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CISFV-NC_034204.1
CISFV-NC_034017.1
CISFV-NC_008604.2
CISFV-NC_021069.1
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                                         --CUAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUC--GAG
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                                                                                                                                                        56
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_001564.2
CISFV-NC_005064.1
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                                      UAGACGUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAG
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                                     AAUUUUUGCCGAGAGACCGUGA---GGUUGAACCCGGCAAGGAAUGAAGAGGAAGG---AAAUCAAAGGAGUUGUUCGGAA---AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA---
                                                                                                                                                        53
53
cISFV-NC 012932.1
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cISFV-NC_034204.1
cISFV-NC_034017.1
cISFV-NC_008604.2
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                                     AAAGUUUUCCCGACCGGAGGUGGUCGGGCCCGGCCCAGAGA-GGGGCUGGGGAAGANNN 113
GAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGA-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
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GUCUGGAGUGGUGGAGAGAGACGUGGAACGCGUCC-UCCUGCCAAUCGUGGAGGAGNNN 113
GAGUAAAUUAUGGGAGAAGGGAAGC-UAUCCCGCC-CCCUCCCAUUAAGGAGAAAANNN 113
                                      CACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGGCUUUGAG--ANNN
                                      AUUUGGAGGCGAGGAAAAGCGCCAGGGAGGAUUUUUCCACCCCGUUCUGGGGAANNN
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                                      0.......70.......80........90........100.......110.......1
                                      .....((((....)))).(((((.....)))).
CISFV-NC_027817.1
CISFV-NC_034204.1
CISFV-NC_034017.1
CISFV-NC_008604.2
CISFV-NC_021069.1
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_001564.2
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NNNN--CUC----CAGAGCAAACAAGGGAUUUAUCCUUAA GUGAUUUGACGACCACCC
NNNNACCUC----CAGAGCAAACGAGGGAGAAAUCCCU-AAGUGAGUUGGCGACCACCC
NNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCCAGGUGACU-GACGCCACCC
NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCC
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NNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCC
                                                                                                                                                       167
                                                                                                                                                        166
                                                                                                                                                       167
                                                                                                                                                 CUC 170
                                                                                                                                                      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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CCG -AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCAGCAGG - UCUCCACCGUUA
CCG -AAAUGUGUAUUGUGACAGGCU -UGCUCGCAGCAGCAAA - UCUCCACCGUUA
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                                                                                                                                                        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
cISFV-NC_001564.2
cISFV-NC_005064.1
cISFV-NC_012932.1
                                                                                                                                                 GGA
GGA
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CCG-UAAUGUGAAAAAGUAUGGGG--AAAAGAACCCAGCUAAAU<mark>CUC</mark>
                                                                                                                                 ACCGUUA
                                                                                                                                                        227
                                                                                                                                                        224
                                                                                                                                 ACCGAUA
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                                                                                                                                 ACCGUUA
                                                                                                                                                        225
                                      80.....190.....200......210......220......230......2
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cISFV-NC_027817.1
cISFV-NC_034204.1
cISFV-NC_034017.1
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                                                                                                       258
258
                                      ACUUUGGGAAAGU<mark>UU</mark>UUCAACAC<mark>A</mark>CGUAGCGUCU
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
CISFV-NC_012671.1
CISFV-NC_024299.2
CISFV-NC_027819.1
CISFV-NC_001564.2
                                       AU-<mark>U</mark>UGUU-GUGA<mark>GC</mark>AUUGCCUU<mark>G</mark>UCUACGU-CU
                                                                                                       258
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UC<mark>C</mark>AUCCAGCAU<mark>UC</mark>UUGGA-AU<mark>C</mark>AU---GG-CU
                                                                                                       258
                                       UUUUGAUUAUCU<mark>UU</mark>UCUAUACC<mark>A</mark>UAGAUGC---
                                                                                                       258
cISFV-NC_005064.1
cISFV-NC_012932.1
                                                                                                       258
                                        UU<mark>U</mark>GGACUAUCU<mark>UU</mark>UCUAUACC<mark>A</mark>UAAAUGCGCU
                                       UU<mark>U</mark>GGAAUACCU<mark>UU</mark>UCUAUACC<mark>A</mark>UAGAUGCGC-
                                      40......250......260.......270...
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