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                    .................(((((..(((((((((((.....................
dISFV-NC_040610.1   -------GAACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAAUU-UCAAUGGUU------ 46
dISFV-NC_024017.1   UC-----GCGCAUUUUGAUAGAGUU---GACUAGAUAACGAAUUAUCGAUGGUG------ 46
dISFV-NC_027999.1   GGAUUAAUUGUUGUUUGAGAGAAC----AGCGAGAAGUACUACU----AUGGUUGAGAAA 52
dISFV-NC_016997.1   AAACAUAACGCAUUAGAAUAGGUUU--------GAAAGAGUUGAAUCAAUGAGAUCAAUG 52
dISFV-NC_017086.1   ----AUAACACGUUUGAAUAGGUUU----AUUGUGAAUCGAAAUACCGAUG--------- 43
TBFV-NC_027709.1    GUGUUGAGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGG------AUGGGCAGGAAG 52
TBFV-NC_001672.1    GUGUUGAGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGG------AUGGGCAAGAAG 52
TBFV-NC_001809.1    GUGUUGAGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGG------AUGGGCAGGAAG 52
TBFV-NC_005062.1    GUGUUGAGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGG------AUGGCCGGGAAG 52
TBFV-NC_003690.1    GUGUUGAGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGA------AUGGCCGGGAAG 52
TBFV-NC_003687.1    ------AGAAAAAGACAGCUUAGG--AGAACAAGAGCUGGGAGUGGUUAUG------AUG 46
                    .........10........20........30........40........50........6
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                    .................(((((.....)))))....((.((.((....)).)))).....
dISFV-NC_040610.1   AGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUUACCACC 106
dISFV-NC_024017.1   ACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAG 106
dISFV-NC_027999.1   ACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUACUG----- 107
dISFV-NC_016997.1   GUCUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUUAAAACGUAAGCUGC-- 110
dISFV-NC_017086.1   GCGAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCU 103
TBFV-NC_027709.1    UCCAUCCUGAAAGGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA-- 110
TBFV-NC_001672.1    GCCAUCCUGAAAGGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA-- 110
TBFV-NC_001809.1    ACCAUCCUGAAAGGCAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA-- 110
TBFV-NC_005062.1    GCCAUUCUGAAAGGAAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA-- 110
TBFV-NC_003690.1    GCCGUUCUAAAAGGAAAGGGGGCGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA-- 110
TBFV-NC_003687.1    ACCACUUCUAAAGGAAAGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAU 106
                    0........70........80........90........100.......110.......1
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                    ......................(((((((.........))))).................
dISFV-NC_040610.1   CGCU---NNNNNNNACCC---CCUCAUUAUUUA-----UAUAACAGCAUAUUGACACCAG 155
dISFV-NC_024017.1   CGUG---NNNNNNNCCCC---ACUGCUGAGAG------CAAAAUAGCAUAUUGACACUUG 154
dISFV-NC_027999.1   -GAA---NNNNNNNUGGCGAACCCCAUUGGAAA-----UUUAGCUUCAUAUUGACACCAG 158
dISFV-NC_016997.1   -------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUA 161
dISFV-NC_017086.1   CGCGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGACACUUG 161
TBFV-NC_027709.1    -------NNNNNNNCUCU----CUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCC 158
TBFV-NC_001672.1    -------NNNNNNNCUCU----CUUCAGGAUUUUUCCUCCUCCUAUACAAAA-UUCCCCC 158
TBFV-NC_001809.1    -------NNNNNNN-UCU----CUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUUCCCCC 158
TBFV-NC_005062.1    -------NNNNNNNCUCU----CUUCAGGAAAUUUCCUCCUCCUAUACCAAA-UUCCCCC 158
TBFV-NC_003690.1    -------NNNNNNNCUCU----CUUCAGGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCC 158
TBFV-NC_003687.1    AAGU---NNNNNNNCU-U----CCUCGGGAUUUUUCCGCCUCCUAUACUAAAUUUCCCCC 158
                    20.......130.......140.......150.......160.......170.......1
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                    ..........)))).))))))))))))).)))))))....(((((((((((.......)
dISFV-NC_040610.1   GG-AUAGACCGGAGAAAU-ACACAAUUUCUGGGCACUAAACAACUACA-AGCACAGAGCG 212
dISFV-NC_024017.1   GGAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACAGAGCG 214
dISFV-NC_027999.1   GGAAAAGACCGGAGAAACUUCUUGCUUACUCGCAGC-AACCACCGAAA-GGCACAGUAUG 216
dISFV-NC_016997.1   GGAA-AGACUGGAGAUAC-CUUUGGUUCUUCGA--CACUAACUCCUAAUGGCACAGAGCG 217
dISFV-NC_017086.1   GGAA-AGACCAGAGGUACUCGCUGAUUCACCGC--CACCAG-ACUACACGGCACAGCGCG 217
TBFV-NC_027709.1    UCGACAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CAUCACCCAGACACAGAUAG 217
TBFV-NC_001672.1    UCGGUAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CAUCACCCAGACACAGGUAG 217
TBFV-NC_001809.1    UCGACAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CAUCACCCAGACACAGAUAG 217
TBFV-NC_005062.1    UCAUCUGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CAUCACCCAGACACAGGCAG 217
TBFV-NC_003690.1    UCGUCAGAGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CAUCACCUAGACACAGAUAG 217
TBFV-NC_003687.1    AGGAAACUGGGGGGGCGGUUCUUGUUCUCCCCUGAGCCAC-CACCAUCCAGGCACAGAUAG 217
                    80.......190.......200.......210.......220.......230.......2
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                    ))))......)))))))).............................
dISFV-NC_040610.1   CUUC--GAGAAGUAGUUGUUUCUUGACGAGAGAAUAU-GUAUAUCUUGG 258
dISFV-NC_024017.1   CCUG--AAGAGAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU--- 258
dISFV-NC_027999.1   CCUG--AAACAUUCGGUGGUUGAUGUAGUAAACAACCAGUUUCU----- 258
dISFV-NC_016997.1   CCAU--AAGAAACGGA-GUUAGUAUGGAAAAAACAUAAGUAUCU----- 258
dISFV-NC_017086.1   CCGG--AAAAGGUAGU-UUGGAGGUUGUAAAACAACAAGUAUCU----- 258
TBFV-NC_027709.1    UCUGACAAGGAGGUGAUGCGUGACUCGGAAAAACACCCGCU-------- 258
TBFV-NC_001672.1    UCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACACCCGCU-------- 258
TBFV-NC_001809.1    UCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACACCCGCU-------- 258
TBFV-NC_005062.1    UCUAACAAGGAGGUGAUGUGUGACUCGGAACAACACCCGCU-------- 258
TBFV-NC_003690.1    UCUGAAAAGGAGGUGAUGCGUGUCUCGGAAAAACACCCGCU-------- 258
TBFV-NC_003687.1    CCUGACAAGGAGAUGGUGUGUGACUCGGAAAAACACCCGCU-------- 258
                    40.......250.......260.......270.......280.......
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