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CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
CISFV-NC_021069.1
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                              AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA-
                              AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC---
                              AAGGCUUGAGUUGUUU - AUAAUAGUCGUUUUUCUCGCAGAA - AUGGGAAAGGACGACG
AA - UUUCGCUACCUU - AUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGAA-
cISFV-NC_012671.1
                              UA--AUUUGCUACUUU--AUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGA--
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_034204.1
CISFV-NC_034017.1
                                                                                                                         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
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                              UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNN
                              GUAAGAAGAAGAGGGCCGGGCUCCUCCGGAUGGU-UGUUACCA--CCGGGGCGCGNNN 113
CISFV-NC_021069.1
CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_034204.1
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                              GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCC-UCCUGCCAAUCGUGGAGGAGNNN 113
                              GAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCC-CCCUCCCAUUAAGGAGAAAANNN 113
                              CACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGGCUUUGAG--ANNN
                              AAAGUUUUCCCGACCGGGAGGUGGUCGGCCCAGAGA-GGGGCUGGGGAAGANNN 113
GAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGCCCAAAGA-GGGGCUGGCGAAGANNN 113
cISFV-NC_027817.1
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CISFV-NC_001564.2
CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
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                              NNNN--UUCAAAACCGGACGGAGCAACAGGGA-AUUUCCCGGGGA--C-GCGUACCCCCU
                                                                                                                         167
                              NNNN--A--AAAAUCGGGCGGAGCAACAGGGAGAAAUCCCGGGGAAUU-GCGAACCCCCU
                                                                                                                         168
                              NNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCCAGGUGACU-GACGCCACCUC
CISFV-NC_021069.1
CISFV-NC_012671.1
                              NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCC
                                                                                                                         172
                              NNNN-CACCGCAAGGUGAGGCGCACAGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCC
                                                                                                                         171
cISFV-NC_024299.2
cISFV-NC_027819.1
cISFV-NC_034204.1
                              NNNN--ACCGCAAGGUGAGGCGCACAGGACGAAAGUCCAGGGUGAGU-GACGACACCCC
                                                                                                                         170
                              NNNNAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCCGGGUGAUU-AGCCACACCCC
                                                                                                                         172
                              NNNN--CUC----CAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCACCC
                                                                                                                         166
cISFV-NC_034017.1
cISFV-NC_027817.1
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                                                                                                                         167
                              NNNNAUGUG----CAGGGCAACAAAGUUCUAACGAACUAGGGUGAGUAG-CGUCACCCC
                                                                                                                         167
                              20......130......140......150......160......170......1
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CISFV-NC_005064.1
CISFV-NC_012932.1
CISFV-NC_008604.2
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                                                                                                                         228
                                                                                                                         225
                                                                                                                   GAG
                                                                                                                         226
                              CCG-AAAUGUGUAUAGUAA<mark>CA</mark>GA-GCA<mark>UG</mark>CCUGCAGCAGC<mark>A</mark>GG<mark>UCU</mark>
                                                                                                     ACCGUUA
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cISFV-NC_021069.1
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cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
                              CCG-AAAUGUGUAUUGUGAC<mark>A</mark>GG-CU-<mark>UG</mark>CUCGCAGCAGCAAAUCU
CCG-AAAUGUGUAUAUAGCCCAU-GUUGAGUCUAUACAAC<mark>A</mark>GGCUU
CCG-AAACGUGAUUUAUAUG<mark>A</mark>UGACAA<mark>GA</mark>AUCAGA-UGGAAGA<mark>UCC</mark>
                                                                                                                  AGA
                                                                                                                         228
                                                                                                     ACCGUUA
                                                                                                      ACCGGUA
                                                                                                                         228
                                                                                                                         230
                                                                                                      ACCGUUA
cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_027817.1
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UCUGUAAUGUUUAAUAAGU<mark>UG</mark>UGCGGAAA-AAUCCAAAGA<mark>A</mark>UG<mark>UCC</mark>
                                                                                                     ACCGAUA
                                                                                                                         225
                                                                                                                         226
                                                                                                      ACCGAUA
                              CCGGUUGUGAAAACGAUUGC<mark>G</mark>ACUAGA<mark>AC</mark>UAAAGUCGAGA<mark>G</mark>UC<mark>UCC</mark>
                                                                                                     ACCGAUA
                                                                                                                         227
                              80......190......200.......210.......220.......230.......2
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--UUUGG-ACUAUCUUUUCUAUACCAUAAAUGCGCU 258
cISFV-NC_001564.2
cISFV-NC_005064.1
cISFV-NC_012932.1
                              --U<mark>U</mark>UGG-AAUAC<mark>CU</mark>UUUCUAUAC<mark>CA</mark>UAGAUGCGC-
cISFV-NC_008604.2
cISFV-NC_021069.1
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U-UUGUU-GUGAGCAUUGC----UUGUCUACGU-CU
                                                                                    258
cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
                              U-U<mark>U</mark>GUU-GUGAG<mark>CA</mark>UUGCC---U<mark>UG</mark>UCUACGU-CU
                                                                                     258
                              CGU<mark>U</mark>GUU-GUGAUGACACAU---GAG-CAACGU-CU
UCCAUCC-AGCAU<mark>UC</mark>UUGGA---A<mark>UC</mark>AU---GG-CU
cISFV-NC_034204.1
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                                                                                     258
cISFV-NC_034017.1
cISFV-NC_027817.1
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                                                                                     258
                              GCU<mark>C</mark>GUC-AGUUU<mark>GU</mark>UUUAG---U<mark>C</mark>CAAUUACG-CU
                              40......250......260......270....
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