

MBFV-NC_018705.3 CCGUA----AACACAGUUUGAACGUUUUUUGGAGAGAG----ACUACUAUGUCAAAACAAA 52
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUGCAUGAAAAACCCC 52
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGAAUUUGGAAACGAGAG-UUUCUGGUGCAUGAAAAA-CCC 51
MBFV-NC_012533.1 AACAGUUUU-----CGUGUUGAGAAAGAAAACAGAU-UCUUCGGAUAUGAAAA--CCC 50
MBFV-NC_002640.1 -----UCUA-ACAGUUUGUUGAAUAGAGAGCAGAUUCUCUGGAAAAAUG----AA--CC 47
MBFV-NC_001474.2 -CGUAGUUCUA-ACAGUUU-UUUAAUUAGAGAGCAGAUUCUCUG-----AUGAAUA--CC 50
MBFV-NC_001477.1 ACGUAGUUCUA-ACAGUUU-UUU-AUUAGAGAGCAGAUUCUCUG-----AUGAACAA--CC 50
MBFV-NC_001475.2 ACGUAGUGCUG-ACAGUUU-UUU-AUUAGAGAGCAGAUUCUCUG-----AUGAACAA--CC 50
MBFV-NC_009026.2 AG-----AAAAGUUUCGAGGAAAGAC-GAGAAGCGAAUUCUCUGAAUGGCU---AGA 49
A--U-----UAACACAGUGCGAGCUGUUUCUUGGCACGAA--GAUCUCGAUGUCU---AAG 49
A--U-----UAACACAGUGCGAGCUGUUUCUAGCAGCAA--GAUCUCGAUGUCU---AAG 49
U--U-----UAACACAGUCUGAAUAGUUUUUAGCAAGGG--AUUCCCAUGUCU---AAA 49
U--U-----UAACACAGUCUGAAUAGUUUUUAGCAGAGG--UACUACCAUGUCU---AAA 49
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUGAAAAACCAUGUCU---AAA 49
MBFV-NC_012534.1 CCGU----AAACACAGUUUGAAC-GUUUUUUGGAGAGA--GAUUAUUAUGUUUAACAAA 52
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAAC-GUUUUUUGGAUAGA--GACAACUAUGUCUAACAAA 52
MBFV-NC_007580.2 ---U----AACUUGACUGCGAACAGUUUUUAGCAGGGAAU-UACCCAUGUCU---AAA 49
MBFV-NC_000943.1 -----UAACGCGGUUUGAACAGUUUUUUGGAGCUUUUGAUUUCAAUGUCU---AAA 49
MBFV-NC_006551.1 ---U----UAACACAGUGCCGGCAGUUUUUUGAGC-GUUGAUUUUCAAUGUCU---AAG 49
MBFV-NC_001437.1 ---U----UAGUGCAGUUUAACAGUUUUUAGAACGGAUAACC-AUGACU---AAA 49
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAAAGUUUUUGAACAG-----CAAUGACC---AAA 49
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAAGUUUUUUGAUCAG-----UGGAUUAUGACU---AAA 49
MBFV-NC_012735.1 -UUUCUACUGAAAGGAGUAAGA-----AAGGAGAUCAUUCCCAUGGCAACAAAG 52
MBFV-NC_008719.1 AUUUCUACUGAAAGG-GUAGAGAA-----AAGGAGUUUGCUUUCUAGGCAACAAGA 52
MBFV-NC_002031.1 UAAUCGUUCGUUGAGCGAUUAGCA-----GAGAACUGAC-CAGAACAUGUC-----45
TBFV-NC_003687.1 -----AGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGAGUGGUUAUG-----AUG 46
TBFV-NC_027709.1 --GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCAGGAAG 52
