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#RELEASE DATE: UniProtKB/Swiss-Prot Release 2012_03 of 18-Apr-2012, UniProtKB/TrEMBL Release 2012_03 of 18-Apr-2012
#DATA FILES ANALYZED: uniprot_sprot_viruses.dat AND uniprot_trembl_viruses.dat
#SUMMARY: Total number of predictions in this clade (115)

>Ostreococcus virus OsV5: Total=1
      A9YW33_OSV5 Window Position=177; Score=52.557 | Prion Domain: TNNYKMNTTTNNNYKVNTNNTNNYKMNTTTNNNYKVNTNNTNNYKMNTTTNNNYKVNTNN
>Heliothis zea virus 1: Total=2
      Q8JKQ9_9VIRU      Window Position=551; Score=88.809 | Prion Domain: NDNSNFNYNNNYNNNYNNSSYSNSSYNNNNYNNNYNNPNFNNSFNHPNSFNQNQNNYNSY
      Q8JKP9_9VIRU      Window Position=501; Score=67.929 | Prion Domain: NNQTSNNQTSSQSNNQNSQSSQSNESTSNDSSSNNSSSNSQSTVVNVNAQSNDNTQSNNN
>Moumouvirus Monve: Total=4
      H2EE98_9VIRU      Window Position=387; Score=69.388 | Prion Domain: NNNNNPMGMNNMNNMMGMNGMMGMNGMNGMNGMNNMMGMNMGNMNGMNNMSAMGNMMGMN
      H2EEG5_9VIRU      Window Position=497; Score=76.823 | Prion Domain: GSSGGFNGNNNNNNNNQNMRNGQNNNNNMRGGQYNNNNNNNNNNNMQGGHHDPWANYGNQY
      H2EE16_9VIRU      Window Position=134; Score=54.622 | Prion Domain: RNNNYNNNFNNNFRNNFRNNNNFRNNNNFRNSDSRFYNNSDPFNKNGFTKNRFNNGQSFM
      H2EF52_9VIRU      Window Position=219; Score=55.120 | Prion Domain: YSSGNNNNSGFGSGSNSRSNNNNSTNTGSRSTANANRNANTMSGNTSGNTSLYTGNNNNN
>Acanthamoeba castellanii mamavirus: Total=9
      G8ECX9_9VIRU      Window Position=34; Score=55.428 | Prion Domain: NNNVHVANNLNNFVQQHKQLQNQMPQFQNQFQNQSMNQSMNQSMNQQINRPMNQSMNQSM
      G8ECX7_9VIRU      Window Position=172; Score=57.024 | Prion Domain: NYNHNFNNNHNFNNNFTNNRMTDYSTDYISHNFNPGNNQYSQQISHEQNINQQQSQNSEN
      G8ED73_9VIRU      Window Position=93; Score=59.911 | Prion Domain: NSSNQSSQSNQVNQSNQSSPSSQISPSSQVNKFNQSSQINQSNQINQSNQINQSNQLNHH
      G8EBH5_9VIRU      Window Position=682; Score=53.032 | Prion Domain: NNVVQQYQSYYDNPSNQQSNQQSNQQSNQQPNQQPNQQPNQQPNQQPNQQSVQQCNQVNQ
      G8EDD3_9VIRU      Window Position=310; Score=51.571 | Prion Domain: NQYNQIGSQGQNRDYRQNNSNHSNNSNHSNHVDRTDRTNQSTYTNQTNLTNQSNRSNYSN
      G8EDT4_9VIRU      Window Position=75; Score=62.116 | Prion Domain: NDNQNLKNFNAINNNFNNDYNNDYDSNNDSNNDSNNDSNNDYDNNSNNYFNNDSNNDSNN
      G8ED52_9VIRU      Window Position=17; Score=83.492 | Prion Domain: IAANNNLLNNKNNQTNQLNNNQYNNQYNNQNYPQNYPQNSQQNFQQNSQQNHQQNYQQNY
      G8EBE3_9VIRU      Window Position=250; Score=50.770 | Prion Domain: SQNNNLSAMNNSLYNNNKAVNTNTINSTNNRNVSSQNNANRNASMATTYNNSVNSANSIN
      G8EDB6_9VIRU      Window Position=264; Score=51.349 | Prion Domain: NNNNTTTSNLNNSTIISGLNNATNTSNLNNAVSTLDNNNTNISNFNNATNTRNNNNNATNN
>Ectocarpus siliculosus virus 1: Total=1
