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                                 .................................(((((((((((((((((((((((
dISFV-NC_027999.1  GGAU--UAAUUGUUGUUUGAGAGAACGCGAGAAGUACUA----CUAUGGUUGAGAAAAC  54
dISFV-NC_016997.1  AAAC--AUAACGCAUUAGAAUAGGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGU  54
dISFV-NC_017086.1  ------AUAACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG--------GC  45
dISFV-NC_040610.1  --GA--ACAAUUUUAAAGAAGUCAA-AACUAGAAAUUUAAUU-UCAAUGGU------UAG  48
dISFV-NC_024017.1  UCGC--GCAUUUUGAUAGAGUU----GACUAGAUAACGAAUUAUCGAUGGU------GAC  48
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_027999.1  UAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUAC--------- 105
dISFV-NC_016997.1  CUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC---- 110
dISFV-NC_017086.1  GAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCG 105
dISFV-NC_040610.1  ACAACUGGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUUACCACCCG 108
dISFV-NC_024017.1  UAAAUCAAGAAAAGCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAGCG 108
TBFV-NC_027709.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA---- 110
TBFV-NC_001672.1   CAUCCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA---- 110
TBFV-NC_001809.1   CAUCCUGAAAGGCAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA---- 110
TBFV-NC_005062.1   CAUUCUGAAAGGAAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA---- 110
TBFV-NC_003690.1   CGUUCUAAAAGGAAAGGGGGGGCGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA--- 110
TBFV-NC_003687.1   CACUUCUAAAGGAAAGGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAA 108
                   0........70........80........90........100.......110.......1
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                   ..................((((((((((((.....))))))..................
dISFV-NC_027999.1  UGGAANNNNNNNUGGCGA-ACCCCAUUGGAAAUUUA----GCUUCAUAUUGA---CACCA 157
dISFV-NC_016997.1  -----NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGA---CACCU 160
dISFV-NC_017086.1  CGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGA---CACUU 160
dISFV-NC_040610.1  CU---NNNNNNN--------ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGA---CACCA 154
dISFV-NC_024017.1  UG---NNNNNNN--------CCCCACUGCUGAGAG-CAAAAUAGCAUAUUGA---CACUU 153
TBFV-NC_027709.1   -----NNNNNNNC-------UCUCUUCAGGAUUUUUCCUCCUCCUAUA--UUCCCC 157
TBFV-NC_001672.1   -----NNNNNNNC-------UCUCUUCAGGAUUUUUCCUCCUCCUAUACAAAA-UUCCCC 157
TBFV-NC_001809.1   -----NNNNNNN--------UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUCCCC 157
TBFV-NC_005062.1   -----NNNNNNNC-------UCUCUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UUCCCC 157
TBFV-NC_003690.1   -----NNNNNNNC-------UCUCUUCAGGGAUAUUUCCUCCUCCUAUACCAAA-UUCCCC 157
TBFV-NC_003687.1   GU---NNNNNNNC------U-UCUCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUCCCC 157
                   20.......130.......140.......150.......160.......170.......1
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                   .........))))))))))))))))))))))).)))))))))...(((((((((.(((...
dISFV-NC_027999.1  GGGAAAAGACCGCAGAAACUUCUUGCUUACUCGCAGCAACCACG--AAAGGCACAGUAU 215
dISFV-NC_016997.1  AGG-AAAGACUGCAGCAUAC-CUUUGGUUCUUCGACACUAACUCCU--AAUGGCACAGAGC 216
dISFV-NC_017086.1  GGG-AAAGACCAGAGGUACUCGCGUCGCAACCGCCACCAG-ACUA--CACGGCACAGAGC 216
dISFV-NC_040610.1  GGG-AUAGACCGCAGAAAUACACAAUUUCUGGGCACUAAACAACUACAA--GGCACAGAGC 211
dISFV-NC_024017.1  GGGAAGAGACCACAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAGGGCACAGAGC 213
TBFV-NC_027709.1   CUCGACAGAGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAGUAUA 216
TBFV-NC_001672.1   CUCGGUAGAGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAGGUA 216
TBFV-NC_001809.1   CUCGACAGAGGGCGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAGGUA 216
TBFV-NC_005062.1   CUCAUCUGAGAGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCCA-GACACAGGCA 216
TBFV-NC_003690.1   CUCGUCAGAGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCAUCACCUA-GACACAGUAU 216
TBFV-NC_003687.1   CAGGAAACUGGGGGGGCGGUUCUUGUUCUCCCUGAGCCACCACCAUCCA-GGCACAGUAU 216
                   80.......190.......200.......210.......220.......230.......2
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                   ))))))........))))))))
dISFV-NC_027999.1  GCCUGAAACAUUCGGUGGUUG----AUGUAGUAAACAACCAGUUUCU 258
dISFV-NC_016997.1  GCCAUAAGAAACGGAGUUAGU----AUGGAA-AAAACAUAAGUAUCU 258
dISFV-NC_017086.1  GCCGGAAAAGGUAGUUUGGAG----GUUGUA-AAACAACAAGUAUCU 258
dISFV-NC_040610.1  GCUUCGAGAAGUAGUUGUUUCUUGACGAGAGAAUAUGUAUAUCUUGG 258
dISFV-NC_024017.1  GCCGAAGA--GAGCGUGUUAUUGUUUAUAAUUUACAAAAAUAACAUAUCU 258
TBFV-NC_027709.1   GUCUG-ACAAGGAGGUGAUGCGUGACUCGGAAAAACACCC----GCU 258
TBFV-NC_001672.1   GUCUG-ACAAGGAGGUGAUGUGUGACUCGGAAAAACACCC----GCU 258
TBFV-NC_001809.1   GUCUG-ACAAGGAGGUGAUGUGUGACUCGGAAAAACACCC----GCU 258
TBFV-NC_005062.1   GUCUA-ACAAGGAGGUGAUGUGUGACUCGGAACAACACCC----GCU 258
TBFV-NC_003690.1   GUCUG-AAAGGAGGUGAUGUGUGUCUCGGAAAAACACCC----GCU 258
TBFV-NC_003687.1   GCCUG-ACAAGGAGAGUGAUGUGUGACUCGGAAAAACACCC----GCU 258
                   40.......250.......260.......270.......280.....
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