```
                                  .........................(((((((((((((((((((((((
dISFV-NC_040610.1  GAACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAAUU-UCA------AUGGUUAGACAAC  53
dISFV-NC_024017.1  UCGCGCAUUUUGAU-AGAGUUGACUAGAUAACGAAUUAUCG------AUGGUGACUAAAU  53
dISFV-NC_027999.1  GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA-CU------AUGGUUGAGAAAA  53
dISFV-NC_016997.1  A-AACAUAACGCAUUAGAAUAGGUUUGAAAGAGUUGAAUCA------AUGAGAUCAAUGG  53
dISFV-NC_017086.1  AUAACACGUUUGAAUAGGUUUAUUGUG-AAUCGAAAUACCG------AUG--------G  44
TBFV-NC_003690.1   GUGUUGAGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGG  53
TBFV-NC_001809.1   GUGUUGAGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGA  53
TBFV-NC_005062.1   GUGUUGAGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGG  53
TBFV-NC_001672.1   GUGUUGAGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGG  53
TBFV-NC_027709.1   GUGUUGAGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGU  53
TBFV-NC_003687.1   ------AGAAAAAG-ACAGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG---AUGACCA  50
                   .........10........20........30........40........50........6


                   ((.(((((.....(..(((((((.............((.((((.
dISFV-NC_040610.1  UUGGA--AAGACC----GUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACCAC  105
dISFV-NC_024017.1  CAAGA--AAGCCC----GCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCA--UCACCAA  105
dISFV-NC_027999.1  CUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA--CUG----  107
dISFV-NC_016997.1  UCUUUCGAGCUAGGAUGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC---  110
dISFV-NC_017086.1  CGAAUAAACCCAAGAAGCCCGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCUUC  104
TBFV-NC_003690.1   CCGUUCUAAAAGGAAAGGGGGGGGGGUC--CCCCUCGACGAGCGUCGAAAGUGGCCCCAA-  110
TBFV-NC_001809.1   CCAUCCUGAAAGGCAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCGA-  110
TBFV-NC_005062.1   CCAUUCUGAAAGGAAAGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCUA-  110
TBFV-NC_001672.1   CCAUCCUGAAAGGUAAGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA-  110
TBFV-NC_027709.1   CCAUCCUGAAAGGGUAAGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA-  110
TBFV-NC_003687.1   C--UUCUAAAGGAAAGGGGGGCGGUCC--CCCUA-GGCGCAAGCUUAAAGUGACCGCAAA  105
                   0........70........80........90........100.......110.......1


                   ...................................)))).)).............)))))))))...
dISFV-NC_040610.1  CCG-CUNNNNNNNACCC----CCUCAUUAUUAUAUAA--CAGCAUAUUGACACCAGGG-  157
dISFV-NC_024017.1  GCG-UGNNNNNNNCCCC----ACUGCUGAGAG-CAAAA--UAGCAUAUUGACACUUGGGA  157
dISFV-NC_027999.1  --G-AANNNNNNNUGGCGA-ACCCCAUUGGAAAUUUAG--CUUCAUAUUGACACCAGGGA  161
dISFV-NC_016997.1  ------NNNNNNNAGGAAACCCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGGA  164
dISFV-NC_017086.1  GCGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUGAAGGGCACUAUAUCGACACUUGGGA  164
TBFV-NC_003690.1   ------NNNNNNNCUCUCUUCAGGGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUC  163
TBFV-NC_001809.1   ------NNNNNNN-UCUCUUCAGGGGUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGAC  163
TBFV-NC_005062.1   ------NNNNNNNCUCUCUUCAGGGAAAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCAUC  163
TBFV-NC_001672.1   ------NNNNNNNCUCUCUUCAGGGAUUUUUCCUCCUCCUAUACAAAA-UUCCCCCUCGGU  163
TBFV-NC_027709.1   ------NNNNNNNCUCUCUUCAGGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGAC  163
TBFV-NC_003687.1   UAA-GUNNNNNNNCU-UCCUCGGGAUUUUUCCGGCCUCCUAUACUAAAUUUCCCCCAGGAA  163
                   20.......130.......140.......150.......160.......170.......1


                   ..)))).)).)))))).....)))))))))).)))))))).....................
dISFV-NC_040610.1  AUAGACCGGGAGAAAU-ACACAAUUUCUGGGCACUAAACAACUACAAGCACAGA  209
dISFV-NC_024017.1  AGAGACCACGAGAUAUGAUGUUUGUUUGUCAAUGCU-GACAACAGCCAUGGCACA  209
dISFV-NC_027999.1  AAAGACCGGGAGAAA-----CUUCUUUGCUUACUCGCGACAACCACCGAAAGGCA  209
dISFV-NC_016997.1  A-AGACUGGGAGAUA-----C-CUUUGUUCUUCGACACUAACUCC-UAAUGGC  209
dISFV-NC_017086.1  A-AGACCAGAGGGUA-----CUCGCUGAUUCACCGCCACCAG-ACU-ACACGGC  209
TBFV-NC_003690.1   AGAGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCAUCACCUAGAC---  209
TBFV-NC_001809.1   AGAGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCAUCACCCAGAC---  209
TBFV-NC_005062.1   UGAGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCAUCACCCAGAC---  209
TBFV-NC_001672.1   AGAGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCAUCACCCAGAC---  209
TBFV-NC_027709.1   AGAGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCAUCACCCAGAC---  209
TBFV-NC_003687.1   ACUGGGGGGGCGGUU----CUUGUUCUCUCCUGAGCCACCACCAUCCAGGC---  209
                   80.......190.......200.......210.......220.......230.
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