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                    .........................................................
cISFV-NC_027817.1   ----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAG 56
cISFV-NC_034204.1   ----AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAA--CGG 54
cISFV-NC_034017.1   ----CUAGUGUUGUUAAAAAUUUCUUUUGGCGACGUACGGAAAGUAUGGGACCAAUC--GAG 54
cISFV-NC_008604.2   AAGGCUUGAGUUUGUUU--AUAAUAGUCGUUUUUCUCGCAGAA--AUGGGAAAGGGACGACG 56
cISFV-NC_021069.1   AA--UUUCGCUACCUU--AUUACCUCUCGGUACUAAAGUGGCACUAUGAAGAAGAAGGA-- 54
cISFV-NC_012671.1   UA--AUUUGCUACUUU--AUUACCUCUCGGUACUAAACGUGGCGCUAUGAAGAAGAAGGA-- 54
cISFV-NC_024299.2   ----CUUUGGUUGUUUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAAG 56
cISFV-NC_027819.1   UAGACGUUUGUCACGG----AAUAGCUUAGGAAGAAGCUUACUUUAUGUCAAGUAAGGAAG 56
cISFV-NC_001564.2   AAUUUUUUGCCGAGAGACCGUGA----GGUUGAACCCGGCAAGGAAUGAAGAGGAAGG--- 53
cISFV-NC_005064.1   AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA--- 53
cISFV-NC_012932.1   AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC--- 53
                    .........10........20........30........40........50......6
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                    .....................((((......((((.............))))).........
cISFV-NC_027817.1   GCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCCUGUGA---UACAAUCACAGGNNN 113
cISFV-NC_034204.1   AAAGUUUUCCCGACCGGGAGGUGGUCGGGCCCGGCCCAGAGA-GGGGCUGGGGAAGANNN 113
cISFV-NC_034017.1   GAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGA-GGGGCUGGCGAAGANNN 113
cISFV-NC_008604.2   GUAAGAAGAAGAAGGGGCCGGGCUCCUCCGGAUGGU-UGUUACCA--CCGGGGCGCGNNN 113
cISFV-NC_021069.1   GUCGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCC-UCCGACCAACCGUGGAGGANNN 113
cISFV-NC_012671.1   GUCUGGAGUGGUGGAGAGGACGCGUGGAACGCGUCC-UCCUGCCAAUCGUGGGAGGAGNNN 113
cISFV-NC_024299.2   GAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCC-CCCUCCCAUUAAGGAGAAAANNN 113
cISFV-NC_027819.1   CACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGGCUUUGAG--ANNN 113
cISFV-NC_001564.2   AUUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAANNN 113
cISFV-NC_005064.1   GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCCGNNN 113
cISFV-NC_012932.1   UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNN 113
                    0........70........80........90........100......110.......1
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                    .......................(((.....)))..(((((.........)))))..
cISFV-NC_027817.1   NNNNAUGUG-----CAGGGCAACAAAGUUCUAACGAACUAGGGUGAGUAG-CGUCACCCC 167
cISFV-NC_034204.1   NNNN--CUC-----CAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCACCC 166
cISFV-NC_034017.1   NNNNACCUC-----CAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUUGCGACCACCC 167
cISFV-NC_008604.2   NNNNUCACCGCAAGGUGAGGGCGCACAGGGAUAGA--AUCCAGGUGACU-GACGCCACCUC 170
cISFV-NC_021069.1   NNNNUCACCGUAAGGUGAGGGCGCACAGGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCC 172
cISFV-NC_012671.1   NNNN-CACCGCAAGGUGAGGGCGCACAGGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCC 171
cISFV-NC_024299.2   NNNN--ACCGCAAGGUGAGGGCGCACAGGACGAAAGGACCAGGGUGUAG-GACGACACCCC 170
cISFV-NC_027819.1   NNNNAAAGGCCAUCCACGCAGGGCAACAGGGAAGAAAUUCUCGGGGAUU-AGCCACACCCC 172
cISFV-NC_001564.2   NNNNCUUUCAAAACUGGACGGGGCAACAGGAGAGAAAUCCCGGGG--UA-GCGAACCUCCU 170
cISFV-NC_005064.1   NNNN--UUCAAAACCGGACGGAGCAACAGGA-AUUUCCGGGGA--C-GCGUACCCCCU 167
cISFV-NC_012932.1   NNNN--A--AAAAUCGGGCGGAGCAACAGGGGAGAAAUCCCGGCGGAAUU-GCGAACCCCU 168
                    20.......130.......140.......150.......160.......170.......1
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                    ......................((((....((((((....(((((...(((((.......))))
