```
                          .............(((((((((((((((((((((((
dISFV-NC_016997.1   AAAC--AUAACGCAUUAGAAUAGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGU  54
dISFV-NC_017086.1   ------AUAACACGUUUGAAUGGUGUUUAUUUGUGAAUCGAAAUACCGAUG--------GC  45
dISFV-NC_040610.1   --GA--ACAAUUUUAAAGAAGUCAAAC-UAGAAAAUUU-AAUUUCAUG--------GU  45
dISFV-NC_024017.1   UCGC--GCAUUUUGAUAGAGUU---GAC-UAGAUAACGAAUUAUCGAUG--------GU  45
dISFV-NC_027999.1   GGAU--UAAUUGUUGUUUUGAGAGAACAGCGAGAAGUACUA----CUAUG---GUUGAGAA  51
TBFV-NC_001809.1    GUGUUGAGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGAC  54
TBFV-NC_005062.1    GUGUUGAGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGC  54
TBFV-NC_027709.1    GUGUUGAGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGG------AUGGCCAGGAAGUC  54
TBFV-NC_001672.1    GUGUUGAGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGGC  54
TBFV-NC_003690.1    GUGUUGAGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGC  54
TBFV-NC_003687.1    ------AGAAAAAGACAGCUUAGGGAGAACAAGAGCUGGGAGUGGUUAUG------AUGAC  48
                    .........10........20........30........40........50.......6

                    ...............(((((((....)))))..)).))...........((.......))........
dISFV-NC_016997.1   CUUUCGAGCUAGGAGGCCCGU---UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC-  110
dISFV-NC_017086.1   GAAUAAACCCAAGAAGGCCCGG---AAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCC  102
dISFV-NC_040610.1   UAGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACC  103
dISFV-NC_024017.1   GACUAAAUCAAGAAGACCGCCGCAAAGCGGGCGUUUAAUAUGCUAACACGCA--UCACC  103
dISFV-NC_027999.1   AACUAGAGUAGGAUCUCGGGUUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA-------  104
TBFV-NC_001809.1    CAUCCUGAAAGGCAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_005062.1    CAUUCUGAAAGGAAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_027709.1    CAUCCUGAAAGGUAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_001672.1    CAUCCUGAAAGGUAAGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_003690.1    CGUUCUAAAAGGAAAGGGGGGCGGUCCCC--CUCGACGAGCGUCGAAAGUGG----CCCC  108
TBFV-NC_003687.1    CACUUCUAAAGGAAAGGGGGGCGGUCCCC--CUAGGCGCAAGCUUAAAGUGA----CCGC  102
                    0........70........80........90........100.......110.......1

                    ...................(((((((((((((...........))))))...
dISFV-NC_016997.1   --------NNNNNNN--AGGAAACCCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGA---CA  157
dISFV-NC_017086.1   UCGCGUCUNNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGA---CA  157
dISFV-NC_040610.1   ACCCGCU-NNNNNNN-------ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGA---CA  151
dISFV-NC_024017.1   AAGCGUG-NNNNNNN-------CCCCACU-GCUGAGAGCAAAAUAGCAUAUUGA---CA  150
dISFV-NC_027999.1   --CUGGAANNNNNNNNUGGCGAACCCCAUUGGAAAUU-----UAGCUUCAUAUUGA---CA  154
TBFV-NC_001809.1    GA------NNNNNNN-------UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUC  154
TBFV-NC_005062.1    UA------NNNNNNN-------CUCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UUC  154
TBFV-NC_027709.1    AA------NNNNNNN-------CUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUC  154
TBFV-NC_001672.1    AA------NNNNNNN-------CUCUCUUCAGGAUUUUUCCUCCUCCUAUACAAA-UUC  154
TBFV-NC_003690.1    AA------NNNNNNN-------CUCUCUUCAGGAUAUUUCCUCCUCCUAUACCAAA-UUC  154
TBFV-NC_003687.1    AAAUAAGUNNNNNNN-------CU-UCCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUC  154
                    20.......130.......140.......150.......160.......170.......1

                    ..........))))))))))).)))))))))))))))).))))))))...(((((((((((.(((....
dISFV-NC_016997.1   CCUAGGAA-AGACUGGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCUA--AUGGCACAG  213
dISFV-NC_017086.1   CUUGGGAA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUAC--ACGGCACAG  213
dISFV-NC_040610.1   CCAGGG-AUAGACCGGAGAAAU-ACACAAUUUCUGGGCACUAAACCAACUACA-AGCACAG  208
dISFV-NC_024017.1   CUUGGGAAGAGACCAGAGAUAUGAUGUUGUUUUGUCAAUGCUGACAACAGCCAUGGCACAG  210
dISFV-NC_027999.1   CCAGGGAAAAGACCGGAGAAACUUCUUGCUUACUCG-CAGCAACACCGAAA-GGCACAG  212
TBFV-NC_001809.1    CCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCAUCACCCA-GACACAG  213
TBFV-NC_005062.1    CCCCUCAUCUGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCAUCACCCA-GACACAG  213
TBFV-NC_027709.1    CCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCAUCACCCA-GACACAG  213
TBFV-NC_001672.1    CCCCUCGGUUGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCAUCACCCA-GACACAG  213
TBFV-NC_003690.1    CCCCUCGUCAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCAUCACCUA-GACACAG  213
TBFV-NC_003687.1    CCCCAGGAAACUGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCACCACCAUCCA-GGCACAG  213
                    80.......190.......200.......210.......220.......230.......2

                    ...))))))......))))))))).........................
dISFV-NC_016997.1   AGCGCCAUA-AGAA--ACGGAGUUAGUAUGGAAAAAACAUAA-GUAUCU---  258
dISFV-NC_017086.1   CGCGCCGGA-AAAG--GUAGUUUGGAGGUUGUAAAACAACAA-GUAUCU---  258
dISFV-NC_040610.1   AGCGCUUCG-AGAAGUAGUUGUUUCUUGACGAGAGAAUAU-GUAUAUCUUGG  258
dISFV-NC_024017.1   AGCGCCUGA-AGAGAGCUGUUGUUUUAUAAUUUUACAAAAAAUAACAUAUCU---  258
dISFV-NC_027999.1   UAUGCCAGA-AACAUUCGGUGGUUU--GAUGUAGUAAAACAACCAGUUUCU---  258
TBFV-NC_001809.1    AUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAACAACAC-------CCGCU  258
TBFV-NC_005062.1    GCAGUCUAACAAGGAGGUGAUGUGUGAGUCUCGGAACAACAC-------CCGCU  258
TBFV-NC_027709.1    AUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAAACAC-------CCGCU  258
TBFV-NC_001672.1    GUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACAC-------CCGCU  258
TBFV-NC_003690.1    AUAGUCUGAAAAGGAGGAUGGUGCGUGUCUCGGAAAAACAC-------CCGCU  258
TBFV-NC_003687.1    AUAGCCUGACAAGGAGGAUGGUGUGUGACUCGGAAAAACAC-------CCGCU  258
                    40.......250.......260.......270.......280.......290
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