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                                        .                                                                                                    .
cISFV-NC_001564.2  AAUUUUUGCCGAGAGACCGUGA----GGUUGAACCCGGCAAGGAAUGAAGAGGAAGG---AUUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCG 100
cISFV-NC_005064.1  AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA---GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGGUU 100
cISFV-NC_012932.1  AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAUGGAUGC---UCUUGAAGAAAGGGCCUUCGGGCCCAAACAUCUGGUCGCCAUCGACC 100
cISFV-NC_008604.2  AAGGCUUGAGUUGUUU--AUAAUAGUCGUUUUUUCUCGCAGAA--AUGGGAAAGGACGACGGUAAGAAGAAGAAGGGGCCGGGCUCCUCCGGAUGGU-UGUUACCA-- 100
cISFV-NC_021069.1  AA--UUUCGCUACCUU--AUUACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GUCGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCC-UCCGACCAAC 100
cISFV-NC_012671.1  UA--AUUUGCUACUUU--AUUACCUCUGGUACUAACGUGGCGCUAUGAAGAAGAAGGA--GUCUGGGAGUGGUGGAGAGAGGACGUGGAACGCGUCC-UCCUGCCAAU 100
cISFV-NC_024299.2  ----CUUUGGUUGUUUGAAAAACUUUUUGCGACGUUGUUGAGAAGUGGAGCGAAAGAAAGGAGUAAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCC-CCCUCCCAUU 100
cISFV-NC_027819.1  UAGACGUUUGUCACGG----AAUAGCUUGAAGAAGAAGCUUACUUAUGUCAAGUAAGGAAGCACUGAAGAAGAAGGGGAAAUUUCCCUGGGAGGGAGGAG-GAAAACGGG 102
cISFV-NC_034204.1  ----AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAA--CGGAAAGUUUUCCCGACCGGGAGGUGGUGGGCCCCGGCCCAGAGA-GGGG 100
cISFV-NC_034017.1  ----CUAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGAGCAAUC--GAGGAAGUUUUCCCGGCCGGGAGGCGGUCGGGCCGGGCCCCAAAGA-GGGG 100
cISFV-NC_027817.1  ----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCCUGUGA---UA 100
                   .........10........20........30........40........50........60........70........80........90........100.....
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                                                                                                     (((((..(((((((....
cISFV-NC_001564.2  UUCUGGGGAANNNNNNNCUUUCAAAACUGGACGGGGCAACAGGGAGAAAUCCCGGGG--UA-GCGAACCUCCUCCGUUAAUGUGAAAAGUAUG-GGGAAAGAACUC 203
cISFV-NC_005064.1  GGACCGCCCGNNNNNNN--UUCAAAACCGGACGGAGCAACAGGGA-AUUUCCCGGGGA---C-GCGUACCCCCUCCG-UAAUGUGAAAAGUAUGGGGGAAAAGAACCC 200
cISFV-NC_012932.1  AGACCGCCGGNNNNNNN--A--AAAAUCGGGCGGAGCAACAGGGAGAAAUCCCGGGGAAUU-GCGAACCCCCUCCG-AAAUGUGAAAAAUUAUGGGGGAAAAGUACCC 201
cISFV-NC_008604.2  CCGGGGCNNNNNNNNUCACCGCAAGGUGAGGGGCGCACAGGAAAUAGA--AUCCAGGUGACU-GACGCCACCUCCCG-AAAUGUGUAUAGUAACAGA-GCAUGCCUGC 202
cISFV-NC_021069.1  CGUGGAGGAANNNNNNNUCACCGUAAGGUGAGGGGCGCACAGGAUCGAAUAUCCGGGUGAGU-GACGCCACCCCCCG-AAAUGUGUAUUGUGACAGG-CU-UGCUCGC 203
cISFV-NC_012671.1  CGUGGAGGAGNNNNNNN--CACCGCAAGGUGAGGGGCGCACAGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCCCCG-AAAUGUGUAUUGUGACAGG-CU-UGCUCGC 202
cISFV-NC_024299.2  AAGGAGAAAANNNNNNN--ACCGCAAGGUGAGGGCGCGCACAGGACGAAAGUCCAGGGUGAGU-GACGACACCCCCCG-AAAUGUGUAUAUAGCCCAU-GUUGAGUCUA 202
cISFV-NC_027819.1  CUUUGAG--ANNNNNNNAAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCCGGGUGAUU-AGCCACACCCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAG 205
cISFV-NC_034204.1  CUGGGCGAAGANNNNNNN--CUC-----CAGAGCAAACAAGGGAUUUAUCUUAAGGUGAUUUGACGACCACCCUCGUAAAUGUUAAUAAAACUCGUGCGGAAA-AAUC 199
cISFV-NC_034017.1  CUGGCGAAGANNNNNNN--ACCUC-----CAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUGGGACGACCACCCUCUCGUAAAUGUUUAAUAAGUUGUGCGACUA-AAUC 200
cISFV-NC_027817.1  CAAUCACAGGNNNNNNNNAUGUG-----CAGGGCAACAAAGUUCUAACGAACUAGGGUGAGUAG-CGUCACCCCCCGGUUGUGAAAACGAUUGCGACUAGAACUAAAG 201
                   ..110.......120.......130.......140.......150.......160.......170.......180.......190.......200.......210..
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                   ..(((((((((((((((.......))))))).)))))))))...)))))))))....)))))))))
cISFV-NC_001564.2  -AUCUUAAACCUCCCACCGUUAGGGAG--UUUUGAUUAAUCUUUCUAUACCAUAGAUGC--- 258
cISFV-NC_005064.1  -AGCUAAAUCUCCCACCGAUAGGGAG--UUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU 258
cISFV-NC_012932.1  -AUCUAAGCCUCCCACCGUUAGGGAG--UUUGGAAUACCUUUUCUAUACCAUAGAUGCGC- 258
cISFV-NC_008604.2  AGCAGCAGGUCUCCCACCGUUAGGGAGAC-UUGUUGCGGGCAAGCUC---UUGUUCACGU-CU 258
cISFV-NC_021069.1  AGCAGCAAAUCUCCCACCGUUAGGGAGAU-UUGUUGUGAGCAUUGC----UUGUCUACGU-CU 258
cISFV-NC_012671.1  AGCAGCAAAUCUCCCACCGUUAGGGAGAU-UUGUUGUGAGCAUUGCC---UUGUCUACGU-CU 258
cISFV-NC_024299.2  UACAACAGGCUUCCCACCGGUAGGGAAGCGUUGUUGUGAUGACAUCAU---GAG-CAACGU-CU 258
cISFV-NC_027819.1  A-UGGAAGAUCCCCCACCGUUAGGGGGAUCCAUCCAGCAUUCUUGGA---AUCAU---GG-CU 258
cISFV-NC_034204.1  CAAAGACUGUCCCCCACCGAUAGGGGAC--UUUGGGAAAGUUUUUCAACACACGUAGCGUCU 258
cISFV-NC_034017.1  CAAAGAAUGUCCCCCACCGAUAGGGGGC--UUUGGGAAAGUUUUUCAACACACGUAGCG-CU 258
cISFV-NC_027817.1  UCGAGAGUCUCCCCACCGAUAGGGGAG--CUCGUC-AGUUUGUUUUUAGUCAUUACG-CU 258
                   .....220.......230.......240.......250.......260.......270...
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