

dISFV-NC_016997.1	AAAC--AUAACGCAUUAAGAAUAGGUUU---GAAAGAGUUGAAUCAAUGAGAUCAAUGGU	54
dISFV-NC_017086.1	-----AUAACACGUUUGAAUAGGUUUUAUUGUGAAUUCGAAUACCGAUG-----GC	45
dISFV-NC_040610.1	--GA--ACAAUUUUAAAAGAAUGCAAAC--UAGAAAUUU--AAUUUCAAUUG-----GU	45
dISFV-NC_024017.1	UCGC--GCAUUUUGAUAGAGUU---GAC--UAGAAUACGAAUUUAUCGAUG-----GU	45
dISFV-NC_027999.1	GGAU--UAAUUGUUGUUUGAGAGAACAGCGAGAAAGUACUA----CUAUG-----GUUGAGAA	51
TBFV-NC_001809.1	GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGG-----AUGGGCAGGAAGA	54
TBFV-NC_005062.1	GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGG-----AUGGCCGGGAAGGC	54
TBFV-NC_027709.1	GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGG-----AUGGGCAGGAAGU	54
TBFV-NC_001672.1	GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGG-----AUGGUCAAGAAGGC	54
TBFV-NC_003690.1	GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGA-----AUGGCCGGGAAGGC	54
TBFV-NC_003687.1	-----AGAAAAAGACAGCUUAGGAGAAACAAAGCUGGGGAGUGGUUAUG-----AUGAC	48
10.....20.....30.....40.....50.....6	

dISFV-NC_016997.1 CUU CGAGCUAGGAGGCCCUU ---UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC- 110
 dISFV-NC_017086.1 GAUUAACCCCAAGAAGCCG ---AAGACGGGCUAUCGAUAUAGUGAGACGUCGCGUUGCC 102
 dISFV-NC_040610.1 UAGACAACUUGGAAAGACCUUUGCAAGCGGUCUGUCAAUAUGCGAAACCGGA--UUAC 103
 dISFV-NC_024017.1 GACUAAAUAAGAAAGCCCGCAAAGCGGGCUGUUAUAUAGCUAACACGCA--UCACC 103
 dISFV-NC_027999.1 AACUAGAGUAGGAUCUCGGUUA AAAACCCGAGGUGUCAUAUAUGAUUAGGCGUA----- 104
 TBFV-NC_001809.1 CAUUCUGAAAGGCAAGGGGCGCGUCCCC--CUCGACGAGUGUCGAAAGAGA---CCGC 108
 TBFV-NC_005062.1 CAUUCUGAAAGGAAAGGGCGCGUCCCC--CUCGACGAGUGUCGAAAGAGA---CCGC 108
 TBFV-NC_027709.1 CAUUCUGAAAGGUUAGGGGCGCGUCCCC--CUCGACGAGUGUCGAAAGAGA---CCGC 108
 TBFV-NC_001672.1 CAUUCUGAAAGGUUAGGGGCGCGUCCCC--CUCGACGAGUGUCGAAAGAGA---CCGC 108
 TBFV-NC_003690.1 CGUUCUAAAAGGAAAGGGGCGGGUCCCC--CUCGACGAGCUGUCGAAAGUGG---CCCC 108
 TBFV-NC_003687.1 CACUUCUAAAGGAAAGGGGCGCGUCCCC--CUAGGCGCAAGCUUAAAGUGA---CCGC 102

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dISFV-NC_016997.1 -----NNNNNNNN--AGGAAACCCCCCAAAAAUGUAUAGGCUAUUAUAUCGA--CA 157
dISFV-NC_017086.1 UCGCGUCUNNNNNNNNAGAGGAGACCCCCCAAAAAUG--AAGGGCACUAUAUCGA--CA 157
dISFV-NC_040610.1 ACCCGCU-NNNNNNNN-----ACCCCCUUAUUUAUUAUAAACAGCAUAUUGA--CA 151
dISFV-NC_024017.1 AAGCGUG-NNNNNNNN-----CCCCACU-UCUGAGAGCAAAAUAGCAUAUUGA--CA 150
dISFV-NC_027999.1 --CUGGAANNNNNNNNNUGGCGAACCCCAUUGGAAAUU-----UAGCUUCAUAUUGA--CA 154
TBFV-NC_001809.1 GA-----NNNNNNNN-----UCUCUUCAGGGUUUUUCCUCCUCCUAUACCAAAUUU 154
TBFV-NC_005062.1 UA-----NNNNNNNN-----CUCUCUUCAGGAAAUUUUCCUCCUCCUAUACCAAA-UUC 154
TBFV-NC_027709.1 AA-----NNNNNNNN-----CUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUC 154
TBFV-NC_001672.1 AA-----NNNNNNNN-----CUCUCUUCAGGAUUUUUCCUCCUCCUAUACCAAA-UUC 154
TBFV-NC_003690.1 AA-----NNNNNNNN-----CUCUCUUCAGGAUAUUUCCUCCUCCUAUACCAAA-UUC 154
TBFV-NC_003687.1 AAAUAAGUNNNNNNNN-----CU-UCCUCGGGAUUUUUCCGCCUCCUAUACAAAUUUC 154
20.....130.....140.....150.....160.....170.....1