      Q8QNJ9_ESV1K      Window Position=21; Score=76.395 | Prion Domain: QHQQQMQQRQAQQRQMQMQMQMEQQQQQRILQQQQQQQQQQQRILQQQQQQQQQQQQQQQQ
>Zantedeschia mild mosaic virus: Total=2
      Q4VU86_9POTV      Window Position=2856; Score=86.671 | Prion Domain: QQQQQQQKQQQQQDQQQQQQQQQNQQQQQQQQQNQQQQQQQNQNKTVDAGNNQNKQKSPA
      Q5GJ53_9POTV      Window Position=83; Score=90.598 | Prion Domain: QQQQQQQQQQQQQDQQQQQQQQQNQQQQQQQQQNQQQQQQQNQNKTVDAGNNQNKQKSPA
>Emiliania huxleyi virus 202: Total=1
      G8DF10_9PHYC      Window Position=48; Score=55.890 | Prion Domain: DRTRGGGGNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNTATVMDLRSATTYQDDPNIFDKQA
>Thermus phage TMA: Total=1
      F9VHW2_9CAUD      Window Position=220; Score=95.846 | Prion Domain: NNQGGQQNQNNNQQIQGNQNSQQNQGSQDSQQNQGGQNNQQNQGQNNSNQQGGQNNQQEQ
>Paramecium bursaria Chlorella virus MT325: Total=1
      A7IUH2_PBCVM      Window Position=35; Score=58.779 | Prion Domain: NGNWGNSGNWGETNNNWWNNGNNNGWNNNNGNNNNGNNNNGNNNNGNNNNGRLNVVKKNN
>Megavirus courdo7: Total=8
      H2EB66_9VIRU      Window Position=60; Score=72.442 | Prion Domain: TTYLDNATNNNNNNNNNNNNNNNNNNNNIYGGKNNWNNNNNQWNNKPKSNNFNNTKNNRT
      H2EBE2_9VIRU      Window Position=101; Score=61.668 | Prion Domain: NNIHDDINDNNDNDSYGDNDNDNNNDNNNDNNNDNNNDNNNDNNNDNNNDNNNDNNNDNN
      H2EBG2_9VIRU      Window Position=40; Score=72.887 | Prion Domain: NNQTNNQHNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQF
      H2EBD5_9VIRU      Window Position=201; Score=72.546 | Prion Domain: NNNNNNTNNTDKNIDDMTNNNNNGNNNSGSNNNNNNNDSNSNSNDSNTTNNTTNNTTNNKNI
      H2EB64_9VIRU      Window Position=221; Score=69.723 | Prion Domain: NNKSGPNNVWNNSQIMGGEDNWNNTNNNMNNNTNNNTNNNTNNNTNNNTNNNWKNNNKNNNNNN
      H2EAX4_9VIRU      Window Position=108; Score=57.880 | Prion Domain: QQVMMQNNIRNGNGQNMNQGMNQGMNQGMNQKMNQMYGQQNQSINPMFLRQQILNNAQSN
      H2EAZ5_9VIRU      Window Position=138; Score=76.252 | Prion Domain: NNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSSSNNRSNS
      H2EBM3_9VIRU      Window Position=156; Score=54.325 | Prion Domain: FQNKNPQYNYYNPRSRNNPTRPNYNYYNSTQNNGYQNVNSMRGYQQNQYQNQNQYQGQNN
>Lactococcus phage u136.13: Total=1
      Q0ILF2_9CAUD      Window Position=104; Score=83.784 | Prion Domain: SRSQGQQQQQQQGNYNQNQNNYQNQGQQNTVQQQHNQGNYQRQPQGNYQNQQQSAQQRAQQ
>Organic Lake phycodnavirus: Total=1
      F2Y2X1_9PHYC      Window Position=145; Score=56.591 | Prion Domain: MNVGNNQRMNNRRMNNQNNNQRMNNQNNNQRMNNQPMNVGNNNRRLSKNNIFMTNRRSNI
>Megavirus chiliensis: Total=10
      G5CT61_9VIRU      Window Position=108; Score=62.274 | Prion Domain: QQVMMQNNIRNGNGQNMNQGMNQGMNQGMNQNMNQMYGQQNQSINPMFLRQQILNNAQSN
