

MBFV-NC_002031.1 -----UAAUCGCUUCGUGAGCGAUUAGCAGAGAACUGAC--CAGAACAUGUC----- 45
MBFV-NC_012735.1 -----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUCCCAUUGGCAACAAA 51
MBFV-NC_008719.1 -----AUUUCUACUGAAAGG--GUAGAGAAAGGAGUUUUGCUUCUACUUGGCAACAAG 51
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAAACCC 51
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA--CC 50
MBFV-NC_012533.1 AACAGUUUU-----CGUGUUGAGUUAAGAAACAGAUUUCUGG--AAUGAAAA--CC 49
MBFV-NC_002640.1 -----UCUA--ACAGUUUGUUUGAAUAGAGAGCAG--AUCUCUGGAAAAAUG---AA--C 46
MBFV-NC_001477.1 ACGUAGUUCUA--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUGAACAA--C 49
MBFV-NC_001475.2 ACGUAGUGCUG--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUGAACAA--C 49
MBFV-NC_001474.2 -CGUAGUUCUA--ACAGUUU--UUUAAUUAGAGAGCAG--AUCUCUG-----AUGAAUA--C 49
MBFV-NC_009028.2 U--U-----UAACACAGUCUGAAUAGUUUUUU--AGCAAGGGA--UUUCC--CAUGUCU---AA 48
MBFV-NC_040776.1 U--U-----UAACACAGUCUGAAUAGUUUUUU--AGCAGAGGU--ACUAC--CAUGUCU---AA 48
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUGAAAAAC--CAUGUCU---AA 48
MBFV-NC_007580.2 ---U-----AACUUGACUGCGAACAGUUUUUU--AGCAGGGAAUUACCC--AAUGUCU---AA 48
MBFV-NC_001437.1 ---U-----UAGUGCAGUUUAAACAGUUUUUU--AGAACCGGAAGAUAA--CAUGACU---AA 48
MBFV-NC_006551.1 ---U-----UAACACAGUGCGCGGAGUUUCUU--UGAGCGUUGAUUUUC--AAUGUCU---AA 48
MBFV-NC_001563.2 A--U-----UAACACAGUGCGAGCUGUUUCUU--GGCACG--AAGAUUCG--GAUGUCU---AA 48
MBFV-NC_009942.1 A--U-----UAACACAGUGCGAGCUGUUUCUU--AGCACG--AAGAUUCG--GAUGUCU---AA 48
MBFV-NC_000943.1 -----UAACGCGGUUGAACAGUUUUUUGGAGCUUUUGAUUUCA--AAUGUCU---AA 48
MBFV-NC_012534.1 CCGU-----AAACACAGUUUGAACGUUUUUUGGAGA---GAGAUUAU--UAUGUUUAACAA 51
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAACGUUUUUUGGAUA---GAGACAAC--UAUGUCUAACAA 51
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAAGUUUUUUGAAC---AG---C--AAUGACC---AA 48
MBFV-NC_032088.2 UC-----UAUAACUCAGUGGAACAAGUUUUUGAUC---AGUGGAU--AAUGACU---AA 48
MBFV-NC_009026.2 -----AGAAAAGUUUUCGAGGAAGACGAGAAGCGAAUUCUGAAUGGCU---AG 48
MBFV-NC_018705.3 CCGU-----AAACACAGUUUGAACGUUUUUUGGAGAGAGACUACU-----AUGUCAACAA 51
TBFV-NC_027709.1 GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAA 51
