTGGATAGGGCGTTTATACAGCCTGAAAGTCTGGAACAACTCCAAAGGACAGGAATGT CCTTTATTCATATCGGAATTCTGCAAGTGAGGATTCAGATTCTGCACCGACAAGAAGAAG GCACCATGGCCTTGGTAGTATTCAGGGATAACAGATGGTCGGGAGATCAGTCTATTTTCG CTCAAATGGAAATAGACCTAACAAAAGGAAGCCAATTGGTGTTCGTTATACCAGACACCAT AAATTGGCAGAACGGAGAAGCCAATCTCCTGATAACACGTGGCATGACAGGGCGACTGT CCAACACTCCCAATGTCGCCTTTGCTTACCAAATTGCCAGCGCAACAGATTACTTGGCAA GTCACGGTGTAAAAGCTATCGCAGGAAAGAAAATGAATCTGCAACACCTGCGAAACCAAC AGTGGATACTACGTCCACCACAAACGGACATCACTCCAATGCAACCAAGATCGGTTGAAA CAAGAAATCTCGTAGATGGAAGTATCTCCATCAGATTCCATGATTATGAGGCAGCTACTTC AGCTTCAAGACCTCACTACAATGAGGAAGATGAAGAAGTGGAATCAGAAACGGAGTCAGA AATAAGGGAGCATACTATTGCAGTCTGGATAGGAGAAGAAGAAATTCCAGACCAGACAGG GAGAAAGAAGGTATGGGAAGATCTAGTAATGGAAATGGAAGATTCTTCCGGTATTACAC TCCTCCACCAACATTTGAAGGGCAAATCATTGCTACAGGATGGGGAAGTGATGATAA TGAAAAAACTCCTCCAAAATGGGATGAAAGCCCAGATGAAGAAGGACCCACAGAACCCAT ATGGGATCAAGAAGAAGAAGATGAATATGATCCCAATGTCTATAGGGCATACTTACA AAAGGAGGAAGATGAGTGGCAAGAAATCACAGCTAGTCTCAGGGAAGAAATGGAGTACC CAAAAAGACGACCACAAACAGAGATGGCGTTCTCTGAAACAGTCGACTATACTCCACCTG GTGACACTATGATGACACCTGTCGGATACCCCCCGGCCTCGTCATCAAGATCAACAGTCA CAACGCCAAGTAGACCGCCACTTTTTGAAGGAAGGACCACACACGTGCCACGATTCCTAA AAAGGGATGAGTACACAGAATGGTGGCAATTACCATCATCGCAAGGCACAACAGGGGCG TTATTTGTAATGCCTAAACAAATGGGCCTATTCCATGAAGTCTTCTCGAGATGGGAATCCA TCACCAAAAACTACGTTGCGGCCCAAGGTTTTACGGACCCAACAGAAAAGATGGAGTTCA AGGCTGAATACCAGCAGCTGCTAACCCAAGCAGATGGGCGACAAGGAACCCAAAATATC TTGTCCCAGATCAAGAGAATCTTTTCTTTAGAAGACCCAGCCTCTGGATCCACGAGAATAC AAGATGCTGCATACAGAGACCTGGAGAGATTAACCTGCCACAACATAAAAGACATAGTTC