TBFV-NC_001672.1 --GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGUCAGAAG 52
TBFV-NC_001809.1 --GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCAGGAAG 52
TBFV-NC_005062.1 --GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCCGGGAAG 52
TBFV-NC_003690.1 --GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCCGGGAAG 52
.....10.....20.....30.....40.....50.....6



MBFV-NC_018705.3 A-AACCAGGAAGACCC---GGGGCAACCCGGGUUGUCAAUUAGCUAAAGCGCGGCGCAUC 108
MBFV-NC_012532.1 A-AAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAUAUAGCUAAACCGCGGAGUAGC 105
MBFV-NC_035889.1 A-AAAAAGAAAU--CC---GGAGGAUUCGGGAUUGUCAUAUAGCUAAACCGCGGAGUAGC 105
MBFV-NC_012533.1 A-AAAAAGG-----GCCGCAUCGCGGCCUAUCUAUAGCUGAAGCGCUUUGGAUC 99
MBFV-NC_002640.1 A-ACGAAAAAAG--GU---G--GUUAGACCACCUUUCUAUAGCUGAAACGCGAGAGAAA 99
MBFV-NC_001474.2 A-ACGGAATAAG--GC---G--AAAAACACGGCCUUUCUAUAGCUGAAACGCGAGAGAAA 102
MBFV-NC_001477.1 A-ACGGAATAAG--AC---G--GGUCGACCGUCUUUCUAUAGCUGAAACGCGCGAGAAA 102
MBFV-NC_001475.2 A-ACGGAAGAAG--AC---G--GGAAACCGGUCUAUCUAUAGCUGAAACGCGUGAGAAA 102
MBFV-NC_009026.2 A-AACCAGGAAGACCC---GGAGGAAACCGGGUCGUCUAUAGCUAAAGCGUACGGCGGC 105
MBFV-NC_001563.2 A-AACCAGGAGGGCCC---GGUAAAAACCGGGCUGUCUAUAGCUAAACCGCGUAUGCC 105
MBFV-NC_009942.1 A-AACCAGGAGGGCCC---GGCAAGAGCCGGGCUGUCUAUAGCUAAACCGCGGAUUGCC 105
MBFV-NC_009028.2 A-AACCAGGAAGUCC---GCUGCAAAGCGGACUGUCUAUAGCUAAACCGCCUGGCUAG 105
MBFV-NC_040776.1 A-AACCAGGAGGACCC---GCCGGAAGCGGGUUGUCUAUAGCUGAAACGCCGACCCAG 105
MBFV-NC_034151.1 A-AACCAGGAGGACCC---GGAAAGAACCGGGUCGUUAUAGCUGAAGCGCACAUCCCG 105
MBFV-NC_012534.1 A-AACCAGGAAACCC---GGGGCAACCCGGGUUGUCAUAUAGCUAAAGCGCGGUGCAUC 108
MBFV-NC_015843.2 A-AACCAGGAAGACCC---GGCUCAGGCGGGUUGUCAUAUAGCUAAAGCGCGGAACGUC 108
MBFV-NC_007580.2 A-AACCAGGAACACCC---GGUAGAAACCGGGUUGUCAUAUAGCUAAACCGCGGUAUUC 105
MBFV-NC_000943.1 A-AACCAGGAGGACCC---GGGAAGCCCCGGGUCGUCUAUAGCUAAACCGCGCAUACC 105
MBFV-NC_006551.1 A-AACCAGGAGGGCCC---GGAAGAAACCGGGCCAUCUAUAGCUGAAACGCGGCAUACC 105
MBFV-NC_001437.1 A-AACCAGGAGGGCCC---GGUAAAAACCGGGCUAUCUAUAGCUGAAACCGGCCUACC 105
MBFV-NC_009029.2 A-AACCAGGAAGGCC---GGUAGAAAUCCGGGCCGUCUAUAGCUAAAGCGCGGUGCGUC 105
