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CISFV-NC_034204.1
CISFV-NC_034017.1
CISFV-NC_008604.2
CISFV-NC_021069.1
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                                          -AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAA--CGG
                                        --CUAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUC--GAG
                                    AAGGCUUGAGUUGUUU - AUAAUAGUCGUUUUUCUCGCAGAA - AUGGGAAAGGACGACGAA - UUUCGCUACCUU - AUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGAA-
                                                                                                                                                  56
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_001564.2
CISFV-NC_005064.1
                                    UA--AUUUGCUACUUU--AUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGA--
                                                                                                                                                  56
                                    UAGACGUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAG
                                    AAUUUUUGCCGAGAGACCGUGA---GGUUGAACCCGGCAAGGAAUGAAGAGGAAGG---AAAUCAAAGGAGUUGUUCGGAA---AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA---
cISFV-NC 012932.1
                                    AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC---
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cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_008604.2
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                                    AAAGUUUUCCCGACCGGGAGG<mark>U</mark>GGUCGGGCCCGGCCCAGAGA-GGGGCUGGGGAAGANNN 113
                                    GAAGUUUUCCCGGCCGGAGGCGGUCGGGCGGCCCCAAAGA-GGGGCUGGCGAAGANNN 113
                                    GUAAGAAGAAGAAGGGCCGGCCUCCUCCGGAUGGU-UGUUACCA--CCGGGGCCGCNNN 113
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_015064.1
                                    GUCGGGAGUAGUGGAGAGAGACGCGGUACGCCCC-UCCGACCAACCGUGGAGGAANNN 113
GUCUGGAGUGGUGGAGAGAGACGUGGAACGCGUCC-UCCUGCCAAUCGUGGAGGAGNNN 113
GAGUAAAUUAUGGGAGAAGGGAAGC-UAUCCCGCC-CCCUCCCAUUAAGGAGAAAANNN 113
                                    CACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGGCUUUGAG--ANNN
                                    AUUUGGAGGCGAGGAAAAGCGCCAGGGAGGAUUUUUCCACCCCGUUCUGGGGAANNN
                                    cISFV-NC_012932.1
                                    UCUUGAAGAAAGGGCCUUCGG<mark>G</mark>CCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNN
                                    0.......70.......80........90........100.......110.......1
                                     .....((((....)))).(((((.....)))).
                                   NNNNAUGUG----CAGGGCAACAAAGUUCUAACGAACUAG GUGAGUAG-CGUCACCC
NNNN--CUC----CAGAGCAAACAAGGGAUUUAUCCUUAA GUGAUUUGACGACCACCC
NNNNACCUC----CAGAGCAAACGAGGGAGAAAUCCCU-AAGUGAGUUGGCGACCACCC
NNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCCAGGUGACU-GACGCCACCC
NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCC
NNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCC
NNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCC
CISFV-NC_027817.1
CISFV-NC_034204.1
CISFV-NC_034017.1
CISFV-NC_008604.2
                                                                                                                                                 167
                                                                                                                                                  166
                                                                                                                                                 167
                                                                                                                                           CUC 170
CISFV-NC_008604.2

CISFV-NC_021069.1

CISFV-NC_012671.1

CISFV-NC_024299.2

CISFV-NC_027819.1

CISFV-NC_001564.2
                                                                                                                                                172
                                                                                                                                                 171
                                   NNNN--ACCGCAAGGUGAGGGCGCACAGGACGCAGGGUGAGU-GACGACAC CC
NNNNAAAGGCCAUCCACGCAGGGCAACAGGAAAAUUCCGGGUGAUU-AGCCACAC CC
NNNNCUUUCAAAACUGGACGGGGCAACAGGGAGAAAUCCCGGGG--UA-GCGAACCU CU
                                                                                                                                                  170
                                                                                                                                                  170
cISFV-NC_005064.1
cISFV-NC_012932.1
                                    NNNN--UUCAAAACCGGACGGAGCAACA<mark>G</mark>GGA-AUUUC<mark>C</mark>CG<mark>GGG</mark>A--C-GCGUAC<mark>CCC</mark>CU
                                                                                                                                                  167
                                    NNNN--A--AAAAUCGGGCGGAGCAACA<mark>G</mark>GGAGAAAUC<mark>C</mark>CG<mark>GGG</mark>AAUU-GCGAAC<mark>CCCC</mark>U
                                                                                                                                                 168
                                    20......130......140......150......160......170......1
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CISFV-NC_034204.1
CISFV-NC_034017.1
CISFV-NC_008604.2
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UCUGUAAUGUUUAAUAAGUU -GUGCGGAA -AAAUCCAAAGAAUGUCCCCACCGAUA
CCG -AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCAGCAGG - UCUCCACCGUUA
CCG -AAAUGUGUAUUGUGACAGGCU - UGCUCGCAGCAGCAAA - UCUCCACCGUUA
                                                                                                                                                  226
                                                                                                                                                  224
                                                                                                                                                  225
227
CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
                                                                                                                                                  228
                                    CCG-AAAUGUGUAUUGUGA<mark>C</mark>AGGCU-U<mark>GC</mark>UCGCAGCAGCA<mark>A</mark>A--<mark>UCU</mark>
                                                                                                                                           GAG
                                                                                                                                                  227
                                                                                                                            ACCGUUA
                                    CCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAACAGG--
CCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAUGGAAGA--
                                                                                                                                                  227
                                                                                                                            ACCGGUA
                                                                                                                                                  229
                                                                                                                            ACCGUUA
                                                                                                                                            GG
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cISFV-NC_005064.1
cISFV-NC_012932.1
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CCG-UAAUGUGAAAAAGUAUGGGG--AAAAGAACCCAGCUAAAU<mark>CUC</mark>
                                                                                                                                           <mark>G</mark>GA
                                                                                                                            ACCGUUA
                                                                                                                                                  227
                                                                                                                                                  224
                                                                                                                            ACCGAUA
                                    CCG-AAAUGUGAAAAAUUA<mark>U</mark>GGGG--A<mark>AA</mark>AGUACCCAUCUAAGC<mark>CUC</mark>
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cISFV-NC_034017.1
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                                                                                                   258
258
                                     CUUUGGGAAAGU<mark>UU</mark>UUCAACACACGUAGCGUCU
CUUUGGGAAAGU<mark>UU</mark>UUCAACAC<mark>A</mark>CGUAGCG-CU
                                                                                                   258
CISFV-NC_008604.2
CISFV-NC_021069.1
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AU-<mark>U</mark>UGUU-GUGA<mark>GC</mark>AUUGCU-U<mark>G</mark>UCUACGU-CU</mark>
                                                                                                  258
                                                                                                   258
cISFV-NC_012671.1
cISFV-NC_024299.2
cISFV-NC_027819.1
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                                                                                                   258
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                                                                                                   258
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                                     UUUUGAUUAUCU<mark>UU</mark>UCUAUACC<mark>A</mark>UAGAUGC---
                                                                                                   258
cISFV-NC_005064.1
cISFV-NC_012932.1
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                                     UU<mark>U</mark>GGAAUACCU<mark>UU</mark>UCUAUACC<mark>A</mark>UAGAUGCGC-
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