TBFV-NC_001672.1 GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGUCAAGAA 51
TBFV-NC_001809.1 GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAA 51
TBFV-NC_005062.1 GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCGGGAA 51
TBFV-NC_003690.1 GUGUUGAGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGA-----AUGGCCGGGAA 51
TBFV-NC_003687.1 -----AGAAAAGACAGCUUAGGAGAACAAGAGCUGGGAGUGGU---UAUG-----AU 45
.....10.....20.....30.....40.....50.....6



MBFV-NC_002031.1 -----UGGUCGUAAAGCUCAGGGGAAAAACCCUGGGCGUCAAUATGGU-A--CGACGAGGAG 98
MBFV-NC_012735.1 GG--GGAUGAAUAAGUCUCGGGUCUGCAUCCCGAGGCGUCAUAUUGGUG--CGGCUAGA-- 106
MBFV-NC_008719.1 AG-GAGCAAGCAAGUCUCGGGUCACGACCCCGGGCGUCAUAUUGGUGA--CGGCCAAG-- 106
MBFV-NC_012532.1 CA-AAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAUAUUGCUAAAACGCGGAGUAG 104
MBFV-NC_035889.1 CA-AAAAAGAAAU--CC---GGAGGAUUCGGAUUGUCAUAUUGCUAAAACGCGGAGUAG 104
MBFV-NC_012533.1 CA-AAAAAGG-----GCCGCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGAU 98
MBFV-NC_002640.1 CA-ACGAAAAAAG--GU---G--GUUAGACCACCUUUCAUAUUGCUGAAACGCGAGAGAA 98
MBFV-NC_001477.1 CA-ACGGA AAAAG--AC---G--GGUCGACCGUCUUUCAUAUUGCUGAAACGCGCGAGAA 101
MBFV-NC_001475.2 CA-ACGGAAGAAG--AC---G--GGAAAAACGCUUAUCAUAUUGCUGAAACGCGGUGAGAA 101
MBFV-NC_001474.2 CA-ACGGA AAAAG--GC---G--AAAAACACGCCUUUCAUAUUGCUGAAACGCGAGAGAA 101
MBFV-NC_009028.2 AA-AACCAGGAAAGUCC--GCUGCAAAGCGGACUGUCAUAUUGCUAAAACGCCUGGCUA 104
MBFV-NC_040776.1 AA-AACCAGGAGGACCC--GCCGGAAGGCGGGUUGUCAUAUUGCUGAAACGCCACGCCA 104
MBFV-NC_034151.1 AA-AACCAGGAGGACCC--GGAAAGAACCAGGUCGUUAUAUGCUGAAGCGCACAUCC 104
MBFV-NC_007580.2 AA-AACCAGGAAACCC--GGUAGAAACCGGGUUGUCAUAUUGCUAAAACGCGGCGUAU 104
MBFV-NC_001437.1 AA-AACCAGGAGGGGCC--GGUAAAAACCGGGCUUAUCAUAUUGCUGAAACGCGGCCUAC 104
MBFV-NC_006551.1 GA-AACCAGGAGGGCCC--GGAAGAACCAGGGCAUCAUAUUGCUGAAACGCGGCAUAC 104
MBFV-NC_001563.2 GA-AACCAGGAGGGCCC--GGUAAAAACCGGGCUUCAUAUUGCUGAAACGCGGUUAUGC 104
MBFV-NC_009942.1 GA-AACCAGGAGGGCCC--GGCAAGAGCCGGGCUGUCAUAUUGCUAAAACGCGGAUUGC 104
MBFV-NC_000943.1 AA-AACCAGGAGGACCC--GGGAAGCCCCGGGUCGUCAAUUGCUGAAACGCGGCAUAC 104
MBFV-NC_012534.1 AA-AACCAGGAAACCC--GGGGCAACCCGGGUUGUCAUAUUGCUGAAGCGCGGUGCAU 107