MBFV-NC_032088.1 A-AACCAGGAAGUCC---GGCAUUCGUGGACUGUCUAUAGCUAAAGCGCGGAGCAUC 105
MBFV-NC_012735.1 G-GGAUGAAUAGUCUCGGGUCGAGUCCGAGGCGUCUAUAGGUAG--CGGCUAG--106
MBFV-NC_008719.1 G-GAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCUAUAGGUGA--CGGCCAG---106
MBFV-NC_002031.1 ---UGGUCGUAAGCUCAGGGAAAAACCCUGGGCGUCUAUAGGU-A--CGACGAGGAGU 99
TBFV-NC_003687.1 ACCACUUCUAAAGGAAAGGGGGGCGUCCCCUAGGC-CAGAGCUAAAGUGACCGCAAAU 106
TBFV-NC_027709.1 UCCACUCCUGAAAGGUAAGGGGGGCGGUCCCCUGAC-CAGUGUCGAAAGAGACCCGAA--110
TBFV-NC_001672.1 GCCAUCCUGAAAGGUAAGGGGGGCGGUCCCCUGAC-CAGUGUCGAAAGAGACCCGAA--110
TBFV-NC_001809.1 ACCAUCCUGAAAGGCAAGGGGGGCGUCCCCUGAC-CAGUGUCGAAAGAGACCCGAA--110
TBFV-NC_005062.1 GCCAUUCUGAAAGGAAAGGGGGGCGGUCCCCUGAC-CAGUGUCGAAAGAGACCCGUA--110
TBFV-NC_003690.1 GCCGUUCUAAAGGAAAGGGGGGGGUGCCCCUGAC-CAGCGUCGAAAGUGGCCCCAA--110
0.....70.....80.....90.....100.....110.....1



MBFV-NC_018705.3 CC-----NNNNNNNN-----CCCCGCGGAAAUCAAAGCAAGGCCCAACCUGG 149
MBFV-NC_012532.1 CCGUG-----NNNNNNNN-----GGUUGAGGAGACCCCCCGGAAACGCAAAACAG 151
MBFV-NC_035889.1 CCGUG-----NNNNNNNN-----GUUAGGAGAGACCCCCCGGAAACGCAAAACAG 150
MBFV-NC_012533.1 UAGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAACAUAACACAG 151
MBFV-NC_002640.1 CCGCGUAUCAANNNNNNNNACUAG--AGGUUAGAGGAGACCCCCCAACA--CAAAAACAG 155
MBFV-NC_001474.2 CCGCGUGU--NNNNNNNN--CUAG--AGGUUAGAGGAGACCCCCCG--CGAAACAAAAACAG 155
MBFV-NC_001477.1 CCGCGUGU--NNNNNNNN--UAG--AGGUUAGAGGAGACCCCCCGCACAACAACAACAG 155
MBFV-NC_001475.2 CCGUGUGU--NNNNNNNNACUAG--AGGUUAGAGGAGACCCCCCG--CAAAUAAAAACAG 155
MBFV-NC_009026.2 CAACG-----NNNNNNNN-----CCCCUUGGCAAAACAGUUAACGCACCAAAAGAAACAG 154
MBFV-NC_001563.2 CCGCG-----NNNNNNNAGAG--AGACCCGUGCCA-AAAACACCA-A-AAGAAACAG 153
MBFV-NC_009942.1 CCGCG-----NNNNNNNN--GG--AGACCCCGUGCCACAACACACCACA-ACAAAACAG 153
MBFV-NC_009028.2 CGUGU-----NNNNNNNNGUGG--AGACCCUCAGCAC-CAAGCGCGAA--ACAAAACAG 153
MBFV-NC_040776.1 CGUUU-----NNNNNNNA-GG--AGACCCUUGGCACACAAGCGCAUUGCA--GACAAACAG 154
MBFV-NC_034151.1 CGCUA-----NNNNNNNN-----AGACCCCUUGGCAGUGAAC-GCAA-UAAAAACAG 150
MBFV-NC_012534.1 CC-----NNNNNNNN--GG--AGACCCUUGCGAGUUAACACCAAA-ACAAAACAG 153
MBFV-NC_015843.2 CC-----NNNNNNNNA-GG--AGACCCUUGCGAGUGAACACACCA-A-GAAACAG 153
