```
        ((((((.............((((((((((((((((((((((((((((............................(
dISFV-NC_016997.1  A-AACAUAACGCAUUAGAAUAGGUUUGAAAGAGUUGAAUCA-----AUGAGAUCAAUGGU 54
dISFV-NC_017086.1  AUAACACGUUUGAAUAGGUUUAUUGUG-AAUCGAAAUACCG-----AUG--------GC 45
dISFV-NC_040610.1  GAACAAUUUUAAAGAAGUCAAAACUAGAGUAUUUAAAU-UCA-----AUG------GUUAG 48
dISFV-NC_024017.1  UCGCGCAUUUUGAU-AGAGUUGACUAGAUAACGAAUCAUCG-----AUG------GUGAC 48
dISFV-NC_027999.1  GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA-CU-----AUGGUUGAGAAAAC 54
TBFV-NC_003690.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGC 54
TBFV-NC_027709.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCAGGAAGUC 54
TBFV-NC_001672.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAGGAAGC 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

        (.((((.........((((((((.........................((.((((............
dISFV-NC_016997.1  CUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC---- 110
dISFV-NC_017086.1  GAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCG 105
dISFV-NC_040610.1  ACAACUUGGAAAGACCGGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACCACC 106
dISFV-NC_024017.1  UAAAUCAAGAAAGACCCGCGUCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCA--UCACCAAG 106
dISFV-NC_027999.1  UAGAGUAGGAUCUCGGUUAAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA--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   CAUCCUGAAAGGCAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCAA-- 110
TBFV-NC_005062.1   CAUUCUGAAAGGAAAGGGGGGCGGUC--CCCCUCGACGAGUGUCGAAAGAGACCGCUA-- 110
TBFV-NC_003687.1   --UUCUAAAAGGAAAGGGGGGCGGUCC--CCCUA-GGCGCAAGCUUAAAGUGACCGCAAAU 106
        0.......70.......80.......90.......100.......110.......1

        .....................................)))).))...........)))))))))).....
dISFV-NC_016997.1  -----NNNNNNNAGGAAACCCCCCCAAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGGAA 165
dISFV-NC_017086.1  CGUCUNNNNNNNNAGAGGAGACCCCCCCAAAAAUGAAGGGCACUAUAUCGACACUUGGGAA 165
dISFV-NC_040610.1  CG-CUNNNNNNNNA------CCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGGG-A 158
dISFV-NC_024017.1  CG-UGNNNNNNNNC------CCCACUGCUGAGAG-CAAAAUAGCAUAUUGACACUUGGGAA 158
dISFV-NC_027999.1  -----NNNNNNNUGGCGAACCCCAUUGGAAAUU---UAGCUUCAUAUUGACACCAGGGAA 162
TBFV-NC_003690.1   -----NNNNNNNCUCUCUUCAGGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUCA 164
TBFV-NC_027709.1   -----NNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUCA 164
TBFV-NC_001672.1   -----NNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGGUA 164
TBFV-NC_001809.1   -----NNNNNNN-UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUCCCCCUCGUCA 164
TBFV-NC_005062.1   -----NNNNNNNCUCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCAUCU 164
TBFV-NC_003687.1   AA-GUNNNNNNNCU-UCCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGAAA 164
        20.......130.......140.......150.......160.......170.......1

        .))))).)).)))))))))))))))))))).)))))..............................)))))......
dISFV-NC_016997.1  -AGACUGGGAGAUACCUUUGGUUCUUCGACACU-AACUCCU--AAUGGCACAGAGCGCCA 220
dISFV-NC_017086.1  -AGACCAGAGGUACUCGCUGAUUCACCGCCACC-AG-ACUA--CACGGCACAGCGCGCCG 220
dISFV-NC_040610.1  UAGACCGGAGAAAU-ACACAAUUUCUGGCACUCUCAAACAACUACAAGCACAGAGCGCCUUCG 217
dISFV-NC_024017.1  GAGACCGAGAGAUAUGAUGUUGUUUGUCAAUGCU-GACAACAGCCAUGGCACAGAGCGCCU 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-AGAGAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU 258
dISFV-NC_027999.1  GA-AACUUCGUGUUGAUGUAGUAAACAACCA--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   AACAAGGAGGUGAUGUGUGACUCGGAAGAACAC----CCGCU 258
TBFV-NC_003687.1   GACAAGGAGAUGGUGUGUGACUCGGAAAAACAC----CCGCU 258
        40.......250.......260.......270.......280
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