      G5CQW2_9VIRU      Window Position=136; Score=51.924 | Prion Domain: NNDDNNDNDNDNNNDDNDNDYNNNDNDDDNNDDNNDDNNDHYNNDDNNNDNDNNENNNNNNF
      G5CQD2_9VIRU      Window Position=65; Score=54.491 | Prion Domain: NDNENENEKNQLDNLSQNNNSNDDQNNSDNNDNNDNDINDNNNDNNNNNNNNDNNNDESS
      G5CT78_9VIRU      Window Position=137; Score=63.311 | Prion Domain: NNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSNSNNRSSSNNRSNNRSNSNDKLDSNNR
      G5CT16_9VIRU      Window Position=35; Score=67.224 | Prion Domain: IDRSDNNQTNNQHNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQFNNQ
      G5CST1_9VIRU      Window Position=314; Score=60.103 | Prion Domain: STNMNGQNNMGMNGMNNMGMNNMGMNNMGMNNMGMNGMNGMNNMGMNGMNNMGMSGMNGMNMGGMN
      G5CST2_9VIRU      Window Position=201; Score=76.572 | Prion Domain: NNNNNTNNTDKNIDDMTNNNNNGNNNSGSNNNNNNNNDNSNSNDSNTTNNTTNNTTNNKN
      G5CSG7_9VIRU      Window Position=243; Score=93.576 | Prion Domain: NNTNNNMNNNTNNNTNNNTNNNWNNNNQNNNNNNNWNNNLMGGENNNTNMNNNSNYWNNNN
      G5CT03_9VIRU      Window Position=90; Score=51.635 | Prion Domain: NNYTGINNFSNNIHDDINDNDDNDSNGDNDNDNNNNDNNDNNNDNDNNNDNNNDNNNDN
      G5CQ80_9VIRU      Window Position=162; Score=54.325 | Prion Domain: FQNKNPQYNYYNPRSRNNPTRPNYNYYNSTQNNGYQNVNSMRGYQQNQYQNQNQYQGQNN
>Helicoverpa zea nudivirus 2: Total=3
      G9I0B0_9VIRU      Window Position=543; Score=89.533 | Prion Domain: NDNSNFNYNNNYNNNYNNSSYNNSSYNNNNYNNNYNNTFNFNNSFNHPNSFNQNQNNYNS
      G9I0C0_9VIRU      Window Position=364; Score=59.542 | Prion Domain: NQTYNQNYNHQNYNQNYNNQNYKQDYNQNYNHQTYNHQNYTPSSYNHHQNYHQKKYPPNY
      G9I0A1_9VIRU      Window Position=504; Score=63.835 | Prion Domain: NNQTSNNQTSSQSNNQNSQSSQSNESTSNDSSSNNSSSNSQSTVVNVNTQSKDNTQSNNN
>Porcine epidemic diarrhea virus: Total=34
      NCAP_PEDV7  Window Position=151; Score=62.790 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSNNRNQSN
      G8GMX6_9ALPC      Window Position=151; Score=56.671 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      E2FYM9_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      Q692L9_9ALPC      Window Position=152; Score=58.085 | Prion Domain: GNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSND
      G9G5U2_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMZ2_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      A7LKV4_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMW1_9ALPC      Window Position=151; Score=54.443 | Prion Domain: RGNGNNRSRSPSNNRGNNHSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      Q9IH81_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMY4_9ALPC      Window Position=151; Score=55.679 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGDSQNRGGNNNNNNKSRNQSKSRNQSN