AATTTCTAAATGACTATGGAAGGCTTGCAGCAAAGAGTGGGCGACTGTTCTTAGGAACTG AGCTCAGTGAAAAGTTATGGATGAAGATGCCACCAGAACTAGGGCATCGAATGAAAGAAG CATTTCAAAAGGAGTATTCAGGCAATGAAGTAGGAGTCTTCCCGCGCATCTTGTTTGCCT ACAGATATCTGGAGCAAGAGTGTAAAGATGCAGCTTTTAAGAGAAGCCTGAAGTCATTAA GCTTCTGCAAAGACATGCCATTAACAGGATACTATGATAAAACCTCCAAGTATGGCATGA GAAAGTCAAGAACCTATAAAGGAAAACCACATGCATCACATGCAAGGGTGGAGAAGAGG AAGCACTTGATCAGAAATAAAAAGTGCAAATGTTATCTCTGCGGGGATGAAGGACACTTT GCCCGGGAATGCCCCAATCAGAGGAGAGATGTCAAGAGAGTAGCCATTTTTGAAGGAAT TGATCTTCCTGAAGGCTTTGATATCGTCTCAGTAGAAGAAGAAGAAGAAGAAGATGC TATTTATAGCATATCTGAAAATGAAGACGGAGAACTTGACACTGAAGTAGTCCATGAGAAA GTCTTCATGATGAGAGAGAGACCAGTCCTATTGGTTAGGAAAGACAAACCATTGGACG GAAATATTAGTGGTGGCCCACATCAACTGCCACTTCTGTAAGCAACCCACTCAGTTAAGG AGTCGAATACACTGTCCCACGTGTCAACTCACCAGTTGCTTCATGTGTGCCCCAATATACT GCAATATGATAGTCCAGCAGCAGCCTAAACCACCAGTACCGTTTAATACTCACACACTGC TCCAACAACAAGCGGCTTATATCCAGTGGTTGGAAAAAGAAAATCAGCGGTTAACTGAAG CCGTTGAATTCTATAAAAAGGAGGCTGAAGAATTAAGGCTCGAAAGAGACTTAGAGCAAG ATAGAAGGAGTCTGGAACCTACGTTGTTAGACAAAGGAAAGAAGGTTCAGATTCTTGATC CAGATGAAGATCAGCACACAGCGTATCTTGAAGAAGATACCATCAGCCGTGTTATCGGCC ATACTGTGGAACAACAAGAGGTTAGAAAGCCCCGTTAAAAAGGGGAAACATGCTCTATAACC TCGATGTGGTGTTACATATCCCAGAGGTAGGAAGACCTATCAAAGTCAAAGCAATTCTAG ACACTGGAGCAACCACATGCTGTATAAACATCAACTCTGTACCACAAACAGCAATTGAACA GAACACTTTTCTGGTACAATTCCGAGGCATAAATTCCACGCAATCTGTGGATAAGAAACTC AAATATGGGAGGATGACTATCAGCAATCACCAGTTCAGAATCCCGTACTGTTATGCCTTTC CTCTATCCCTTGGTGATGGAATAGAGATGATCCTAGGGTGTAATTTCATCCGTGGGATGT ATGGCGGTTTGCGTATTGAAGGTCACACAATCACCTTCTACAAAAATGTCACTACAATCCA AACCCGCCTTGCTGCCGTAATGGTTGGTGGTACAACCGCTTCTGAGTTAGGGGGAGGGG AGGAGTCCAAATCCGATTCTGAATCCATGTTTGACCTCTCAGAAACAGAAGAATTTGACTC AGAAACCCACCAGCAGATTGTGAGTCATGTTGCAGCCCAAGCCCAACAACAAAATTGGA TCCAAAACTCCAACAACTAATGGTCCAACTTCAGGATCAGGGCTTTATTGGGGAAAATCCT ATGCAACATTGGGCTAAAAACAAGATCCTATGTCGACTAGATATCAAGAATCCTGATCTTA TAATAGAAGACAAGCCCATCAAGCATCTAACACCGGCCATGGAGAAACAGTTCCAGAAGCACATCAAAGCACTCCTGGACATTGGTGTTATCAGGCCTAGTAAGTCAAAACACAGGACTA