MBFV-NC_007580.2 CCGCG-----NNNNNNNN--UU--AGAGGAGACCCCUUGCCGUUAACGC-AAACAACAG 153
MBFV-NC_000943.1 CCGCG-----NNNNNNNN--G--ACCCACUCUCAAAAGCAUC-----AAACAACAG 147
MBFV-NC_006551.1 CCGCG-----NNNNNNNN-----ACCCGCGA-UCAUAAGCAUCA-----AAAAACAG 146
MBFV-NC_001437.1 CCGCG-----NNNNNNNN--U--AGAGGAGACCCCGCAUUGCA-----UCA-AACAG 147
MBFV-NC_009029.2 CCGCG-----NNNNNNNAGAGG--AGACCCUCCCAAAAGAAGCGCAACACAAAACAG 157
MBFV-NC_032088.1 CCGCG-----NNNNNNNAGAGGUUAGAGGAGACCCCCACAACCGCAA-ACAAAACAG 158
MBFV-NC_012735.1 -----GUGANNNNNNNN-----C-CCAUCCUCCCGAGCACACAUAGC-GGAC 147
MBFV-NC_008719.1 -----GCCANNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UUAUGC-GGAC 147
MBFV-NC_002031.1 UCGCUCCUUGUNNNNNNN-----GAGACCCUCCAGGGAACAAUAGUGGGAC 146
TBFV-NC_003687.1 AAGU-----NNNNNNNNCU-UCCUCGGGAUUUUUCCGCUCCU-----AUAC 146
TBFV-NC_027709.1 -----NNNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCU-----AUAC 147
TBFV-NC_001672.1 -----NNNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCU-----AUAC 147
TBFV-NC_001809.1 -----NNNNNNNN-UCUCUUCAGGGUUUUUCCUCCUCCU-----AUAC 146
TBFV-NC_005062.1 -----NNNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCU-----AUAC 147
TBFV-NC_003690.1 -----NNNNNNNNCUCUCUUCAGGAUUUUUCCUCCUCCU-----AUAC 147
20.....130.....140.....150.....160.....170.....1



MBFV-NC_018705.3 CAAAGCUGUAAACCAGGGGAAGGACUAGAGGUUAGAGGAGACCCCUUGCGAGUGAACAC 209
MBFV-NC_012532.1 CAUAUUGAGC--UGGGA--AAGACCAG-AGACUCCAUGAGUUUCCACCACGUGGCCGCC 206
MBFV-NC_035889.1 CAUAUUGAGC-CUGGGA--AAGACCAG-AGACUCCAUGAGUUUCCACCACGUGGCCGCC 206
MBFV-NC_012533.1 CAUAUUGACACCUGGGA--AAGACCGG-AGACUCUGUUGGCUUCCCCACACCAGCCCC 208
MBFV-NC_002640.1 CAUAUUGAGC-CUGGGA--AAGACCAG-AGAUCUCUGC-GUCUCUGCAACAUA--AUCC 208
MBFV-NC_001474.2 CAUAUUGAGC-CUGGGA--AAGACCAG-AGAUCUCUGC-GUCUCCUCAGCAUA--UUCC 208
MBFV-NC_001477.1 CAUAUUGAGC-CUGGGA--GAGACCAG-AGAUCUCUGC-GUCUCUACAGCAUA--UUCC 208
MBFV-NC_001475.2 CAUAUUGAGC-CUGGGA--GAGACCAG-AGAUCUCUGC-GUCUCCUCAGCAUA--UUCC 208
MBFV-NC_009026.2 CAUAUUGACACCUGGGA--UAGACCGGAGAUUUGCUGCC--UCGCAACACCUCCACC 209
MBFV-NC_001563.2 CAUAUUGACACCUGGGA--UAGACUAGGGGAUCUUCUGCU--CUGCACAACCAGCCACA 208
