```
                       ...........................(((((((((((((((((((((((...............................((
dISFV-NC_040610.1  --GA--ACAAUUUUAAAGAAG-UCAAAACUAGAAAUUUAAUU-UCAAUG---------GU  45
dISFV-NC_024017.1  UCGC--GCAUUUUGAUAGAGU-U---GACUAGAUAACGAAUUAUCGAUG---------GU  45
dISFV-NC_016997.1  AAAC--AUAACGCAUUAGAAUAGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGU  54
dISFV-NC_017086.1  ------AUAACACGUUUAGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG---------GC  45
dISFV-NC_027999.1  GGAU--UAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA----CUAUG---GUUGAGAA  51
TBFV-NC_003690.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGC  54
TBFV-NC_005062.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGC  54
TBFV-NC_001809.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGAC  54
TBFV-NC_027709.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGUC  54
TBFV-NC_001672.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGGC  54
TBFV-NC_003687.1   ------AGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG------AUGAC  48
                   .........10........20........30........40........50.......6

                   ( . (((( . . . . . . . . (((((((( . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
dISFV-NC_040610.1  UAGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACC  103
dISFV-NC_024017.1  GACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCA--UCACC  103
dISFV-NC_016997.1  CUUUCGAGCUAGGAGGCCCGU---UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC-  110
dISFV-NC_017086.1  GAAUAAACCCAAGAAGCCCGG---AAGACGGGCUAUCGAUAUAGUGAGACGGUGCGUUGCC  102
dISFV-NC_027999.1  AACUAGAGUAGGGAUCUCGGGUUAAAACCCGAGGGUGUCAAUAUGAUUAGGCGUA-------  104
TBFV-NC_003690.1   CGUUCUAAAAGGAAAGGGGGGGGGUCCCC--CUCGACGAGCGUCGAAAGUGG----CCCC  108
TBFV-NC_005062.1   CAUUCUGAAAGGAAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_001809.1   CAUCCUGAAAGGCAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_027709.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_001672.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC  108
TBFV-NC_003687.1   CACUUCUAAAGGAAAGGGGGGCGGUCCCC--CUAGGCGCAAGCUUAAAGUGA----CCGC  102
                   0........70........80........90........100.......110.......1

                   ...........................................................
dISFV-NC_040610.1  ACCCG-CUNNNNNNNN-------UUAGAGGAGACCCCCUCAU-UAUUUAUAUAACAGCAUA  154
dISFV-NC_024017.1  AAGCG-UGNNNNNNNN-------UAGAGGAGACCCCACUGC-UGAGAGCAAAAUAGCAUA  153
dISFV-NC_016997.1  --------NNNNNNNN--AUCGGUUAGAGGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUA  160
dISFV-NC_017086.1  UCGCGUCUNNNNNNNNCUAGCGGUUAGAGGAGACCCCCCC--AAAAAUGAAGGGCACUAUA  160
dISFV-NC_027999.1  --CUGGAANNNNNNNN--AAUGGGAAGUGGCGAACCCCAU---UGGAAAUUUAGCUUCAUA  157
TBFV-NC_003690.1   AA------NNNNNNNN-------AAUGGCAACUCUCUUCAGGAUAAUUUCCUCCUCCUAU  154
TBFV-NC_005062.1   UA------NNNNNNNN-------AAUUGGCAGCUCUCUUCAGGAUAAUUUCCUCCUCCUAU  154
TBFV-NC_001809.1   GA------NNNNNNNN-------AAUUGGCAGCUCUCUUCAGGGUUUUUUCCUCCUCCUAU  153
TBFV-NC_027709.1   AA------NNNNNNNN-------AAUUGGCAGCUCUCUUCAGGAUUUUUUCCUCCUCCUAU  154
TBFV-NC_001672.1   AA------NNNNNNNN-------AAUUGGCAGCUCUCUUCAGGAUUUUUUCCUCCUCCUAU  154
TBFV-NC_003687.1   AAAUAAGUNNNNNNNN-------AUUGGCAAUCUUCCUCGGGAUUUUUUCCGCCUCCUAU  153
                   20.......130.......140.......150.......160.......170.......1

                   ........))))))))........))))..).)))))))))))))))))))).)))))).........
dISFV-NC_040610.1  UUGACACCAGGG-AUAGACCGGAGAAAUACACAAUUUCUCGGGCACUAAACAACUACAAGC  213
dISFV-NC_024017.1  UUGACACUUGGGGAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUG  213
dISFV-NC_016997.1  UCGACACCUAGGAA-AGACUGGAGAUAC-CUUUGGUACCACACUAAACUCCU-----A  213
dISFV-NC_017086.1  UCGACACUUGGGAA-AGACCAGAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA-----C  213
dISFV-NC_027999.1  UUGACACCAGGGAAAAGACCGGAGAAAACUUCUUGCUUACUCGCAGCAACCACCGA----A  213
TBFV-NC_003690.1   ACCAAA-UUCCCCCUCGUCAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACC  213
TBFV-NC_005062.1   ACCAAA-UUCCCCCUCAUCUGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACC  213
TBFV-NC_001809.1   ACCAAAUUUCCCCCUCAUCAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACC  213
TBFV-NC_027709.1   ACCAAA-UUCCCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACC  213
TBFV-NC_001672.1   ACAAAA-UUCCCCCUCGGUAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACC  213
TBFV-NC_003687.1   ACUAAAUUUCCCCAGGAAACUGGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCACCAUC  213
                   80.......190.......200.......210.......220.......230.......2

                   ....
dISFV-NC_040610.1  ACAG  217
dISFV-NC_024017.1  GCAC  217
dISFV-NC_016997.1  AUGG  217
dISFV-NC_017086.1  ACGG  217
dISFV-NC_027999.1  AGGC  217
TBFV-NC_003690.1   UAGA  217
TBFV-NC_005062.1   CAGA  217
TBFV-NC_001809.1   CAGA  217
TBFV-NC_027709.1   CAGA  217
TBFV-NC_001672.1   CAGA  217
TBFV-NC_003687.1   CAGG  217
                   40..
```