```
                                                                              ((((.......((((.......
         ....----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCCUGUGA---UA
cISFV-NC_027817.1  ----UUGAAAGGUUCAAACUCUGGUGUCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAGGCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCCUGUGA---UA 100
cISFV-NC_034204.1  ----AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAGAUGGAGCAAAA--CGGAAAGUUUUCCCGACCGGGAGGUGGUCGGGCCCGGCCCAGAGA-GGGG 100
cISFV-NC_034017.1  ----CUAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGUAUGGGAGCAAUC--GAGGAAGUUUUCCCGGCCCGGGAGGCGGUCGGGCCGGCCCCAAAGA-GGGG 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--AUUACCUCUGGUACUAAACGUGGCGCUAUGAAGAAGAAGGA--GUCUGGGAGUGGUGGAGAGAGGACGUGGAACGCGUCC-UCCUGCCAAU 100
cISFV-NC_024299.2  ----CUUUGGUUGUUUGUGAAAAACUUUUUGCGACGUUUGUUGAGAAGAGCGAAAAAAGGAGUAAAAUUAUGGGAGAAGGGAAAGC--UAUCCCGCC-CCCUCCCAUU 100
cISFV-NC_027819.1  UAGACGUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUUAUGUCAAGUAAGGAAGCACUGAAGAAGAAGAGGGAAUUUCCCUGGGAGGAGGAG-GAAAACGGGG 102
cISFV-NC_001564.2  AAUUUUUUGCCGAGAGACCCGUGA----GGUUGAACCCGGCCAAGGAAUGAAGAGGAAGG---AUUUGGAGGCGCGAGGAAAAGCCCGCAGGGGAGGGAUUUUUCCACCCCG 100
cISFV-NC_005064.1  AAAUCAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUCGUUAUGAAUAGAGGAA---GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGGUU 100
cISFV-NC_012932.1  AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUAUCUGGUUCGUAUGAAUAGGAUGC---UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACC 100
               .........10........20........30........40........50........60........70........80........90........100.....

               ))))............................................(((.....))).(((((........)))))..................(((((....((((((..
cISFV-NC_027817.1  CAAUCACAGGNNNNNNNNAUGUG-----CAGGGCAACAAAGUUCUAACGAACUAGCGUGAGUAG-CGUCACCCCCCGGUUGUGAAAACGAUUGC-GACUAGAACUAAA 200
cISFV-NC_034204.1  CUGGGGAAGANNNNNNN--CUC-----CAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCACCCUCUGAAAUGUUAAUAAAACU-GUGCGGAA-AAAU 198
cISFV-NC_034017.1  CUGGCGAAGANNNNNNNNACCUC-----CAGAGCAAACGAGGGAGAAAUCCCU-AGGUGAGUUGGCGACCACCCUCUGUAAUGUUUAAUAAGUU-GUGCGGAA-AAAU 199
cISFV-NC_008604.2  CCGGGGCGCGNNNNNNNUCACCGCAAGGUGAGGGCGCACAGGAUAGA--AUCCACGUGACU-GACGCCACCUCCCG-AAAUGUGUAUAUGUAACAGAGCAUGCCUGCA 203
cISFV-NC_021069.1  CGUGGGAGGAGNNNNNNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAGUAUUCCGGGUGAGU-GACGCCACCCCCCG-AAAUGUGUAUUGUGACAGGCU-UGCUCGCA 204
cISFV-NC_012671.1  CGUGGGAGGAGNNNNNNN-CACCGCAAGGUGAGGGCGCACAGGAUCGAGUAUCCGGGUGAGU-GACGCCACCCCCG-AAAUGUGUAUUGUGACAGGCU-UGCUCGCA 203
cISFV-NC_024299.2  AAGGAGAAAANNNNNNN--ACCGCAAGGUGAGGGCGCACAGGACGAAAGUCCAGGGUGAGU-GACGCACACCCCCG-AAAUGUGUAUAUAGCCAUGUUGAGUCUAU 203
cISFV-NC_027819.1  CUUUGAG--ANNNNNNNAAAAGGCCAUCCACGCAGGGCAACAGGAAGAAAUUCCGGGUGAUU-AGCCACACCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAG 205
cISFV-NC_001564.2  UUCUGGGGGAANNNNNNNCUUUCAAAACUGGACGGGGCAACAGGGAGAAAUCCGGGGG---UA-GCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG---AAAGAAC 201
cISFV-NC_005064.1  GGACCGCCCGNNNNNNN--UUCAAAACGGACGGACGAACAGGGAGAAAUCCCGCGGGA--C-GCGAACCCCUCCCG-UAAUGUGAAAAAUUAUGGGG--AAAGAAC 198
cISFV-NC_012932.1  AGACCGCCGGNNNNNNN--A--AAAAAUCGGGCGGAGCAACAGGGAGAAAUCCCCGCGGGAUU-GCGAACCCCCUCCG-AAAUGUGAAAAAUUUAUGGGG--AAAGUAC 199
               ..110.......120.......130.......140.......150.......160.......170.......180.......190.......200.......210..

               ..(((((...(((((.......)))))..))))))..)))))).)))))))))..))))))).)).))
cISFV-NC_027817.1  GUCGAGAGUCUCCCCACCGAUAGGCGGAGCUCGUCAGUUUGUUUUAGUCCAAUUA--CGCU 258
cISFV-NC_034204.1  CCAAAGACUGUCCCCACCGAUAGGCGGGACUUUGGGAAAGUUUUUCAACACACGUAGCGUCU 258
cISFV-NC_034017.1  CCAAAGAAUGUCCCCACCGAUAGGCGGGCUUUGGGAAAGUUUUUCAACACACGUAGCG-CU 258
cISFV-NC_008604.2  GCAGCAGG--UCUCCACCGUUAGGAGAGACUUGUU-GCGGGCAAGCUCUUGUUCACGU-CU 258
cISFV-NC_021069.1  GCAGCAAA--UCUCCACCGUUAGGAGAU-UUGUU-GUGAGCAUUGCU-UGUCUACGU-CU 258
cISFV-NC_012671.1  GCAGCAAA--UCUCCACCGUUAGGAGAU-UUGUU-GUGAGCAUUGCU-UGUCUACGU-CU 258
cISFV-NC_024299.2  ACAACAGG--CUUCCACCGGUUAGGAAGCGUUGUU-GUGAUGACACAUGAG-CAACGU-CU 258
cISFV-NC_027819.1  AUGGAAGA--UCCCCACCGUUAGGGGAUCCAUCCAGCAUUCUUGGA-AUCAU---GG-CU 258
cISFV-NC_001564.2  UCAUCUUAACCUCCCCACCGUUAGGGGAGUUUUGAUUAUCUUUUCUAUACCAUAGAUGC--- 258
cISFV-NC_005064.1  CCAGCUAAAUCUCCCCACCGAUAGGGGAGUUGGACUAUCUUUUCUAUACCAUAAAUGCGCU 258
cISFV-NC_012932.1  CCAUCUAAGCUCUCCCCACCGUUAGGGGAGUUUGGAAUACCUUUUCUAUACCAUAGAUGCGC- 258
               .....220.......230.......240.......250.......260.......270..
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