MBFV-NC_009942.1 CAUAUUGACACCUGGGA--UAGACUAGGAGAUUUCUGCU--CUGCACAACCAGCCACA 208
MBFV-NC_009028.2 CAUAUUGAGCGCCUGGGA--AAGACCGGGAGAUUUCUGCU--UUCCAACACCAGCCACU 208
MBFV-NC_040776.1 CAUAUUGACACCUGGGA--AAGACCGGGAGACCCUCUGCU--UCGCAACACCAGCCACU 209
MBFV-NC_034151.1 CAUAUUGACACCUGGGA--UAGACUGGGAGACCCUCUGCUU-AUUCAACACCAGCCACU 207
MBFV-NC_012534.1 CAUAUUGACACCUGGGA--GAGACUAGGAGACCCUCUG-UC-CUAACAACACCAGCCACU 209
MBFV-NC_015843.2 CAUAUUGACACCUGGGA--UAGACUAGGAGACCCUCUG-UC-CUAACAACACCAGCCACU 209
MBFV-NC_007580.2 CAUAUUGACACCUGGAAAGA---CAGGAGAUCCCUUGCUU--UUUCAACACCAGCCACA 207
MBFV-NC_000943.1 CAUAUUGACACCUGGGAA-AAGACUAGGAGAUUUCUGCUCUAUCCAACAUAGUCACA 206
MBFV-NC_006551.1 CAUAUUGACACCUGGGAAUAGACUAGGAGAUUUCUGCUCUAUCCAACAUCAACACACA 206
MBFV-NC_001437.1 CAUAUUGACACCUGGGAAUAGACUGGGAGAUUUCUGCUCUAUCCAACAUCAAGCUACU 206
MBFV-NC_009029.2 CAUAUUGACACCUGGGA--AAGACUAGGGGAUUGCUGCUC--UGGACUUCGGCU-CU 211
MBFV-NC_032088.1 CAUAUUGACACCUGGGA--UAGACUAGGAGAUCCGUGCUC--UGGAACCCAGU-CU 212
MBFV-NC_012735.1 CAUAUUGAGCGCCAGGGA--AAGACCGGAGACACUCCUUGAUUUCACCUUUCUGCCCUA 205
MBFV-NC_008719.1 CAUAUUGAGCGCCUGGGA--AAGACCGGAGACACUCCUUGAUUUCACCUUUCACCCUUC 205
MBFV-NC_002031.1 CAUAUUGAGCGCCAGGGA--AAGACCGGAGUGGUUCUCUGCUUUUC-CUCCAGGUGUCU 203
TBFV-NC_003687.1 UAAAUUCCCCCAGGAACUGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCACCAUCC 205
TBFV-NC_027709.1 CAAA-UUCCCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCAUACCC 205
TBFV-NC_001672.1 AAAA-UUCCCCCUCGGUAGAGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCAUACCC 205
TBFV-NC_001809.1 CAAA-UUCCCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCAUACCC 205
TBFV-NC_005062.1 CAAA-UUCCCCCUCGACAGAGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCAUACCC 205
TBFV-NC_003690.1 CAAA-UUCCCCCUCGUCAGAGGGGGGGCGGUUCUUGUUCUC-CUGAGCCACCAUACCU 205
80.....190.....200.....210.....220.....230.....2



MBFV-NC_018705.3 ACACAAAAACAGC-----AUAUUGACACCUGGGAAAGACUAGGAGACCCUCUGU 258
MBFV-NC_012532.1 AGGCACAGAUCGCC---GAACUUCGGCGGCCGGUGUGGGGAAAUCCAUGGUUUCU 258
MBFV-NC_035889.1 AGGCACAGAUCGCC---GAAUAGCGGCGGCCGGUGUGGGGAAAUCCAUGGGUCUU 258
MBFV-NC_012533.1 UGGCACAGAGCGCC---AGAUCGGGGUUGGUGUGGAGCAACCCAACAGGACU 258