MBFV-NC_015843.2 AA-AACCAGGAAGACCC--GGCUCAGGCCGGGUUGUCAUAUUGCUAAAGCGCGGAACGU 107
MBFV-NC_009029.2 AA-AACCAGGAAGGCC--GGUAGAAAUCCGGGCCGUCAAUUGCUAAGCGCGGUGCGU 104
MBFV-NC_032088.2 AA-AACCAGGAAGUCC--GGCAUUCGUCGACUGUCAUAUUGCUAAGCGCGGAGCAU 104
MBFV-NC_009026.2 AA-AACCAGGAAGACCC--GGAGGAACCCGGGUCGUCAAUUGCUGAAGCGUACGGCGG 104
MBFV-NC_018705.3 AA-AACCAGGAAGACCC--GGGGCAACCCGGGUUGUCAUAUUGCUAAGCGCGGCGCAU 107
TBFV-NC_027709.1 GUCCAUCCUGAAAGGUAAGGGGGGGCGGUC-----CCCCGACGAGUGUCGAAAGAGACCGCAA 110
TBFV-NC_001672.1 GGCCAUCCUGAAAGGUAAGGGGGGGCGGUC-----CCCUCGACGAGUGUCGAAAGAGACCGCAA 110
TBFV-NC_001809.1 GACCAUCCUGAAAGGCAAGGGGGGGCGGUC-----CCCUCGACGAGUGUCGAAAGAGACCGCGA 110
TBFV-NC_005062.1 GGCCAUUCUGAAAGGAAAGGGGGGGCGGUC-----CCCUCGACGAGUGUCGAAAGAGACCGCUA 110
TBFV-NC_003690.1 GGCCGUUCUAAAGGAAAGGGGGGGGGGUC-----CCCUCGACGAGUGUCGAAAGUGGCCCAA 110
TBFV-NC_003687.1 GACCACUUCUAAAGGAAAGGGGGGGCGGUC-----CCCUGAGGCGCAAGCUUAAAGUGACCGCAA 105
0.....70.....80.....90.....100.....110.....1



MBFV-NC_002031.1 UUCGCUCCUUGUNNNNNNN-----GAGACCCUCCAGGGAACAAAUAGUGGGGA 145
MBFV-NC_012735.1 -----GUGANNNNNNN-----C-CCAUUCCUCCCGAGCACACAUAG--GGA 146
MBFV-NC_008719.1 -----GCCANNNNNNN-----GAGGAGACCCUCCCGAGCAC-UAUAGC--GGA 146
MBFV-NC_012532.1 CCGGUG-----NNNNNNNN-----GGUAGAGGAGACCCCCCGGAAAACGCAAAACA 150
MBFV-NC_035889.1 CCGGUG-----NNNNNNNN-----GUUAGAGGAGACCCCCCGGAAAACGCAAAACA 149
MBFV-NC_012533.1 CUAGCGCUGGCCNNNNNNNN-----GAGACCCCGCGCCCAUAAACCAACUAAACAA 150
MBFV-NC_002640.1 ACCGCGUAUCAANNNNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCAACA--CAAAAACA 154
MBFV-NC_001477.1 ACCGCGUGU---NNNNNNNN---UAGAGGUUAGAGGAGACCCCCCGCACAAACAACAAACA 154
MBFV-NC_001475.2 ACCGUGUGU---NNNNNNNN---ACUAGAGGUUAGAGGAGACCCCCCG--CAAAUAAAAACA 154
MBFV-NC_001474.2 ACCGCGUGU---NNNNNNNN---CUAGAGGUUAGAGGAGACCCCCCG--CGAAAACAAAAACA 154
MBFV-NC_009028.2 GCGUGU-----NNNNNNNN---GUGGAGACCCUCAGCAC--CAAGCGCGAA--ACAAACA 152
MBFV-NC_040776.1 GCGUUU-----NNNNNNNN---A-GGAGACCCCUUGGCACACAAGCGCAAA--GACAAACA 153
MBFV-NC_034151.1 GCGCUA-----NNNNNNNN-----AGACCCUUGCGAGUGAA--CGCAAA--UAAAAACA 149
MBFV-NC_007580.2 CCCGCG-----NNNNNNNNUAGAGGAGACCCCUUGCCGUUAA-----CGC--AAACAACA 152
MBFV-NC_001437.1 CCCGCG-----NNNNNNNN--UAGAGGAGACCCCC--CAU---UGCAUC-----AAACA 146