      Q4PKJ1_9ALPC      Window Position=151; Score=53.836 | Prion Domain: RGNGNNRSRSPSDNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSGNQSN
      D3JVE5_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      F5CHU8_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      F5AU33_9ALPC      Window Position=151; Score=51.314 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNPQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      A2IB79_9ALPC      Window Position=151; Score=62.790 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSNNRNQSN
      Q6QIL5_9ALPC      Window Position=141; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMY2_9ALPC      Window Position=151; Score=54.893 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGYNNNNNKSRNQSKSRNQSN
      G8GMX9_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMY9_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMX8_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      F5CIA0_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      F2VPY5_9ALPC      Window Position=151; Score=62.790 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSNNRNQSN
      E2FYM7_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      B7UCL9_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMX7_9ALPC      Window Position=151; Score=56.671 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGYNNNNNKSRNQSKNRNQSN
      B7UCM9_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      B7UCL7_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
      F5AU32_9ALPC      Window Position=151; Score=55.590 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNREGNNNNNNKSRNQSKNRNQSN
      G8GMY6_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNSRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      G8GMY3_9ALPC      Window Position=151; Score=50.940 | Prion Domain: RGNGNSRSRSPSNNRGNNHSRGNSQNRGNNQGRGASQNRGGYNNNNNKSRNQSKSRNQSN
      B7UCN1_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      B7UCM7_9ALPC      Window Position=151; Score=52.600 | Prion Domain: RGNGNNRSRSPSDNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      B7UCM1_9ALPC      Window Position=151; Score=56.618 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKSRNQSN
      Q2I0J9_9ALPC      Window Position=151; Score=58.396 | Prion Domain: RGNGNNRSRSPSNNRGNNQSRGNSQNRGNNQGRGASQNRGGNNNNNNKSRNQSKNRNQSN
>Ralstonia phage RSL1: Total=1
      B2ZXU0_9CAUD      Window Position=162; Score=66.197 | Prion Domain: NNNANANSNVNGNTNLNGNSNHNQSSASNSNANTAQGGAGGNASNTNTNSNANNGNNSAQ
>Bathycoccus sp. RCC1105 virus BpV2: Total=1
