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                    .........|.........|.........|.........|.........|.........|
cISFV-NC_001564.2   GAGAGACCGUGAGGUUGAACCCGGCAAGGAAUGAAGAGGAAGG--AUUUGGAGGCGCGAG 58
cISFV-NC_005064.1   AGUUGUUCGGAAAAGUGACCUUGGUUCGUUAUGAAUAGAGGAA--GCUUGAAGAAGGGGC 58
cISFV-NC_012932.1   AGUUGUUUGGAAAAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC--UCUUGAAGAAAGGGC 58
cISFV-NC_034204.1   GCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGGAGCAAAACG--GAAAGUUUUCCCGAC 58
cISFV-NC_034017.1   UAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUCGA--GGAAGUUUUCCCGGC 58
cISFV-NC_008604.2   UUGUUUAUAAUAGUCGUUUUUCUCGCAGAAAUGGGAAAGGACGACGGUAAGAAGAAGAAG 60
cISFV-NC_021069.1   CCUUAUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GUCGGGAGUAGUGG 58
cISFV-NC_012671.1   CUUUAUUACCUCUGGUACUAAACGUGGCGUCAUGAAGAAGAAGGA--GUCUGGAGUGGUGG 58
cISFV-NC_024299.2   UUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAAGGAGUAAAUUAUGGG 60
cISFV-NC_027819.1   UCACGGAAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAGCACUGAAGAAGAAG 60
cISFV-NC_027817.1   CAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGCCUUCUUCCCCUC 60
                    .........10........20........30........40........50........6
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                    .........|.........|.........|.........|.........|.........|
cISFV-NC_001564.2   GAAAAGCGCCAGGGGAGGGAUUUUUCCACCCCGUUCUGGGGGAANNNNNNN--AGCGAACCU 116
cISFV-NC_005064.1   CUCCGGGGCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCCGNNNNNNN----CGUACCC 114
cISFV-NC_012932.1   CUUCGGGGCCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNNNNNN-UUGCGAACCC 117
cISFV-NC_034204.1   CGGGAGGGUGGUCGGGCCCGGCCCAGAGAGGGGGCUGGGGAAGANNNNNNN---ACGACCAC 115
cISFV-NC_034017.1   CGGGAGGCGGUCGGGCGGGCCCAAAGAGGGGCUGGCGAAGANNNNNNN--GCGACCAC 115
cISFV-NC_008604.2   GGGCCGGGCUCCUCCGGAUGGUUGUUACCA--CCGGGGCGCGNNNNNNN-CUGACGCCAC 117
cISFV-NC_021069.1   AGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAANNNNNNN-GUGACGCCAC 117
cISFV-NC_012671.1   AGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAGNNNNNNN-GUGACGCCAC 117
cISFV-NC_024299.2   AGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAAANNNNNNN----ACGACAC 114
cISFV-NC_027819.1   GGAAUUUCCCUGGGAGGGAGGAGGAAAACGGGGCUUUGAG--ANNNNNNN-UUAGCCACAC 117
cISFV-NC_027817.1   CGAGGGAAGAA--AAAGAAAGCCCCUGUGAUACAAUCACAGGNNNNNNNNAGUAGCGUCAC 118
                    0........70........80........90........100.......110.......1
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                    .........|.........|.........|.........|.........|.........|
cISFV-NC_001564.2   CCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAACUCAUCUUAACCUCCCAC-- 167
cISFV-NC_005064.1   CCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCCAGCUAAAUCUCCCACCG 167
cISFV-NC_012932.1   CCUCCG-AAAUGUGAAAAAUUAUGGGGAAAAGUACCCAUCUAAGCCUCCCA--- 167
cISFV-NC_034204.1   CCUCUG-AAAUGUUAAUAAACUGUGCGGAAAAAUCCAAAGACUGUCCCCACC- 167
cISFV-NC_034017.1   CCUCUG-UAAUGUUUAAUAAGUUGUGCGGAAAAAUCCAAAGAAUGUCCCCACC- 167
cISFV-NC_008604.2   CUCCCG-AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCAGCAGGUCUCCA--- 167
cISFV-NC_021069.1   CCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAAAUCUCCAC-- 167
cISFV-NC_012671.1   CCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAAAUCUCCAC-- 167
cISFV-NC_024299.2   CCCCCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAACAGGCUUCCACCG 167
cISFV-NC_027819.1   CCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAUGGAAGAUCCCCA--- 167
cISFV-NC_027817.1   CCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAAAGUCGAGAGUCUC---C 167
                    20.......130.......140.......150.......160.......170..
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