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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---
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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
                               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
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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
                               NNNNUCACCGCAAGGUGAGGCGCACAGGAUAGA--AUCCAGGUGACU-GACGCCACCUC
CISFV-NC_021069.1
CISFV-NC_012671.1
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                                                                                                                              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
                               NNNNACCUC----CAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUGGCGACCACCC
cISFV-NC_034017.1
                                                                                                                              167
cISFV-NC 027817.1
                               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
                                                                                                                              226
                                                                                                                        GAG
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CCG-AAAUGUGUAUUGUGACAGG-CU-UGCUCGCAGCAGCAAAUCU
                                                                                                         ACCGUUA
                                                                                                                              228
cISFV-NC_021069.1
                                                                                                         ACCGUUA
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cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
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CCG-AAAUGUGUAUAUAGCCCAU-GUUGAGUCUAUACAA<mark>C</mark>AGGCUU
CCG-AAACGUGAUUUAUAU<mark>GA</mark>UGACAAGAAUCAGA-UG<mark>GA</mark>AGA<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>UGC GAAA-AAUCCAAAG<mark>AA</mark>UG<mark>UCC</mark>
                                                                                                         ACCGAUA
                                                                                                                              225
                                                                                                                              226
                                                                                                          ACCGAUA
                               CCGGUUGUGAAAACGAUUGC<mark>G</mark>ACU<mark>A</mark>GA<mark>A</mark>CUAAAGUCGA<mark>GAG</mark>UC<mark>UCC</mark>
                                                                                                         ACCGAUA
                                                                                                                              227
                               80.....190.....200......210......220......230......2
                               ).))))))...))))
                               --UUUUGAUUAUCUUUUCUAUACCAUAGAUGC--
--UUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU
--UUUGGAAUACCUUUUCUAUACCAUAGAUGCGC-
C-UUGUUGCGGGCAAGCUC---UUGUUCACGU-CU
U-UUGUUGUGAGCAUUGC---UUGUCUACGU-CU
U-UUGUUGUGAGCAUUGCC---UUGUCUACGU-CU
CGUUGUUGUGAUGACACAU--GAG-CAACGU-CU
UCCAUCCAGCAUUCUUGA--AUCAU---GG-CU
cISFV-NC_001564.2
cISFV-NC_005064.1
cISFV-NC_012932.1
                                                                                       258
258
                                                                                       258
cISFV-NC_008604.2
cISFV-NC_021069.1
                                                                                       258
cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
                                                                                       258
                                                                                       258
258
cISFV-NC_034204.1
                               C--U<mark>UU</mark>GGGAAAG<mark>U</mark>UU<mark>U</mark>UCAACA<mark>CA</mark>CGUAGCGUCU
                                                                                       258
cISFV-NC_034017.1
cISFV-NC_027817.1
                                                                                       258
                               C--UUUGGGAAAGUUUUUCAACACACACGUAGCG-CU
                               G--CUCGUC-AGUUUGUUUUAGUCCAAUUACG-CU
                                                                                       258
                               40......250......260......270...
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