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dISFV-NC_016997.1 CUAGGAA-AGACUGGAGAUAC-CUUGGUUCUUGGACAUAAUCCUA--AUGGCACAG 213
 dISFV-NC_017086.1 UUGGGAA-AGACCAAGGUACUCUCUGAUUCCAGGCCACAG-ACUAC--ACGGCACAG 213
 dISFV-NC_040610.1 CAGGG-AUAGACCGGAGAAU-ACAUAUUUCUGGGCACUAAACAACUACA-AGCACAG 208
 dISFV-NC_024017.1 UUGGGGAAGACCAAGAGAUAGAUAUUGUUGUUGUCAAUGUGACAACACGCAUUGGCACAG 210
 dISFV-NC_027999.1 CAGGGAAAAGACCGGAGAAAUUUGUUGCUUACUCG-CAGCAACCACCGAAA-GGCACAG 212
 TBFV-NC_001809.1 CCCUCGACAGAGGGGGGCGGUUUGUUCUCUCUGAGCACCACUACCCA-GACACAG 213
 TBFV-NC_005062.1 CCCUCAUCUGAGGGGGGCGGUUUGUUCUCUCUGAGCCACCAUCACCCA-GACACAG 213
 TBFV-NC_027709.1 CCCUCGACAGAGGGGGGCGGUUUGUUCUCUCUGAGCACCACUACCCA-GACACAG 213
 TBFV-NC_001672.1 CCCUCGGUAGAGGGGGGCGGUUUGUUCUCUCUGAGCCACCAUCACCCA-GACACAG 213
 TBFV-NC_003690.1 CCCUCGUCAGAGGGGGGCGGUUUGUUCUCUCUGAGCACCACUACCCA-GACACAG 213
 TBFV-NC_003687.1 CCCAGGAAACUGGGGGGCGGUUUGUUCUCUCUGAGCCACCAUCAUCCA-GGCACAG 213

dISFV-NC_016997.1	AGCGCCAUA-AGAA--ACGAGUUAGUAUGGAAAAACAUA-AGUAUCU---	258
dISFV-NC_017086.1	CGCGCCGGA-AAAG--GUAGUUUGGAGGUUGUAAAAACAACAA-AGUAUCU---	258
dISFV-NC_040610.1	AGCGCUUCG-AGAAGUAGUUGUUUCUUGACGAGAGAAUAU-AGUAUAUCUUGG	258
dISFV-NC_024017.1	AGCGCCUGA-AGAGAGCUUGUUAUAAUUUACAAAAAUAACAUAUCU---	258
dISFV-NC_027999.1	UAUGCCUGA-AACAUCGGGUUU--GAUGUAGUAAACAACCGAUUCU---	258
TBFV-NC_001809.1	AUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACAC-----CCGCU	258
TBFV-NC_005062.1	GCAGUCUAAACAAGGAGGUGAUGUGUGACUCGGAAACAAC-----CCGCU	258
TBFV-NC_027709.1	AUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAAACAC-----CCGCU	258
TBFV-NC_001672.1	GUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAAACAC-----CCGCU	258
TBFV-NC_003690.1	AUAGUCUGAAAAAGGAGGUGAUGCGUGUCUCGGAAAAACAC-----CCGCU	258
TBFV-NC_003687.1	AUAGCCUGACAAGGAGAUGGUGUGACUCGGAAAAACAC-----CCGCU	258
	40.....250.....260.....270.....280.....290	