cISFV-NC_027817.1   CCGGUUGUGAAAACGAUUGC-GACUAGAACUAAAGUCGAGAGUCUCCCACCGAUAGGGG 226
cISFV-NC_034204.1   UCUGAAAUGUUAAUAAAACU-GUGCGGAA-AAAUCCAAAGACUGUCCCCACCGAUAGGGG 224
cISFV-NC_034017.1   UCUGUAAUGUUAAUAAGUU-GUGCGGAA-AAAUCCAAAGAAUGUCCCCACCGAUAGGGG 225
cISFV-NC_008604.2   CCG-AAAUGUGUAUAAGUAACAGAGCAUGCCUGCAGCAGCAGG--UCUCCACCGUUAGGAG 227
cISFV-NC_021069.1   CCG-AAAUGUGUAUUGUGACAGGCU-UGCUCGCAGCAGCAAA--UCUCCACCGUUAGGAG 228
cISFV-NC_012671.1   CCG-AAAUGUGUAUUGUGACAGGCU-UGCUCGCAGCAGCAAA--UCUCCACCGUUAGGAG 227
cISFV-NC_024299.2   CCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAACAGG--CUUCCACCGGUAGGAA 227
cISFV-NC_027819.1   CCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAUGGAAGA-CUCCCACCGUUAGGGG 229
cISFV-NC_001564.2   CCGUUAAUGUGAAAAGUAUGGGG---AAAGAACUCAUCUUAAACUCUCCCACCGUUAGGGA 227
cISFV-NC_005064.1   CCG-UAAUGUGAAAAAGUAUGGGG---AAAAGAACCCGACUAAAUCUCUCCCACCGUUAGGGA 224
cISFV-NC_012932.1   CCG-AAAUGUGAAAAAUUAUGGGG--AAAAGUACCCAUCUAAGCCUCUCCACCGUUAGGGA 225
                    80.......190.......200.......210.......220.......230.......2
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                    )..)))))..)))))))....)))))...)))))....)))
cISFV-NC_027817.1   AGCUCGUCAGUUUGUUUUUAGUCCAAUUA--CGCU 258
cISFV-NC_034204.1   ACUUUGGGAAAGUUUUUCAACACACGUAGCGUCU 258
cISFV-NC_034017.1   GCUUUGGGAAAGUUUUUCAACACACGUAGCG-CU 258
cISFV-NC_008604.2   AC-UUGUU-GCGGGCAAGCUCUUGUUCACGU-CU 258
cISFV-NC_021069.1   AU-UUGUU-GUGAGCCAUUGCU-UGUCUACGU-CU 258
cISFV-NC_012671.1   AU-UUGUU-GUGAGCCAUUGCU-UGUCUACGU-CU 258
cISFV-NC_024299.2   GCGUUGUU-GUGAUGACACAUGAG-CAACGU-CU 258
cISFV-NC_027819.1   AUCCAUCCAGCAUUCUUGGA-AUCAU---GG-CU 258
cISFV-NC_001564.2   GUUUUGAUUAUCUUUUCUAUACCAUAGAUGC--- 258
cISFV-NC_005064.1   GUUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU 258
cISFV-NC_012932.1   GUUUGGAAUAUCCUUUUCUAUACCAUAGAUGCGC- 258
                    40.......250.......260.......270..
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