MBFV-NC_002640.1 AGGCACAGAGCGCC---GCAAGAUG--GAUUGGUGUUGUUGAUCCAACAGGUUCU 258
MBFV-NC_001474.2 AGGCACAGAACGCC---AGAAAAUG--GAAUGGUGUGUUGAUCAACAGGUUCU 258
MBFV-NC_001477.1 AGGCACAGAACGCC---AAAAAUG--GAAUGGUGUGUUGAUCAACAGGUUCU 258
MBFV-NC_001475.2 AGGCACAGAACGCC---AGAAAAUG--GAAUGGUGUGUUGAUCAACAGGUUCU 258
MBFV-NC_009026.2 CGGCACAGAACGCC---GACA-UGGUGGAGGGGUCGUA--GACACCAGAUUCU 258
MBFV-NC_001563.2 CGGCACAGUGCGCC---GACAUAGGUGGCUGGUGGUGCUA--GAACACAGGAUCU 258
MBFV-NC_009942.1 CGGCACAGUGCGCC---GACAAUGGUGGCUGGUGGUGCGA--GAACACAGGAUCU 258
MBFV-NC_009028.2 AGGCACAGAUCCGC---GCAAGUAGUGGCUGGUGGUGGAA--AACACAUUGAUUCU 258
MBFV-NC_040776.1 AGGCACAGAUCCGC---GAAAUUGUGGCUGGUGGUGCGAA--AACACA-GGAUCU 258
MBFV-NC_034151.1 CGGCACAGAACGCC---GUAAGU-GUGGCUGGUGGUGAAAAUAAACACAGAGUCU 258
MBFV-NC_012534.1 CGGCACAGAGCGCC---GUAAGU-GUGGCUGGUGGUGGUA--GAACACAGUGUCU 258
MBFV-NC_015843.2 UGGCACAGAUCCGC---GAAAGU-GUGGCUGGUGGUGUA--GAACACAGGAUCU 258
MBFV-NC_007580.2 AGGCACAGAGCGCC---GUAAGUGUGGCUGGUGGUGGAA--AAAAUCACAGGAUCU 258
MBFV-NC_000943.1 AGGCACCGAGCGCC---GAACAUUGAGUGAUUGGGGAGAGACCACAGGAUCU 258
MBFV-NC_006551.1 AGGCACAGAGCGCC---GAAAAUUGUGCUGGUGGGGAAAGACACACAGGAUCU 258
MBFV-NC_001437.1 AGGCACAGAGCGCC---GAAGAUUGUAGCUGGUGGUGAGGAAGAACACAGGAUCU 258
MBFV-NC_009029.2 CGGCACAGAACGCC---GUUGAGGA-----GCCGGAGGCCAAAACACCAGAUUC 258
MBFV-NC_032088.1 AGGCACAGAUCCGC---G-UAAUGA-----ACUGGGUGGCAAAAACACCGAUUC 258
MBFV-NC_012735.1 AAGCACAGAUUGCUU--GAUUGAGGGCGGGGAAGUUGGAAACCAACUAGUGUCU 258
MBFV-NC_008719.1 AAGCACAGAUUGCUU--GAUUGCAGGGUGGGGAAGUUGGGAACCAACUAGUGUCU 258
MBFV-NC_002031.1 GAGCACAGUUUGCCUAGAAUAGACAGACCCUUUGGAUGACAACCAACAAACCCACU 258
TBFV-NC_003687.1 AGGCACAGAUAGCUGACAAGGAGGUGUGUGUGACUCGGAACAAACACCCG--CU 258
TBFV-NC_027709.1 AGACACAGAUAGCUGACAAGGAGGUGAGUGUGUGACUCGGAACAAACACCCG--CU 258
TBFV-NC_001672.1 AGACACAGGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAACAAACACCCG--CU 258
TBFV-NC_001809.1 AGACACAGUAGUCUACAAGGAGGUGAUGUGUGACUCGGAACAAACACCCG--CU 258
TBFV-NC_005062.1 AGACACAGAUAGUCUGAAAAGGAGGUGAUGCGUGUCUCGGAACAAACACCCG--CU 258
TBFV-NC_003690.1 AGACACAGAUAGUCUGAAAAGGAGGUGAUGCGUGUCUCGGAACAAACACCCG--CU 258
40.....250.....260.....270.....280.....290...