MBFV-NC_006551.1 CCCGCG-----NNNNNNNN-----ACCCG---CAUCAUAAGCAUCA--AAAAACA 145
MBFV-NC_001563.2 CCCGCG-----NNNNNNNN---GAGGAGACCCCGUGCCA--AAAACACCA--A--AAGAAACA 152
MBFV-NC_009942.1 CCCGCG-----NNNNNNNN---GGAGACCCCGUGCCACAAAACACCACA--ACAAAACA 152
MBFV-NC_000943.1 CCCGCG-----NNNNNNNN-----GACCCCA---CUCUCAAAGCAUC--AAACAACA 146
MBFV-NC_012534.1 CCC-----NNNNNNNN-----GGAGACCCUUGCGAGUUAACACCAAA--ACAAAACA 152
MBFV-NC_015843.2 CCC-----NNNNNNNN---AGGAGACCCCUUGCGAGUGAACACCACA--A--GAAACA 152
MBFV-NC_009029.2 CCCGCG-----NNNNNNNNAGAGG--AGACCCUCCCAACAAGAGCGCAAAACAAAAACA 156
MBFV-NC_032088.2 CCCGCG-----NNNNNNNNAGAGGUUAGAGGAGACCCCAACAAGCGCAAA--ACAAAACA 157
MBFV-NC_009026.2 CCAACG-----NNNNNNNNCCCCUUGG-----CAAAACAGUUAACGCACCAAAAGAAACA 153
MBFV-NC_018705.3 CCC-----NNNNNNNN-----CCCCGCGGAAAUCAAAGCAAGGCCCAACCUG 148
TBFV-NC_027709.1 -----NNNNNNNNCUCUC-----UUCAGGAUUUUUCCUCCUCCUAUACCAA--U 152
TBFV-NC_001672.1 -----NNNNNNNNCUCUC-----UUCAGGAUUUUUCCUCCUCCUAUACCAA--U 152
TBFV-NC_001809.1 -----NNNNNNNN--UCUC-----UUCAGGGUUUUUCCUCCUCCUAUACCAAU 152
TBFV-NC_005062.1 -----NNNNNNNNCUCUC-----UUCAGGAUUUUUCCUCCUCCUAUACCAA--U 152
TBFV-NC_003690.1 -----NNNNNNNNCUCUC-----UUCAGGAUUUUUCCUCCUCCUAUACCAA--U 152
TBFV-NC_003687.1 UAAGU-----NNNNNNNNCU--UC-----CUCGGGAUUUUUCCGCCUCCUAUACUAAAU 152
20.....130.....140.....150.....160.....170.....1



MBFV-NC_002031.1)))))).)))))(((((..... 201
MBFV-NC_012735.1 CCUAU--UUGACGCGCAGGGA--AAGACCGGAGUGGUUCUGCUUUUC--CUCCAGAGGUCU 201
MBFV-NC_008719.1 CCUAU--UUGACGCGCAGGGA--AAGACCGGAGACACUCCUUGAUUUCACCUUUCUGGCC 203
MBFV-NC_012532.1 CCUAU--UUGACGCGCAGGGA--AAGACCGGAGACACUCCUUGAUUUCACCUUUCUACCC 203
MBFV-NC_035889.1 GCAUA--UUGACG--UGGGA--AAGACCAG--AGACUCCAUGAGUUUCCACCACGCUGGCCG 204
MBFV-NC_012533.1 GCAUA--UUGACG--CUGGGA--AAGACCAG--AGACUCCAUGAGUUUCCACCACGCUGGCCG 204
MBFV-NC_002640.1 GCAUA--UUGACACCCUGGGA--AAGACCGG--AGACUCUGUUGGUUUUCCCAACACGACCC 206
MBFV-NC_001477.1 GCAUA--UUGACG--CUGGGA--AAGACCAG--AGAUCUCUGCU--GUCUCUGCAACAUCA--AU 206
MBFV-NC_001475.2 GCAUA--UUGACG--CUGGGA--GAGACCAG--AGAUCUCUGCU--GUCUCUACAGCAUCA--UU 206
MBFV-NC_001474.2 GCAUA--UUGACG--CUGGGA--GAGACCAG--AGAUCUCUGCU--GUCUCUCAGCAUCA--UU 206
MBFV-NC_009028.2 GCAUA--UUGACG--CUGGGA--AAGACCAG--AGAUCUCUGCU--GUCUCUCAGCAUCA--UU 206