      E5ER44_9PHYC      Window Position=145; Score=54.408 | Prion Domain: TNNVNNVNNVNNGTRLNNNVNNGNNVNNGTRLNNNVNNGTRLNNNVNNGTRLNNNLNNGT
>Acanthocystis turfacea Chlorella virus 1: Total=1
      A7K957_9PHYC      Window Position=3; Score=69.616 | Prion Domain: TQSKKQEPVYNTRKNNNYNNNNNNNNNNNNNNYNNNNNYNNNYNNNNDDWNWNSWNEGNN
>Paramecium bursaria Chlorella virus FR483: Total=1
      A7J7F5_PBCVF      Window Position=13; Score=56.270 | Prion Domain: FNNNNNFNNNNNNGDWDWNNWENGNWGNNSNWGETNNNWWNNGNNNNGNNNNNGNNNNNGR
>Acanthamoeba polyphaga mimivirus: Total=19
      YR558_MIMIV Window Position=75; Score=50.350 | Prion Domain: NDNQNLKNFNTTDNNFNDDYNNDYDSNNDSNNDSNNDSNNDYDNESNNYFNNDSNNDSNN
      YL265_MIMIV Window Position=34; Score=55.428 | Prion Domain: NNNVHVANNLNNFVQQHKQLQNQMPQFQNQFQNQSMNQSMNQSMNQQINRPMNQSMNQSM
      YR722_MIMIV Window Position=682; Score=53.032 | Prion Domain: NNVVQQYQSYYDNPSNQQSNQQSNQQSNQQPNQQPNQQPNQQPNQQPNQQSVQQCNQVNQ
      YR692_MIMIV Window Position=256; Score=50.770 | Prion Domain: SQNNNLSAMNNSLYNNNKAVNTNTINSTNNRNVSSQNNANRNASMATTYNNSVNSANSIN
      YL356_MIMIV Window Position=93; Score=59.911 | Prion Domain: NSSNQSSQSNQVNQSNQSSPSSQISPSSQVNKFNQSSQINQSNQINQSNQINQSNQLNHH
      YL264_MIMIV Window Position=546; Score=57.024 | Prion Domain: NYNHNFNNNHNFNNNFTNNRMTDYSTDYISHNFNPGNNQYSQQISHEQNINQQQSQNSEN
      YL399_MIMIV Window Position=264; Score=51.349 | Prion Domain: NNNNTTTSNLNNSTIISGLNNATNTSNLNNAVSTLDNNNTNISNFNNATNTRNNNNATNN
      YR335_MIMIV Window Position=20; Score=94.934 | Prion Domain: NNNLLNNKNNQTNQLNNNQYNNQYNNQNNNQNYPQNYPQNSQQNFQQNSQQNHQQNYQQN
      YL417_MIMIV Window Position=352; Score=51.571 | Prion Domain: NQYNQIGSQGQNRDYRQNNSNHSNNSNHSNHVDRTDRTNQSTYTNQTNLTNQSNRSNYSN
      E3VZY1_MIMIV      Window Position=682; Score=53.032 | Prion Domain: NNVVQQYQSYYDNPSNQQSNQQSNQQSNQQPNQQPNQQPNQQPNQQPNQQSVQQCNQVNQ
      E3VZI3_MIMIV      Window Position=546; Score=57.024 | Prion Domain: NYNHNFNNNHNFNNNFTNNRMTDYSTDYISHNFNPGNNQYSQQISHEQNINQQQSQNSEN
      E3VXS9_MIMIV      Window Position=352; Score=51.571 | Prion Domain: NQYNQIGSQGQNRDYRQNNSNHSNNSNHSNHVDRTDRTNQSTYTNQTNLTNQSNRSNYSN
      E3VZZ3_MIMIV      Window Position=20; Score=94.934 | Prion Domain: NNNLLNNKNNQTNQLNNNQYNNQYNNQNNNQNYPQNYPQNSQQNFQQNSQQNHQQNYQQN
      E3VXK5_MIMIV      Window Position=264; Score=51.349 | Prion Domain: NNNNTTTSNLNNSTIISGLNNATNTSNLNNAVSTLDNNNTNISNFNNATNTRNNNNATNN
      F8V6N0_MIMIV      Window Position=250; Score=50.770 | Prion Domain: SQNNNLSAMNNSLYNNNKAVNTNTINSTNNRNVSSQNNANRNASMATTYNNSVNSANSIN
      E3VZI4_MIMIV      Window Position=34; Score=55.428 | Prion Domain: NNNVHVANNLNNFVQQHKQLQNQMPQFQNQFQNQSMNQSMNQSMNQQINRPMNQSMNQSM
