```
CISFV-NC_008604.2
CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
                          AAGGCUUGAGUUGUUUAUAAUAGUCGUUUUUCUCGCAGAA--AUGGGAAAGGACGACGGU
                          AA--UUUCGCUACCUUAUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GU
                                                                                                       56
                          UA--AUUUGCUACUUUAUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGGA--GU
                          UAGACGUUUGUCACGGAAUAGCUUGAGAAGAAGCUUACU--UAUGUCAAGUAAGGAAGCA
                                                                                                       58
CISFV-NC_027019.1
CISFV-NC_001564.1
CISFV-NC_012932.1
CISFV-NC_034204.1
CISFV-NC_034017.1
                          56
                          AA--AUCAAAGGAGUUGUUCGGAAAAGUGACCUUGGUUCGUUAUGAAUAGAGGAAG--CU
                          AA--AAUCUAGAAGUUGUUUGGAAAAGAGAUAUCUGGUUCGUAUGAAUAGGAUGCU--CU
                                                                                                       56
                                                                                                       56
56
                         AU-UUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAA-CCAGAUG-GAGCAAAACG-GAA
CU-AGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGA-AAGUAUG-GAGCAAUCGA-GGA
cISFV-NC 027817.1
                          -U-UGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGC
                           cISFV-NC_008604.2
cISFV-NC_021069.1
cISFV-NC_012671.1
cISFV-NC_024299.2
                          AAGAAGAAGAAGGGCCCGGCUC-CUCCGGAUGGUUGUUACCACCGGGG-CGCGNNNNNN 116
                          CGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAANNNNNN 116
                          CUGGAGUGGUGGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAGNNNNNN
                          GUAAAUUAUGGGAGAAGGGAAGCUAUCC-CGCCC-CCUCCCAUUAAGGAGAAAANNNNN 116
CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_001564.2

CISFV-NC_005064.1

CISFV-NC_012932.1

CISFV-NC_034204.1

CISFV-NC_034017.1
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                          UGGAGGCGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAANNNNN 116
                          UGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNNNNN
                         AGUUUUCCCGACCGGGAGGUGGUCGGCCCGGCCCAGAGAGGGCUGGGGAAGANNNNNN AGUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGGCGAAGANNNNNN
                                                                                                       116
                                                                                                       116
cISFV-NC_027817.1
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                          0.......70.......80........90.......100......110.......1
CISFV-NC_008604.2

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_001564.2

CISFV-NC_005064.1

CISFV-NC_012932.1

CISFV-NC_034204.1
                         NUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCCAGGUGACU-GACGCCACCUCCCG 173
                         175
                                                                                                       174
                         N--ACCGCAAGGUGAGGGCGCACAGGACGAAAGUCCAGGGUGAGU-GACGACCCCCCCG 173
                         NAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCCGGGUGAUU-AGCCACACCCCCCG
                         NCUUUCAAAACUGGACGGGCAACAGGGAGAAAUCCCGGGG--UA-GCGAACCUCCUCCG
                                                                                                       173
                         N--UUCAAAACCGGACGGAGCAACAGGGA-AUUUCCCGGGGA--C-GCGUACCCCUCCG
N--A-AAAUCGGGCGGAGCAACAGGGAGAAAUCCCGGGGAAUU-GCGAACCCCCUCCG
                                                                                                       170
                                                                                                       169
                         N-----CUCCAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCACCCUCU
cISFV-NC_034017.1
cISFV-NC_027817.1
                         N----ACCUCCAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUGGCGACCACCCUCU
                                                                                                       170
                         N----AUGUGCAGGGCAACAAAGUUCUAACGAACUAGGGUGAGUAG-CGUCACCCCCG
                          20......130......140......150......160......170......1
                          .....))))...))))
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CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
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                          -AAAUGUGUAUUGUGACA-G-GCUUGCUCGCAGCAGCAAAUCUCCACCGUUAGG-AGAUU 230
                          -AAAUGUGUAUAUAGCCC-AUGUUGAGUCUAUACAACAGGCUUCCACCGGUAGGAAGCGU
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CISFV-NC_024299.1
CISFV-NC_027819.1
CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
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                                                                                                       232
                          UUAAUGUGAAAAAGUAUG-GGG-AAAGAACUCA<mark>U</mark>CUUAACCUCCCACCGUUAGGGAGUUU
                                                                                                       231
                          -UAAUGUGAAAAAGUAUG-GGGAAAAGAACCCAGCUAAAUCUCCCACCGAUAGGGAGUUU
-AAAUGUGAAAAAUUAUG-GGGAAAAGUACCCAUCUAAGCCUCCCACCGUUAGGGAGUUU
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cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_027817.1
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                                                                                                       228
                                                                                                       229
                          GUUGUGAAAACGAUUGCGACUAGAACUAAAGUCGAGAGUCUCCCCACCGAUAGGGGAGCU
                                                                                                       230
                          80......190......200.......210.......220.......230.......2
cISFV-NC_008604.2
cISFV-NC_021069.1
cISFV-NC_012671.1
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UG-UUGUGAGCAUUG-CUUGUCUACGU-CU 258
UG-UUGUGAGCAUUGCCUUGUCUACGU-CU 258
CISFV-NC_U1Z6/1.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_034204.1
                          UG-UUGUGAUGACACAUGAG-CAACGU-CU
                          CA-U--CCAGCAUUCUUGGAAUCAUGG-CU
                                                                 258
                          UGAUUAUCUUUUCUAUACCAUAGAUGC---
                                                                 258
                         GGACUAUCUUUUCUAUACCAUAAAUGCGCU
GGAAUACCUUUUCUAUACCAUAGAUGCGC-
                                                                 258
                                                                 258
                          UGGGAAAGUUUUUCAACACACGUAGCGUCU
                                                                 258
cISFV-NC_034017.1
cISFV-NC_027817.1
                          UGGGAAAGUUUUUCAACACACGUAGCG-CU
                                                                 258
                          CGUCAGUUUGUUUUAGUCCAAUUA-CG-CU
                          40......250......260.....
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