MBFV-NC_040776.1 GCAUA--UUGACGCCUGGGA--AAGACCGGGAGAUCCUCUGCU---UCCAUCACCAGCCA 206
MBFV-NC_034151.1 GCAUA--UUGACACCCUGGGA--AAGACCGGGAGACCCUCUGCU---CUGCAACACCAGCCA 207
MBFV-NC_007580.2 GCAUA--UUGACACCCUGGGA--UAGACUGGGAGACCCUCUGCU--UAUUUACACACCAGCCA 205
MBFV-NC_001437.1 GCAUA--UUGACACCCUGGAA--AGAC--AGGAGAUCCCUUGCUUU--UUCAACACCAGCCA 205
MBFV-NC_006551.1 GCAUA--UUGACACCCUGGGAA--UAGACUGGGAGAUUUCUGCUCUAUUCUACAACAGCUA 204
MBFV-NC_001563.2 GCAUA--UUGACACCCUGGGA--UAGACUAGGGGAUCUUCUGCUCUG---CACAACCAGCCA 206
MBFV-NC_009942.1 GCAUA--UUGACACCCUGGGA--UAGACUAGGAGAUUUCUGCUCUG---CACAACCAGCCA 206
MBFV-NC_000943.1 GCAUA--UUGACACCCUGGGA--AAGACUAGGAGAUUUCUGCUCUAUUCUACAACUAGUCA 204
MBFV-NC_012534.1 GCAUA--UUGACACCCUGGGA--GAGACUAGGAGAUUUCUGGUCUA--ACAACACCAGCCA 207
MBFV-NC_015843.2 GCAUA--UUGACACCCUGGGA--UAGACUAGGAGACCCUCUGUCCUA--ACAACACCAGCCA 207
MBFV-NC_009029.2 GCAUA--UUGACACCCUGGGA--AAGACUAGGGGAUUUGCUGCUCUG---GACUUCGGCGU 209
MBFV-NC_032088.2 GCAUA--UUGACACCCUGGGA--UAGACUAGGAGAUCCGCGUCUCUG---GAACCCAGGU 210
MBFV-NC_009026.2 GCAUA--UUGACACCCUGGGA--UAGACCGGAGAAUUUGCUGCCUCG---CAACACCUCCCA 207
MBFV-NC_018705.3 GCAAGCUGUAAACCAGGGGAAGGACUAG--AGGUUAGAGGAGACCCCUUGCGAGUGAACA 207
TBFV-NC_027709.1 UCCCCCUCGACAGAGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
TBFV-NC_001672.1 UCCCCCUCGACAGAGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
TBFV-NC_001809.1 UCCCCCUCGACAGAGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
TBFV-NC_005062.1 UCCCCCUCGACAGAGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
TBFV-NC_003690.1 UCCCCCUCGACAGAGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
TBFV-NC_003687.1 UCCCCCAGGAAACUGGGG-----GGGCGGUUCUUGUUCUCCUAGGCCACCAUAC 203
80.....190.....200.....210.....220.....230.....2



MBFV-NC_002031.1 GUGAGCACAGUUUGCUCAAGAAUAAGCAGACCUUUGGAUGACAAACACAAAACCACU 258
MBFV-NC_012735.1 UAAAGCACAGAUUGCUU--GAAUAGCAGGGCGGGGAAGUUGAGAACCAACUAGUGUCU 258
MBFV-NC_008719.1 UUAAGCACAGAUUGCUU--GAAUAGCAGGGUGGGGAAGUUGGGAACCAACUAGUGUCU 258
MBFV-NC_012532.1 CCAGGCACAGAUUCGCCGAACUUCGGCGGCCGGUGUGG--GGAAAUCCA--GGGUUCU 258
MBFV-NC_035889.1 CCAGGCACAGAUUCGCCGAUAGCGGGCGGCCGGUGUGG--GGAAAUCCA--GGGUUCU 258