      E3VZ03_MIMIV      Window Position=75; Score=50.350 | Prion Domain: NDNQNLKNFNTTDNNFNDDYNNDYDSNNDSNNDSNNDSNNDYDNESNNYFNNDSNNDSNN
      E3VZS4_MIMIV      Window Position=256; Score=50.770 | Prion Domain: SQNNNLSAMNNSLYNNNKAVNTNTINSTNNRNVSSQNNANRNASMATTYNNSVNSANSIN
      E3W038_MIMIV      Window Position=93; Score=59.911 | Prion Domain: NSSNQSSQSNQVNQSNQSSPSSQISPSSQVNKFNQSSQINQSNQINQSNQINQSNQLNHH
>Paramecium bursaria Chlorella virus AR158: Total=1
      A7RBX9_PBCVA      Window Position=20; Score=50.085 | Prion Domain: VNNKTSNNSNDNWNNSNWNWNNDDNAWNNWGNWNNSNWNNGNWNNNTNNNNNNNESNNKN
>Molluscum contagiosum virus subtype 1: Total=2
      Q85285_MCV1 Window Position=585; Score=65.176 | Prion Domain: AQQAQAQQAQQAQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQAQQ
      Q98177_MCV1 Window Position=558; Score=61.936 | Prion Domain: QQAQAQQAQAQQAQAQQAQAQQAQAQQARAQQAQAQQAQQAQQAQQAQQAQQAQQAQQAQQ
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>Thermus phage phiYS40: Total=1
      A0MNC5_9CAUD      Window Position=220; Score=90.827 | Prion Domain: NNQNSQQNQGKQDNDNQQNQEENNQQNQGNNQQNQGGNNQQNQGVQNNNNQEQNNNQQGQ
>Staphylococcus phage CNPH82: Total=1
      A1BTY4_9CAUD      Window Position=69; Score=51.391 | Prion Domain: NNDTTSNKSESTSKNDNSKTQSNNGNNERSQGKQAQATQQKNNQQQANNNQQMQNNNGYM
>Camelpox virus: Total=1
      Q8V2P6_CAMPM      Window Position=75; Score=91.777 | Prion Domain: NTNNANTDSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSNSN
>Chrysodeixis chalcites nucleopolyhedrovirus: Total=1
      Q4KT76_9ABAC      Window Position=956; Score=72.008 | Prion Domain: SYNNGYNNGRYNNGGYNNGGYNNGGYNNRYNNGYNNGYNGYNNYNNGYNNGNNGYDGVYN
>Apocheima cinerarium nucleopolyhedrovirus: Total=1
      D7F5T6_9ABAC      Window Position=35; Score=62.796 | Prion Domain: NYHRQFQQQNQQQQPNNIGSQFQQNNQQRFNSGGGNMGHHQFGNSNNAMNIGGNNAGSSN
>White spot syndrome virus: Total=3
      Q91LF6_WSSV Window Position=1031; Score=62.943 | Prion Domain: QQQQQQRDQQQQEQQQREQQQQQQQREQQQQREQQQQQQQQQSDQFRQQLLQQQQQFQQL
      Q8VB53_WSSVS      Window Position=1030; Score=62.943 | Prion Domain: QQQQQQRDQQQQEQQQREQQQQQQQREQQQQREQQQQQQQQQSDQFRQQLLQQQQQFQQL
      Q8QTE3_WSSV Window Position=1029; Score=62.943 | Prion Domain: QQQQQQRDQQQQEQQQREQQQQQQQREQQQQREQQQQQQQQQSDQFRQQLLQQQQQFQQL
>Vibrio phage VP16C: Total=1
      Q6VSX6_9CAUD      Window Position=261; Score=68.325 | Prion Domain: QQQQQQQQQQQQQWGQPQQQQQQQQQQQQWGQPQQQQQQQYEQPQQQQQQQQYEQPQQQQYNN
>Paramecium bursaria Chlorella virus 1: Total=1
      Q98476_PBCV1      Window Position=18; Score=66.067 | Prion Domain: NNNNNNNNNWNNNNNNNNNNGWDPNAWNNWNNSNNNYSNNWNFEEWGNFNNYNWNNNGNNNG
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