```
CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
CISFV-NC_021069.1
                                AAUUUUUGCCGAGAGACCGUGA----GGUUGAACCCGGCAAGGAAUGAAGAGGAAGG
                                AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA-
                                AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC---
                                AAGGCUUGAGUUGUUU - AUAAUAGUCGUUUUUCUCGCAGAA - AUGGGAAAGGACGACG
AA - UUUCGCUACCUU - AUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGAA-
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_034204.1
CISFV-NC_034017.1
                                UA--AUUUGCUACUUU--AUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGA--
                                                                                                                                  56
                                ----CUUUGGUUGUUUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAG
                                UAGACGUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAG
                                                                                                                                  56
                                ---AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAA--CGG
---CUAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUC--GAG
                                                                                                                                  54
cISFV-NC 027817.1
                                ----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAG
                                       .....10.......20.......30........40.......50........6
                                 cISFV-NC_001564.2
cISFV-NC_005064.1
cISFV-NC_012932.1
cISFV-NC_008604.2
                                AUUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGAUUUUUCCACCCCGUUCUGGGGAANNN 113
                                UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNN
                                GUAAGAAGAAGAGGGCCGGGCUCCUCCGGAUGGU-UGUUACCA--CCGGGGCGCGNNN 113
CISFV-NC_008604.2

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_034204.1

CISFV-NC_034017.1
                                GUCGGGAGUAGUGGAGAGAGGACGCGGUACGCCC-UCCGACCAACCGUGGAGGAANNN 113
                                GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCC-UCCUGCCAAUCGUGGAGGAGNNN 113
                                GAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCC-CCCUCCCAUUAAGGAGAAAANNN 113
                                CACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGGCUUUGAG--ANNN
                                AAAGUUUUCCCGACCGGGAGGUGGUCGGCCCAGAGA-GGGGCUGGGGAAGANNN 113
GAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGCCCAAAGA-GGGGCUGGCGAAGANNN 113
cISFV-NC_027817.1
                                GCCUUCUUCCCCUCGAGGGAAGAAAAGAAAGCCCCUGUGA---UACAAUCACAGGNNN
                                0.......70.......80........90........100.......110.......1
                                 .....))).(((((.....))))).
                               NNNNCUUUCAAAACUGGACGGGGCAACAGGGAAAUCCCG GG-UA-GCGAACCU CU NNNN--UUCAAAACCGGACGAACAGGGA-AUUUCCCG GGA-C-GCGUACCC CU NNNN--A-AAAAUCGGGCGAGCAACAGGGAGAAAUCCCG GGAAUU-GCGAACCC CU NNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA-AUCCA GUGACU-GACGCCAC UC NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCCG GUGAGU-GACGCCAC CC NNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAAUAUCCG GUGAGU-GACGCCAC CC NNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAAUAUCCG GUGAGU-GACGCCACCC
CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
                                                                                                                                 170
                                                                                                                                 167
                                                                                                                                 168
CISFV-NC_008604.2

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_034204.1
                                                                                                                                 172
                                                                                                                                 171
                               NNNN--ACCGCAAGGUGAGGGCGCACAGGACGAAAGUCCAGGGUGAGU-GACGACAC CC
NNNNAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCCGGGUGAUU-AGCCACAC CC
NNNN--CUC----CAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCA CC
                                                                                                                                 170
                                                                                                                                  172
                                                                                                                                 166
cISFV-NC_034017.1
cISFV-NC_027817.1
                                NNNNACCUC----CAGAGCAAA<mark>C</mark>GAGG<mark>G</mark>AGAAAUCCC<mark>U</mark>-A<mark>G</mark>GUGAGUUGGCGACCA<mark>C</mark>CC
                                                                                                                                  167
                                NNNNAUGUG----CAGGGCAACAAGUUCUAACGAACUA<mark>G GU</mark>GAGUAG-CGUC<mark>AC C</mark>C
                                                                                                                                 167
                                20......130......140......150......160......170......1
                                CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
                                228
                                                                                                                                  225
                                                                                                                                  226
                                                                                                                                  227
CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
                                                                                                                                  228
                                CCG-AAAU<mark>G</mark>UGUAUUGUGA<mark>C</mark>A-G-GCUUGCUCGCAGCAGAAUCUCC</mark>ACCGUUA
CCG-AAAU<mark>G</mark>UGUAUAUAGC<mark>C</mark>C-AUGUUGAGUCUAUACAACAGGCUUCCACCGGUA
CCG-AAACGUGAUUUAUAUGA-UGACAAGAAUCAGAUGGAAGA<mark>U</mark>CCCCACCGUUA
                                                                                                                         G-A
                                                                                                                                  227
                                                                                                                         GAA
                                                                                                                                  228
                                                                                                                                  227
                                                                                                            ACCGUUA
cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_027817.1
                                UCUGAAAU<mark>G</mark>UUAAUAAAAC<mark>U</mark>GUGCGGAA-AAAUCCAAAGACUGUCC
UCUGUAAU<mark>G</mark>UUUAAUAAGU<mark>UG</mark>UGCGGAA-AAAUCCAAAGAAUGUCC
                                                                                                            ACCGAUA
                                                                                                                                  225
                                                                                                            ACCGAUA
                                                                                                                         GGG
                                                                                                                                  226
                                CCGGUUG<mark>UG</mark>AAAACGAUUG<mark>CG</mark>ACUAGAACUAAAGUCGAGAGUC<mark>U</mark>CC<mark>CC</mark>ACCGAUA
                                80......190......200.......210.......220.......230.......2
                                ))))).......
cISFV-NC_001564.2
cISFV-NC_005064.1
cISFV-NC_012932.1
                                UUUUGAUUAUCUUUUCUAUACCAUAGAUGC-- 258
UUUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU 258
                                UUUG<mark>G</mark>AAUACCUUUUCUAUACCAUAGAUGCGC-
                                                                                      258
cISFV-NC_008604.2
cISFV-NC_021069.1
                                ACUU<mark>G</mark>-UUGCGGGCAAGCUCUUGUUCACGU-CU
                                AUUUG-UUGUGAGCAUUG-CUUGUCUACGU-CU
                                                                                      258
cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
                                AUUU<mark>G</mark>-UUGUGAGCAUUGCCUUGUCUACGU-CU
                                                                                      258
                                CGUU<mark>G</mark>-UUGUGAUGACACAUGAG-CAACGU-CU
GGAUC-CAUCCAGCAUUCUUGGAAUCAUGG-CU
cISFV-NC_034204.1
                                CUUUGGGAAAGUUUUUCAACACACGUAGCGUCU
                                                                                      258
cISFV-NC_034017.1
cISFV-NC_027817.1
                                                                                      258
                                CUUU<mark>G</mark>GGAAAGUUUUUCAACACACGUAGCG-CU
                                GCUCGUCAGUUUGUUUUAGUCCAAUUA--CGCU
                                                                                      258
                                40......250......260......270.
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