MBFV-NC_012533.1 CCUGGCACAGAGCGCC--AGAUCCGGGGUGUGGUGG--AGCAACCCAA--CAGGACU 258
MBFV-NC_002640.1 CCAGGCACAGAGCGCCGCAAGAUG--GAUUGGUGUG--UUGAUCCAAC--AGGUUCU 258
MBFV-NC_001477.1 CCAGGCACAGAACGCCAAAAAUG--GAAUGGUGCUG--UUGAAUCAAC--AGGUUCU 258
MBFV-NC_001475.2 CCAGGCACAGAACGCCAGAAAAUG--GAAUGGUGCUG--UUGAAUCAAC--AGGUUCU 258
MBFV-NC_001474.2 CCAGGCACAGAACGCCAGAAAAUG--GAAUGGUGCUG--UUGAAUCAAC--AGGUUCU 258
MBFV-NC_009028.2 CUAGGCACAGAUUCGCCGAAGUAG--UGGCUGGUGGUGAAAA--ACACA--U--GGA--UCU 258
MBFV-NC_040776.1 CUAGGCACAGAUUCGCCGAUUCUG--UGGCUGGUGCUGGAAA--ACACA----GGA--UCU 258
MBFV-NC_034151.1 CUCGGCACAGAACGCCGUAAGU--G--UGGCUGGUGUGAAAAUAAACA--C--AGAGUCU 258
MBFV-NC_007580.2 CAAGGCACAGAGCGCCGUAAAGUG--UGGCUGGUGGUGAA--AAAAUCA--C--AGGAUCU 258
MBFV-NC_001437.1 CUAGGCACAGAGCGCCGAAGUAG--UAGCUGGUGGUGAGGAAGAACCA--C--AGGAUCU 258
MBFV-NC_006551.1 CAAGGCACAGAGCGCCGAAAAUUG--UGGCUGGUGGGAUACUAGAACCA--C--AGGAUCU 258
MBFV-NC_001563.2 CACGGCACAGUGCGCCGACAUAGG--UGGCUGGUGGUG--CUAGAACA--C--AGGAUCU 258
MBFV-NC_009942.1 CACGGCACAGUGCGCCGACAAUGG--UGGCUGGUGGUG--CGAGAACA--C--AGGAUCU 258
MBFV-NC_000943.1 CAAGGCACCGAGCGCCGAACACUG--UGACUGAUGGGGGAGGAAGAACCA--C--AGGAUCU 258
MBFV-NC_012534.1 UCCGGCACAGAGCGCCGUAAG--UG--UGGCUGGUGGUG--GUAGAACA--C--AGUGUCU 258
MBFV-NC_015843.2 CUUGGCACAGAUUCGCCGAAG--UG--UGGCUGGUGGUG--GUAGAACA--C--AGGAUCU 258
MBFV-NC_009029.2 CUCGGCACAGAACGCCGUAGGA-----GCCGGAGGCCCAAAACA--C--CAGAUUCU 258
MBFV-NC_032088.2 CUAGGCACAGAACCGCCG--UAAUGA-----ACUGGGUGGCCCAAAACA--C--CGAAUCU 258
MBFV-NC_009026.2 CCCGGCACAGAACGCCGACA-----UGGUGGGAGGGGUCGUAAGACACC--AGAUCU 258
MBFV-NC_018705.3 CCACACAAAAACAGC-----AAUUGACACCUGGGAAGACUAGGAGACCCUCUGU 258
TBFV-NC_027709.1 CCAGACACAGAUAGUCUGACAAGGAGG--UGAUGCGUGACUCGGAAGAACACCCCGCU 258
TBFV-NC_001672.1 CCAGACACAGGUAGUCUGACAAGGAGG--UGAUGGUGACUCGGAAGAACACCCCGCU 258
TBFV-NC_001809.1 CCAGACACAGAUAGUCUGACAAGGAGG--UGAUGGUGACUCGGAAGAACACCCCGCU 258
TBFV-NC_005062.1 CCAGACACAGGAGUCUAAAAGGAGG--UGAUGGUGACUCGGAAGAACACCCCGCU 258
TBFV-NC_003690.1 CUAGGCACAGAUAGCCUGACAAGGAGA--UGGUGUGUGACUCGGAAGAACACCCCGCU 258
TBFV-NC_003687.1 CCAGGCACAGAUAGCCUGACAAGGAGA--UGGUGUGUGACUCGGAAGAACACCCCGCU 258
40.....250.....260.....270.....280.....290.....