CGGCCTTCATTGTGGAATCAGGCACTGTTATTGATCCAGTCACGAAAAAAGACTATACACG GTAAAGAACGTTTGGTCTTCAACTACAAACGCCTGAACGACAACACTGAAAAGGACCAGT ACTCGCTACCCGGTATACAGACCATCCTAAAGCGGGTGGGCAACAAGAAGGTCTTCAGC AAGTTCGATCTAAAATCGGGCTTCCATCAAGTTGCCATGGCGGAAGAGTCCATCCCTTGG ACTGCTTTCTGGGTACCGCAGGGCCTTTATGAATGGTTAGTGATGCCCTTTGGGCTCAAA AACGCTCCTGCAGTATTTCAAAGAAAAATGGACCAATGCTTCAAAGGTACAGAAGAATTCA TTGCAGTGTACATTGATGACATTTTGGTCTTCAGCGAAAATATGGCAGAGCATACCAAGCA CATTGGAATCATGCTCAAGATCTGCCAAGAAAATGGGCTGGTACTAAGCCCAAGTAAGAT ATGTCTTGCCCAACGCGAGATTGAATTTTTGGGCACAGTCATCTCACAAGGACAAATGAA GCTTCAGGCCCATGTAATCAAAAAGATAGTCAACAAAGCCAACATAGAGCTAGAAACAAC AAAAGGCCTGAGATCCTTTTTGGGCCTCCTCAACTATGCCCGTATCTACATACCCAACCT GGGTAGAAAGCTAAGTCCACTATATGCCAAGACTAGTCCCACCGGAGAGAAAAGGTTTAA TCGACAAGATTGGCATCTGATAAAGGAAATCAAAGATATGGTCCAAAAGCTCCCAAACCT CGCTATCCCACCAGCAAGATGCTACATTATCATTGAAAGCGATGGCTGCATGGAAGGATG GGGGGCCGTATGTAAGTGGAAATTAGCAAAAGAAGATTCCCGCACTACTGAAAAGATTTG TGCGTACGCTAGTGGGAAATTCGGTGTTGTCAAGTCTACCATCGACGCCGAAATTTACGC ACTCATAAAAGCATTGGAATCTTTTAAAATCTTCTATCTGGACAAAAAACATTTGGTGGTGC GAACAGACTGTCAAGCGATAGTGACATTTTACAACAAGACAAGTACTCATAAACCCAGTA GAATACGTTGGATCACCTTTTCCGACTATATAACGGGGGTTAGGAGTTCCGGTTACTATCG AACACATAGACGGAAAAGAGAACCAGTTGGCTGATACACTAAGCAGACTCGTGTATACAA CATGGAACCAGTCCCAAACTCACCAACCGGAGGAAGAAGAGCTGGAGAAGTCCCAACAT CTCAGCTTCGCGGGGCTAGCTATCCCTATAGCTTGGCCTATGATGGGCTCCTACAACAA AGAAGGACGCCATTACTCACGGGACAATCACTCTGGCAACGGAACAAGCCATCTCAGCA CAGCTCTACCGCATCGAAGAGCAGGCAGCCAAGAAAGCACTATTGGCCCTACGTGACCT ACAGGGCGTACTCCACTTCAAGAGAGACTATTTGGCCGCTACTGCCACTAGAGACAACTG GGCTAGCGACAGACTGCCAGCCCAACAAGACTCAGCCGCCCTAGACCAACATGCTG GCGTGATTAACGCCATTATTGAAAGGGCTGTCCAACCCTAGTTTGGACGGTAGTAGTAGG TGTAATAATAGTTAGGTGTGCTTTACTTTTCCAAGCTGTCACTCATTATAGAGTAGACA TGATGATCGACGATGGGGCCCAATGAGCACCCGGATCACCATTCTCCCATCTATAAATGA GAGTTTGTAAGGCTTAGCCATCAGAGAGTGAAAACTACTCAACTGATCCTAAGTGTTAGA GTTTGTATTTTCCTAAGAGTTTGTAAGATTTTTATGAAATAAAGAGTCTACTTTGTGTTTATC

