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                   ............................................................
cISFV-NC_001564.2  NGAGAGACCGUGAGGUUGAACCCGGCAAGGAAUGAAGAGGAAGG--AUUUGGAGGCGCGA 58
cISFV-NC_005064.1  NAGUUGUUCGGAAAAGUGACCUUGGUUCGUUAUGAAUAGAGGAA--GCUUGAAGAAGGGG 58
cISFV-NC_012932.1  NAGUUGUUUGGAAAAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC--UCUUGAAGAAAGGG 58
cISFV-NC_034204.1  NGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAACG--GAAAGUUUUCCCGA 58
cISFV-NC_034017.1  NUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUCGA--GGAAGUUUUCCCGG 58
cISFV-NC_008604.2  NUUGUUUAUAAUAGUCGUUUUUCUCGCAGAAAUGGGAAAGGACGACGGUAAGAAGAAGAA 60
cISFV-NC_021069.1  NCCUUAUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GUCGGGAGUAGUG 58
cISFV-NC_012671.1  NCUUUAUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGGA--GUCUGGAGUGGUG 58
cISFV-NC_024299.2  NUUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAAGGAGUAAAUUAUGG 60
cISFV-NC_027819.1  NUCACGGAAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAGCACUGAAGAAGAA 60
cISFV-NC_027817.1  NCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGCCUUCUUCCCCU 60
                   .........10........20........30........40........50........6
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                   ......((((((........)))))).................................((
cISFV-NC_001564.2  GGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAANNNNNNN--AGCGAACC 116
cISFV-NC_005064.1  CCUCCGGGCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCCGNNNNNNN----CGUACC 114
cISFV-NC_012932.1  CCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGGNNNNNNN-UUGCGAACC 117
cISFV-NC_034204.1  CCGGGAGGUGGUCGGGCCCGGCCCAGAGAGGGGCUGGGGAAGANNNNNNN---ACGACCA 115
cISFV-NC_034017.1  CCGGGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGGCGAAGANNNNNNN---GCGACCA 115
cISFV-NC_008604.2  GGGGCCGGGCUCCUCCGGAUGGUUGUUACCA--CCGGGGCGCGNNNNNNN-CUGACGCCA 117
cISFV-NC_021069.1  GAGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAANNNNNNN-GUGACGCCA 117
cISFV-NC_012671.1  GAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAGNNNNNNN-GUGACGCCA 117
cISFV-NC_024299.2  GAGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAAANNNNNNN----ACGACA 114
cISFV-NC_027819.1  GGGAAUUUCCCUGGGAGGAGGAGAAACGGGGCUUUGAG--ANNNNNNNN-UUAGCCACA 117
cISFV-NC_027817.1  CCGAGGGAAGAA--AAAGAAAGCCCCUGUGAUACAAUCACAGGNNNNNNNAGUAGCGUCA 118
                   0........70........80........90........100.......110.......1
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                   (...........)))........................................
cISFV-NC_001564.2  UCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAACUCAUCUUAACCUCCCAC-- 168
cISFV-NC_005064.1  CCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCCAGCUAAAUCUCCCACCG 168
cISFV-NC_012932.1  CCCUCCG-AAAUGUGAAAAAUUAUGGGGAAAAGUACCCAUCUAAGCCUCCCA--- 168
cISFV-NC_034204.1  CCCUCUG-AAAUGUUAAUAAAACUGUGCGGAAAAAUCCAAAGACUGUCCCCACC- 168
cISFV-NC_034017.1  CCCUCUG-UAAUGUUUAAUAAGUUGUGCGGAAAAAUCCAAAGAAUGUCCCCACC- 168
cISFV-NC_008604.2  CCUCCCG-AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCAGCAGGUCUCCA--- 168
cISFV-NC_021069.1  CCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAAAUCUCCAC-- 168
cISFV-NC_012671.1  CCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAAAUCUCCAC-- 168
cISFV-NC_024299.2  CCCCCCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAACAGGCUUCCACCG 168
cISFV-NC_027819.1  CCCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAUGGAAGAUCCCCA--- 168
cISFV-NC_027817.1  CCCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAAAGUCGAGAGUCUC---C 168
                   20.......130.......140.......150.......160.......170...
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