```
                    .........(((((((((((((((((((((((((..............................(
dISFV-NC_016997.1  A-AACAUAACGCAUUAGAAUAGGUUUGAAAGAGUUGAAUCA-----AUGAGAUCAAUGGU  54
dISFV-NC_017086.1  AUAACACGUUUGAAUAGGUUUAUUGUG-AAUCGAAAUACCG-----AUG--------GC  45
dISFV-NC_040610.1  GAACAAUUUUAAAGAAGUCAAAAACUAGAAUUUUAAUU-UCA-----AUG------GUUAG  48
dISFV-NC_024017.1  UCGCGCAUUUUGAU-AGAGUUGACUAGAUAACGAAUUAUCG-----AUG------GUGAC  48
dISFV-NC_027999.1  GGAUUAAUUGUUGUUUGAGAGAACAGCGGAGAAGUACUA-CU-----AUGGUUGAGAAAAC  54
TBFV-NC_003690.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGC  54
TBFV-NC_027709.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGUC  54
TBFV-NC_001672.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAGAAGGC  54
TBFV-NC_001809.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCAGGAAGAC  54
TBFV-NC_005062.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGC  54
TBFV-NC_003687.1   ------AGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG---AUGACCAC  51
                   .........10.......20.......30.......40.......50.......6
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                   (.((((.........((((((((..............................
dISFV-NC_016997.1  CUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC----  110
dISFV-NC_017086.1  GAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCG  105
dISFV-NC_040610.1  ACAACUUGGAAAGACCGGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACCACC  106
dISFV-NC_024017.1  UAAAUCAAGAAAGCCCGCCGCCAAAGCGGGCUGUUAAAUACGCAACACGCA--UCACCAAG  106
dISFV-NC_027999.1  UAGAGUAGGAUCUCGCGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA--CUGGAA--  110
TBFV-NC_003690.1   CGUUCUAAAAGGAAAGGGGGGGGGUC--CCCCUCGACGAGCGUCGAAAGUGGCCCCAA--  110
TBFV-NC_027709.1   CAUCCUGAAAGGUAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA--  110
TBFV-NC_001672.1   CAUCCUGAAAGGUAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA--  110
TBFV-NC_001809.1   CAUCCUGAAAGGCAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA--  110
TBFV-NC_005062.1   CAUUCUGAAAGGAAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCUA--  110
TBFV-NC_003687.1   --UUCUAAAAGGAAAGGGGGGCGGUCC--CCCUA-GGCGCAAGCUUAAAGUGACCGCAAAU  106
                   0.......70.......80.......90.......100.......110.......1
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                   ..............................................))))))))).....
dISFV-NC_016997.1  -----NNNNNNNAGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGGAA  165
dISFV-NC_017086.1  CGUCUNNNNNNNNAGAGGAGACCCCCCCAAAAAUGAAGGGCACUAUAUCGACACUUGGGAA  165
dISFV-NC_040610.1  CG-CUNNNNNNNA------CCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGGG-A  158
dISFV-NC_024017.1  CG-UGNNNNNNNC------CCCACUGCUGAGAG-CAAAAUAGCAUAUUGACACUUUGGGAA  158
dISFV-NC_027999.1  -----NNNNNNNUGGCGAACCCCAUUGGAAAUU---UAGCUUCAUAUUGACACCAGGGAA  162
TBFV-NC_003690.1   -----NNNNNNNCUCUCUUCAGGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUCA  164
TBFV-NC_027709.1   -----NNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGACA  164
TBFV-NC_001672.1   -----NNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGGUA  164
TBFV-NC_001809.1   -----NNNNNNN-UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUCCCCCUCGACA  164
TBFV-NC_005062.1   -----NNNNNNNCUCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCAUCU  164
TBFV-NC_003687.1   AA-GUNNNNNNNCU-UCCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGAAA  164
                   20.......130.......140.......150.......160.......170.......1
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                   .))))).)).))))))))))))))))))))).))))).......................
dISFV-NC_016997.1  -AGACUGGAGAUACCUUUGGUUCUUCGACACU-AACUCCU--AAUGGCACAGAGCGCCA  220
dISFV-NC_017086.1  -AGACCAGAGAGGUACUCGCUGAUUCACCGCCACC-AG-ACUA--CACGGCACAGCGCGCCG  220
dISFV-NC_040610.1  UAGACCGGAGAAAU-ACACAAUUUCUGCUCUCUCACAAAUACAACUACAAGCACAGAGCGCUUCG  217
dISFV-NC_024017.1  GAGACCGAGAUAUGAUGUUGUUUGUCAAUGCU-GACAGCCAUGGCACAGAGCGCCU  217
dISFV-NC_027999.1  AAGACCGGAGAAACUUCUUGCUUACUCG-CAGCAACCACCG--AAAGGCACAGUAUGCCU  219
TBFV-NC_003690.1   GAGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCAUCACCU----AGACACAGAUAGUCU  220
TBFV-NC_027709.1   GAGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCAUCACCC----AGACACAGAUAGUCU  220
TBFV-NC_001672.1   GAGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCAUCACCC----AGACACAGGUAGUCU  220
TBFV-NC_001809.1   GAGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCAUCACCC----AGACACAGAUAGUCU  220
TBFV-NC_005062.1   GAGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCAUCACCC----AGACACAGGCAGUCU  220
TBFV-NC_003687.1   CUGGGGGGGCGGUUCUUGGUUCUCCCUGAGCCACCACCAUCC----AGGCACAGAUAGUCCU  220
                   80.......190.......200.......210.......220.......230.......2
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                   ..................................................
dISFV-NC_016997.1  UA-AG--AAACGGAGUUAGUAUGGAAAAAACAUAA-GUAUCU  258
dISFV-NC_017086.1  GA-AA--AGGUAGUUUGGAGGUUGUAAAACAACAA-GUAUCU  258
dISFV-NC_040610.1  AG-AAGUAGUUGUUUCUUGACGAGAGAAUAUGUAUAUCUUGG  258
dISFV-NC_024017.1  GA-AGAGAGCUGUUGUUUAUAAUUUACAAAAAAUAACAUAUCU  258
dISFV-NC_027999.1  GA-AACUUCGGUGUUGAUGGUAGUAAAACAACCA--GUUUCU  258
TBFV-NC_003690.1   GAAAAGGAGGUGAUGCGUGUCUCGGAAAAACAC----CCGCU  258
TBFV-NC_027709.1   GACAAGGAGGUGAUGCGUGACUCGGAAAAACAC----CCGCU  258
TBFV-NC_001672.1   GACAAGGAGGUGAUGUGUGACUCGGAAAAACAC----CCGCU  258
TBFV-NC_001809.1   GACAAGGAGGUGAUGUGUGACUCGGAAAAACAC----CCGCU  258
TBFV-NC_005062.1   AACAAGGAGGUGAUGUGUGACUCGGAACAAAACAC----CCGCU  258
TBFV-NC_003687.1   GACAAGGAGAUGGUGUGUGACUCGGAAAAACAC----CCGCU  258
                   40.......250.......260.......270.......280
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