TCTTTGTTTCACCTGGGATTTAAACAGTTTTTGTTTTTCCGCACCATCGGTTTGCGCCCGA TCGATGT

Structure in dot-bracket format:	
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Energy: -2056.9

RNA-Structure:



Virus Domain(s)

Number of proteins:

Protein 1

Number of domains:

Aminoacidsequence:

MAGAASPCANGCGPSAPSDAEVVHLCRSLEVGTVMTLFYSKKSQRPERKTFQVKLETRQIT WSRGADKIEGAIDIREIKEIRPGKTSRDFDRYQEDPAFRPDQSHCFVILYGMEFRLKTLSLQAT SEDEVNMWIRGLTWLMEDTLQAATPLQIERWLRKQFYSVDRNREDRISAKDLKNMLSQVNY RVPNMRFLRERLTDLEQRTSDITYGQFAQLYRSLMYSAQKTMDLPFLEASALRAGERPELCR VSLPEFQQFLLEYQGELWAVDRLQVQEFMLSFLRDPLREIEEPYFFLDEFVTFLFSKENSIWN SQLDEVCPDTMNNPLSHYWISSSHNTYLTGDQFSSESSLEAYARCLRMGCRCIELDCWDGP DGMPVIYHGHTLTTKIKFSDVLHTIKEHAFVASEYPVILSIEDHCSIAQQRNMAQYFKKVLGDTL LTKPVDIAADGLPSPNQLKRKILIKHKKLAEGSAYEEVPTSVMYSENDISNSIKNGILYLEDPVN HEWYPHYFVLTSSKIYYSEETSSDQGNEDEEEPKEASGSTELHSNEKWFHGKLGAGRDGRH IAERLLTEYCIETGAPDGSFLVRESETFVGDYTLSFWRNGKVQHCRIHSRQDAGTPKFFLTDN LVFDSLYDLITHYQQVPLRCNEFEMRLSEPVPQTNAHESKEWYHASLTRAQAEHMLMRVPR DGAFLVRKRNEPNSYAISFRAEGKIKHCRVQQEGQTVMLGNSEFDSLVDLISYYEKHPLYRK MKLRYPINEEALEKIGTAEPDYGALYEGRNPGFYVEANPMPTFKCAVKALFDYKAQREDELTF TKSAIIQNVEKQEGGWWRGDYGGKKQLWFPSNYVEEMVSPAALEPEREHLDENSPLGDLLR GVLDVPACQIAVRPEGKNNRLFVFSISMASVAHWSLDVAADSQEELQDWVKKIREVAQTADA RLTEGKMMERRKKIALELSELVVYCRPVPFDEEKIGTERACYRDMSSFPETKAEKYVNKAKG KKFLQYNRLQLSRIYPKGQRLDSSNYDPLPMWICGSQLVALNFQTPDKPMQMNQALFLAGG **HCGYVLQPSVMRDEAFDPFDKSSLRGLEPCAICIEVLGARHLPKNGRGIVCPFVEIEVAGAEY** DSIKQKTEFVVDNGLNPVWPAKPFHFQISNPEFAFLRFVVYEEDMFSDQNFLAQATFPVKGLK TGYRAVPLKNNYSEGLELASLLVKIDVFPAKQENGDLSPFGGASLRERSCDASGPLFHGRAR EGSFEARYQQPFEDFRISQEHLADHFDGRDRRTPRRTRVNGDNRL

Starting nucleotide position:

Ending nucleotide position: 1337



Domain 1 Starting aminoacid position: 245 Ending aminoacid position: 318 Domain description: Phosphoinositide-specific phospholipase C, efhand-like Identifier: EF-hand_like Domain 2 Starting aminoacid position: 322 Ending aminoacid position: Domain description: Phosphatidylinositol-specific phospholipase C, X domain Identifier: PI-PLC-X Domain 3 Starting aminoacid position: Ending aminoacid position: 639 Domain description: SH2 domain Identifier: SH₂ Domain 4 Starting aminoacid position: 668 Ending aminoacid position: 741

Domain description:

SH2 domain

Identifier:

SH₂

Domain 5

Starting aminoacid position: 797

Ending aminoacid position:

843

Domain description: SH3 domain
Identifier: SH3_1
Domain 6
Starting aminoacid position: 952
Ending aminoacid position: 1070
Domain description: Phosphatidylinositol-specific phospholipase C, Y domain
Identifier: PI-PLC-Y
Domain 7
Starting aminoacid position: 1090
Ending aminoacid position: 1177
Domain description: C2 domain
Identifier: C2