```
CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
CISFV-NC_021069.1
                          AAUUUUUGCCGAGAGACCGUGA----GGUUGAACCCGGCAAGGAAUGAAGAGGAA-GGA-
                          AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGG-AAG-
                                                                                                        54
                          AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAU-GCU-
                          AAGGCUUGAGUUGUUU -- AUAAUAGUCGUUUUUCUCGCAGAA -- AUGGGAAAGGA - CGACAA -- UUUCGCUACCUU -- AUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAA - GGA
CISFV-NC_012671.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_034204.1
CISFV-NC_034017.1
                          UA--AUUUGCUACUUU--AUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAA-GGA-
                          ----CUUUGGUUGUUUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAA-GAAA
                          UAGACGUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAA-GGAA
                                                                                                        55
                                                                                                        56
56
                          A---UUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAA-CCAGAUGGAGCAAAACGGAA
C---UAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGA-AAGUAUGGAGCAAUCGAGGA
cISFV-NC 027817.1
                          ----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUG-AGUGGAUUAGGA
                              CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
                          -UUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGAUUUUUCCACCCCGUUCUGGGGAANNN 113
                          GGUAAGAAGAAGAGGGCCCGGGCUCCUCCGGAUGGUUGUUACCA--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
                          -GUCGGGAGUAGUGGAGAGAGGCCCGGUACGCCCUCCGACCAACCGUGGAGGAANNN 113
                          -GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAGNNN 113
                          GGAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAAANNN 113
                          GCACUGAAGAAGAAGGAAUUUCCCUGGGAGGAGGAGGAAAACGGGGCUUUGAG--ANNN
                          AG---UUUUCCCGACCGGGAGGUGGUCGGCCCGGCCCAGAGAGGGGCUGGGGAAGANNN 113
AG---UUUUCCCGGCCGGGAGGCGGUCGGGCCGGCCCAAAGAGGGGCUGGCGAAGANNN 113
cISFV-NC_027817.1
                          GGCCUUCUUCCCCUCCGAGGGAAGAAAAGAAAGCCCCUGUGAUACAAUCAC--AGGNNN 113
                          0.......70.......80........90........100.......110.......1
CISFV-NC_001564.2

CISFV-NC_005064.1

CISFV-NC_012932.1

CISFV-NC_008604.2

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_034204.1
                          NNNNCUUUCAAAACUGGACGGGCAACAGGGAGAAAUCCC---GGGG--UAGCGAACCUC 168
                         NNNN--UUCAAAACCGGACGGACAACAGGGA-AUUUCCC---GGGGA--CGCGUACCCC 165
NNNN--A--AAAAUCGGGCGGAGCAACAGGGAGAAAUCCC---GGGGAAUUGCGAACCCC 166
NNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCC---AGGUGACUGACGCCACC 168
NNNNUCACCGUAAGGUGAGGGCGCACAGGAUAGAAUAUCC CGCUGACGCACCACC 168
                          NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCC---GGGUGAGUGACGCCACC 170
                          NNNN-CACCGCAAGGUGAGGCGCACAGGAUCGAGUAUCC---GGGUGAGUGACGCCACC
                                                                                                       169
                          NNNN--ACCGCAAGGUGAGGGCGCACAGGACGAAAGUCCA---GGGUGAGUGACGACACC
                                                                                                        168
                          NNNNAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCC---GGGUGAUUAGCCACACC
                                                                                                        170
                          NNNN-----CUCCAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCAC
                                                                                                        164
cISFV-NC_034017.1
cISFV-NC_027817.1
                          NNNN-----ACCUCCAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUGGCGACCAC
                                                                                                        165
                          NNNN------AUGUGCAGGGCAACAAGUUCUAACGAACUAGGGUGAGUAG-CGUCACC
                                                                                                       165
                          20......130......140......150......160......170......1
                          CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
                          CUCCGUUAAUGUGAAAAAGUAU-G--GGGAAAGAACUCAUCUUAACCUCCCACCGUUAGG 225
                                                                                                        \frac{1}{2} \frac{1}{2} \frac{1}{2}
                          CUCCGU-AAUGUGAAAAAGUAUGG--GGAAAAGAACCCAGCUAAAUCUCCCACCGAUAGG
                          CUCCGA-AAUGUGAAAAAUUAUGG--GGAAAAGUACCCAUCUAAGCCUCCCACCGUUAGG
                                                                                                        223
                          UCCCGA-AAUGUGUAUAGUAACAG--AGCAUGCCUGCAGCAGCAGGUCUCCACCGUUAGG
                                                                                                       225
CISFV-NC_021069.1
CISFV-NC_021069.1
CISFV-NC_024299.2
CISFV-NC_024299.2
                          CCCCGA-AAUGUGUAUUGUGACAG-GCU-UGCUCGCAGCAGCAAAUCUCCACCGUUAGG
                                                                                                        226
                          CCCCGA-AAUGUGUAUUGUGACAG-GCU-UGCUCGCAGCAGCAAAUCUCCACCGUUAGG
                                                                                                        225
                          CCCCGA-AAUGUGUAUAUAGCCCA--UGUUGAGUCUAUACAACAGGCUUCCACCGGUAGG
                                                                                                        225
                          CCCCGA-AACGUGAUUUAUAUGAU-GACAAGAAUCAGAUGGAAGAUCCCCACCGUUAGG
                                                                                                        227
cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_027817.1
                          CCUCUGAAAUGUUAAUAAAACUGUGCGGAA--AAAUCCAAAGACUGUCCCCACCGAUAGG
                                                                                                        222
                          CCUCUGUAAUGUUUAAUAAGUUGUGCGGAA--AAAUCCAAAGAAUGUCCCCACCGAUAGG
                                                                                                        223
                          CCCCGGUUGUGAAAACGAUUGCGACUAGAACUAAAGUCG-AGAGUCUCCCCACCGAUAGG
                          CISFV-NC_001564.2

CISFV-NC_005064.1

CISFV-NC_012932.1

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_034204.1

CISFV-NC_034017.1

CISFV-NC_027817.1
                          GAG--UUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU 258
GAG--UUUGGAAHACCHIIIIIICUAHACCAUAAAUGCGCU 258
                          GAG--UUUGGAAUACCUUUUCUAUACCAUAGAUGCGC- 258
                          AGAC-UUGUUG-CGGGCAAGCUCU--UGUUCACGU-CU 258
                          AGAU-UUGUUG-UGAGCAUUGC-U--UGUCUACGU-CU 258
                          AGAU-UUGUUG-UGAGCAUUGCCU--UGUCUACGU-CU 258
                          AAGCGUUGUUG-UGAUGA-CACAU--GAGCAACGU-CU
                          GGAUCCAUCCAGCAUUCUUGG-AA--UCAU---GG-CU
                          GGAC--UUUGGGAAAGUUUUUCAACACACGUAGCGUCU
                                                                           258
                                                                            258
                          GGGC--UUUGGGAAAGUUUUUCAACACACGUAGCG-CU
                          GGAG--CUCGUCAGUUUGUUUUAGUCCA-AUUACG-CU
                          40......250......260......270.....
```