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                    .........10........20........30........40........50.......6
cISFV-NC_008604.2   AAGGCUUGAGUUGUUUAUAAUAGUCGUUUUUCUCGCAGAA--AUGGGAAAGGACGACGGU 58
cISFV-NC_021069.1   AA--UUUCGCUACCUUAUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GU 56
cISFV-NC_012671.1   UA--AUUUGCUACUUUAUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGGA--GU 56
cISFV-NC_001564.2   AA--UUUUUGCCGAGAGACCGUGAGGUUGAACCCGGCAAGGAAUGAAGAGGAAGGAUUUG 58
cISFV-NC_005064.1   AA--AUCAAAGGAGUUGUUCGGAAAAGUGACCUUGGUUCGUUAUGAAUAGAGGAAGCUUG 58
cISFV-NC_012932.1   AA--AAUCUAGAAGUUGUUUUGGAAAAGAGAUAUCUGGUUCGUAUGAAUAGGAUGCUCUUG 58
cISFV-NC_027819.1   UAGACGUUUGUCACGGAAUAGCUU--GAGAAGAAGCUUACUUUAUGUCAAGUAAGGAAGCA 58
cISFV-NC_034204.1   AUUUUCUUUAGCGA--CAGAAGGAACCCAAAAAGGGAACCAGAUGGAGC-----AAAACG 53
cISFV-NC_034017.1   CUAGUGUUGUUAAA--AAUUUCUUUGGCGACGUACGGAAAGUAUGGAGC-----AAUCGA 53
cISFV-NC_024299.2   CUUUGGUUGUUUGA--AAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAAGGA 58
cISFV-NC_027817.1   UUGAAAGGUUCAAA--CUCUGGUGUCCAACGUUUUGUAUUAAAAUGAGUGGAUU---AGGA 55
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                    .........10........20........30........40........50.......6
cISFV-NC_008604.2   AAGAAGAAGAAGGGGCCGGGCUCCUCCGGAUGGUUGUUA---CCA--CCGGGGCGCGNNN 113
cISFV-NC_021069.1   CGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCCUCCGA---CCAACCGUGGAGGAANNN 113
cISFV-NC_012671.1   CUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCCUCCUG---CCAAUCGUGGAGGAGNNN 113
cISFV-NC_001564.2   GAGGCG--CGAGGAAAAGCGCCAGGGAGGGAUUUUUCCA---CCCCGUUCUGGGGAANNN 113
cISFV-NC_005064.1   AAGAAG---GGGCCUCCGGGCCCAACAUCCAACCGUCCGU---CGGUUGGACCGCCCGNNN 113
cISFV-NC_012932.1   AAGAAA--GGGCCUUCGGGCCAAACAUCUGGUCGGCCAU---CGACCAGACCGCCGGNNN 113
cISFV-NC_027819.1   CUGAAGAAGAAGGGAAUUUCCCUGGGGAGGAGGAGGAAAA---CGGGGCUUUGAG--ANNN 113
cISFV-NC_034204.1   GAAAGUUUUCCCGACCGGGGAGGUGGUCGGGCCCGGCCCAGAGAGGGGGCUGGGGAAGANNN 113
cISFV-NC_034017.1   GGAAGUUUUCCCGCCGGGAGGCGGUCGGGCGGGCCCAAAGAGGGCGGCUGGCGAAGANNN 113
cISFV-NC_024299.2   GUAAAUUAUGGGGAGAAGGGAAGCUAUCCCGCCCCCUCCCAUUAAGGA-----GAAANNN 113
cISFV-NC_027817.1   GGCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCC--UGUGAUACAAUCACAGGNNN 113
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                    0........70........80........90........100.......110.......1
cISFV-NC_008604.2   NNNNCUG-ACGCCACCUCCCGAAAUGUGUAUAGUAACAGAGC-AUGCCUGCAGCAGCAGG 171
cISFV-NC_021069.1   NNNNGUG-ACGCCACCCCCCGAAAUGUGUAUUGUGACAG-GC-UUGCUCGCAGCAGCAAA 170
cISFV-NC_012671.1   NNNNGUG-ACGCCACCCCCCGAAAUGUGUAUUGUGACAG-GC-UUGCUCGCAGCAGCAAA 170
cISFV-NC_001564.2   NNNNAG-CGAACCUCCUCCGUUAAUGUGAAAAGUAUGGGG-AAAGAACUCAUCUUAA-- 169
cISFV-NC_005064.1   NNNN---CGUACCCCCUCCG-UAAUGUGAAAAGUAUGGGGAAAAGAACCCAGCUAAAUC 169
cISFV-NC_012932.1   NNNNUUGCGAACCCCCUCCG-AAAUGUGAAAAUUAUGGGGAAAAGUACCCAUCUA---A 169
cISFV-NC_027819.1   NNNNUUA-GCCCACACCCCCGAAACGUGAUUUAU-AUGAUGACAAGAAUCGAUGGAAGA 171
cISFV-NC_034204.1   NNNNACG---ACCACCCUCUGAAAUGUUAAUAAAA-CUGUGCGGAAAAAUCCAAAGACUG 169
cISFV-NC_034017.1   NNNNGCG---ACCACCCUCUGUAAUGUUUAAUAAG-UUGUGCGGAAAAAUCCAAAGAAUG 169
cISFV-NC_024299.2   NNNNACG---A-CACCCCCCGAAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAACAGGC 169
cISFV-NC_027817.1   NNNNAGU---AGC-GUCACCCCCCGGUUGUGAAAACGAUUGCGACUAGAACUAAAGUCGA 169
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                    20.......130.......140.......150.......160.......170.......1
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                    ........
cISFV-NC_008604.2   UCUCCA-- 177
cISFV-NC_021069.1   UCUCCAC- 177
cISFV-NC_012671.1   UCUCCAC- 177
cISFV-NC_001564.2   CCUCCCAC 177
cISFV-NC_005064.1   UCCCACCG 177
cISFV-NC_012932.1   GCCUCCCA 177
cISFV-NC_027819.1   UCCCCA-- 177
cISFV-NC_034204.1   UCCCCACC 177
cISFV-NC_034017.1   UCCCCACC 177
cISFV-NC_024299.2   UUCCACCG 177
cISFV-NC_027817.1   GAGUCUCC 177
                    80......
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