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......
                                        dISFV-NC_016997.1
dISFV-NC_017086.1
dISFV-NC_040610.1
dISFV-NC_024017.1
dISFV-NC_027999.1
AISFV-NC_027999.
TBFV-NC_001809.1
TBFV-NC_005062.1
TBFV-NC_027709.1
TBFV-NC_001672.1
TBFV-NC_003690.1
 TBFV-NC 003687.1
                                             ......10.......20.......30........40.......50........6
                                         dISFV-NC_016997.1
dISFV-NC_017086.1
dISFV-NC_040610.1
dISFV-NC_024017.1
dISFV-NC_027999.1
TBFV-NC_001809.1
TBFV-NC_005062.1
TBFV-NC_027709.1
TBFV-NC_001672.1
TBFV-NC_001672.1
TBFV-NC_003690.1
TBFV-NC_003687.1
                                         CA<mark>CU</mark>UCUAAAGGAAAG<mark>G</mark>GG<mark>-G</mark>CGGUCCCC--CUAGGCGCAAGCUUAAAGUGA----CCGC
TBFV-NC_003687.1
                                         0.......70.......80........90.......100......110.......1
dISFV-NC_016997.1
dISFV-NC_017086.1
dISFV-NC_040610.1
dISFV-NC_024017.1
dISFV-NC_027999.1
TBFV-NC_001809.1
TBFV-NC_005062.1
TBFV-NC_027709.1
TBFV-NC_001672.1
TBFV-NC_001672.1
                                         -----NNNNNN--AGGAAACCCCCCAAAAAUGUAUAGGCUAUUAUAUCGA---CA 157
                                         UCGCGUCUNNNNNNAGAGGAGACCCCCCAAAAAUG-AAGGGCACUAUAUCGA--CA 157
ACCCGCU-NNNNNNN-----ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGA--CA 151
AAGCGUG-NNNNNNN-----CCCCACU-GCUGAGAGCAAAAUAGCAUAUUGA--CA 150
--CUGGAANNNNNNNUGGCGAACCCCAUUGGAAAUU
                                         GA-----NNNNNN-------UCUCUUCAGGGUUUUUCCUCCUCUAUACCAAAUUU<mark>C</mark>
                                                                                                                                                                   154
                                         UA-----NNNNNN------CUCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UU<mark>C</mark>
                                                                                                                                                                   154
                                         AA----NNNNNNN-----CUCUCUUCAGGAUUUUUCCUCCUCUUAUACCAAA-UUC
AA----NNNNNNN-----CUCUCUUCAGGAUUUUUCCUCCUCUAUACAAAA-UUC
AA----NNNNNNNN-----CUCUCUUCAGGAUUUUUCCUCCUCUAUACAAAA-UUC
                                                                                                                                                                    154
                                                                                                                                                                   154
TBFV-NC_003690.1
TBFV-NC_003687.1
                                                                                                                                                                   154
                                         AAAUAAGUNNNNNNN------CU-UCCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUU<mark>C</mark>
                                         20......130......140......150......160......170......1
                                         )))))).....((((...
dISFV-NC_016997.1
dISFV-NC_017086.1
dISFV-NC_040610.1
dISFV-NC_024017.1
dISFV-NC_027999.1
TBFV-NC_001809.1
TBFV-NC_005062.1
TBFV-NC_027709.1
                                          CUAGGAA-AGACUGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCUA--AUGGCACAG 213
UUGGGAA-AGACCAGAGGUACUCCCUGAUUCACCGCCACCAG-ACUAC-ACGGCACAG 213
CAGGG-AUAGACCGGAGAAAU-ACACAAUUUCUGGGCACUAAACAACUACA-AGCACAG 208
UUGGGAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACAG 210
CAGGGAAAAGACCGGAGAAACUUUUUGCUUACUCG-CAGCAACCACCGAAA-GGCACAG 212
                                          CAGGGAAAAGACCAGAGAAACUU UUGUUCUCG-CAGCAAACCACCGAAA-GGCACAG
CCCUCGACAGAGGGGGGGCGGUUUUGUUCUCCCUGAGCACACCAUCACCCA-GACACAG
CCCUCGACAGAGGGGGGGCGGUUUUGUUCUCCCUGAGCCACAUCACCCA-GACACAG
CCCUCGACAGAGGGGGGGCGGUUUUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG
CCCUCGGUAGAGGGGGGGGGGGGGUUUUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG
CCCUCGUCAGAGGGGGGGGGCGGUUUUUGUUCUCCCUGAGCCACCAUCACCUA-GACACAG
CCCAGGAAACUGGGGGGGCGGUUUUUGUUCUCCCUGAGCCACCAUCACCUA-GACACAG
                                                                                                                                                                    213
TBFV-NC_001672.1
TBFV-NC_003690.1
TBFV-NC_003687.1
                                                                                                                                                                    213
                                                                                                                                                                    213
                                         80......190......200.......210.......220.......230.......2
                                         dISFV-NC_016997.1
dISFV-NC_017086.1
dISFV-NC_040610.1
dISFV-NC_024017.1
dISFV-NC_027999.1
TBFV-NC_001809.1
TBFV-NC_005062.1
TBFV-NC_027709.1
TBFV-NC_001672.1
TBFV-NC_003690.1
TBFV-NC_003687.1
                                         AGCGCCAUA-AGAA-ACGGA<mark>GU</mark>UAGUAUGGAAAAAACAUAA-GUAUCU--- 258
CGCGC<mark>C</mark>GGA-AAA<mark>G</mark>-GUAGUU<mark>U</mark>GGAGGUUGUAAAACAACAA-GUAUCU--- 258
                                         AGCGC<mark>U</mark>UCG-AGA<mark>A</mark>GUAGUUGU<mark>U</mark>UCUUGACGAGAGAAUAU-GUAUAUCUUGG 258
                                         AGCGC<mark>C</mark>UGA-AGA<mark>G</mark>AGCUGUU<mark>GU</mark>UAUAAUUUACAAAAAAUAACAUAUCU---
                                         UAUGCCUGA-AACAUUCGGUG<mark>GU</mark>U--GAUGUAGUAAACAACCAGUUUCU---
                                         AUAGU<mark>C</mark>UGACAAG<mark>G</mark>AGGUGAU<mark>GU</mark>GUGACUCGGAAAAACAC-----CCGCU 258
                                         GCAGUCUAACAAGGAGGUGAUGUGUGACUCGGAACAACAC-----CCGCU 258
AUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAAACAC-----CCGCU 258
GUAGUCUGACAAGGAGGUGAUGUGUGUGACUCGGAAAAACAC-----CCGCU 258
                                         AUAGU<mark>C</mark>UGAAAAG<mark>G</mark>AGGUGAU<mark>G</mark>CGUGUCUCGGAAAAACAC-----CCGCU 258
                                         AUAGCCUGACAAGGAGAUGGUGUGUGACUCGGAAAAACAC-----CCGCU 258
                                         40......250......260......270......280......290
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