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                                    .........(((((((((((((((((((((((((............................(
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--AUUACGCAUUAGAAUAGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGU   54
dISFV-NC_017086.1  ------AUAACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG---------GC   45
dISFV-NC_027999.1  GGAU--UAAUUGUUGUUUUGAGAGAACAGCGAGAAGUACUA----CUAUG---GUUGAGAA 51
TBFV-NC_027709.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGUC 54
TBFV-NC_001672.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGGC 54
TBFV-NC_001809.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGGC 54
TBFV-NC_005062.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGC 54
TBFV-NC_003690.1   GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGC 54
TBFV-NC_003687.1   ------AGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG------AUGAC 48
                   .........10........20........30........40........50.......6
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                   (.((((.........(((((((.(((.....................(((((((..(((.((........
dISFV-NC_040610.1  UAGACAACUUGGAAAGACCCUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACC 103
dISFV-NC_024017.1  GACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCA--UCACC 103
dISFV-NC_016997.1  CUUUCGAGCUAGGAGGCCCGU---UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC- 110
dISFV-NC_017086.1  GAAUAAACCCAAGAAGCCCCG---AAGACGGGCUAUCGAUAUAUGAGACGCGUGCGUUGCC 102
dISFV-NC_027999.1  AACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA------- 104
TBFV-NC_027709.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC 108
TBFV-NC_001672.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC 108
TBFV-NC_001809.1   CAUCCUGAAAGGCAAGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGA----CCGC 108
TBFV-NC_005062.1   CAUUCUGAAAGGAAAGGGGGGCGGUCCCC--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
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                   ....................))).))))...)))))))))..............)))....))
dISFV-NC_040610.1  ACCCG-CUNNNNNNNN--------ACCCCCUCAUUAUUUUAUAUAACAGCAUAUUGA---CA 151
dISFV-NC_024017.1  AAGCG-UGNNNNNNNN--------CCCCACUGCUGAGAG-CAAAAUAGCAUAUUGA---CA 150
dISFV-NC_016997.1  --------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGA---CA 157
dISFV-NC_017086.1  UCGCGUCUNNNNNNNNAGAGGAGACCCCCCAAAAAUG--AAGGGCACUAUAUCGA---CA 157
dISFV-NC_027999.1  --CUGGAANNNNNNNNUGGCGA-ACCCCAUUGGAAAUUU----AGCUUCAUAUUGA---CA 154
TBFV-NC_027709.1   AA------NNNNNNNC-------UCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA--UUC 154
TBFV-NC_001672.1   AA------NNNNNNNC-------UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAA-UUC 154
TBFV-NC_001809.1   GA------NNNNNNNN-------UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUC 154
TBFV-NC_005062.1   UA------NNNNNNNC-------UCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA--UUC 154
TBFV-NC_003690.1   AA------NNNNNNNC-------UCUCUUCAGGAUAUUUCCUCCUCCUAUACCAAA--UUC 154
TBFV-NC_003687.1   AAAUAAGUNNNNNNNC-------U-UCCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUC 154
                   20.......130.......140.......150.......160.......170.......1
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                   )))))).......)))).)))))))))))))))))).)))))))..........
dISFV-NC_040610.1  CCAGGG-AUAGACCGGAGAAAUACACAAUUUCUGGGCACUAAACAACUACAA--GCACAG 208
dISFV-NC_024017.1  CUUGGGAAGAGACCAGAGAUAUGAUGUUGUUUUGUCAAUGCUGACAACAGCCAUGGCACAG 210
dISFV-NC_016997.1  CCUAGGAA-AGACUGGAGAGAUAC-CUUUGGUUCCUUCCACCACUCCU--AAUGGCACAG 213
dISFV-NC_017086.1  CUUGGGAA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA--CACGGCACAG 213
dISFV-NC_027999.1  CCAGGGAAAAGACCGGAGAAACUUCUUGCUUACUCGCAGCAACCACCG--AAAGGCACAG 212
TBFV-NC_027709.1   CCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG 213
TBFV-NC_001672.1   CCCCUCGGUAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG 213
TBFV-NC_001809.1   CCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG 213
TBFV-NC_005062.1   CCCCUCAUCUGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAG 213
TBFV-NC_003690.1   CCCCUCGUCGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCUA-GACACAG 213
TBFV-NC_003687.1   CCCCAGGAAACUGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCACCAUCCA-GGCACAG 213
                   80.......190.......200.......210.......220.......230.......2
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                   ..........................................................
dISFV-NC_040610.1  AGCGCUUC--GAG-AAGUAGUUGUUUCUUGACGAGAGAAUAUGUAUAUCUUGG 258
dISFV-NC_024017.1  AGCGCCUG--AAG-A--GAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU 258
dISFV-NC_016997.1  AGCGCCAU--AAG-AAACGGAGUUAGUAUGGAAAAAACAUAAGUA-----UCU 258
dISFV-NC_017086.1  CGCGCCGG--AAA-AGGUAGUUUGGAGGUUGUAAAACAACAAGUA-----UCU 258
dISFV-NC_027999.1  UAUGCCUG--AAACAUUCUGUGGUUUGAUGUAGUAAAACAACCAGU-----UUCU 258
TBFV-NC_027709.1   AUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAAACACCCG--------CU 258
TBFV-NC_001672.1   GUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACACCCG--------CU 258
TBFV-NC_001809.1   AUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACACCCG--------CU 258
TBFV-NC_005062.1   GCAGUCUAACAAGGAGGUGAUGUGUGACUCGGAACAACACCCG--------CU 258
TBFV-NC_003690.1   AUAGUCUGAAAAGGAGGUGAUGUGUGUCUCGGAAAAACACCCG--------CU 258
TBFV-NC_003687.1   AUAGCCUGACAAGGAGAUGGUGUGUGACUCGGAAAAACACCCG--------CU 258
                   40.......250.......260.......270.......280.......290.
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