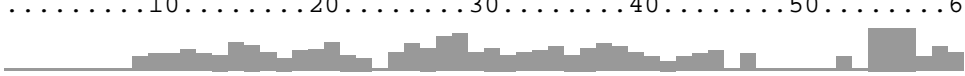


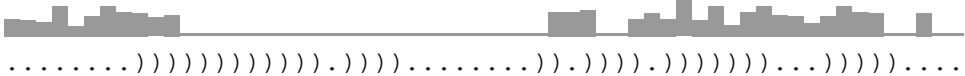
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MBFV-NC_009942.1 A---U---UAACACAGUGCGAGC-UGUUUUCUUGGCACGAAGA--U-C--U--CGAUGUCU 46
MBFV-NC_001437.1 U---U---AGUGCAGUUUAAACA-GUUUUUUAGAACGGAAGA--U-A--A--CCAUGACU 46
MBFV-NC_007580.2 U---A---ACUUGACUUGCGAACA-GUUUUUUAGCAGGGAAUU--A-C--C--CAAUGUCU 46
MBFV-NC_034151.1 U---U---UAACACAGUGUGAAU-AGUUUUUUAGAACAGAGUGAA--U-G--A--CCAUGUCU 46
MBFV-NC_009028.2 U---U---UAACACAGUCUGAAU-AGUUUUUU-AGCAAGGGA-UU-U--C--CCAUGUCU 46
MBFV-NC_040776.1 U---U---UAACACAGUCUGAAU-AGUUUUUU-AGCAGAGU-AC-U--A--CCAUGUCU 46
MBFV-NC_000943.1 U---U---UAACCGCGUUGGAAU-AGUUUUUUUGGAGCUUUUGAU-U--C--AAAUGUCU 46
MBFV-NC_006551.1 ---U---UAACACAGUGCCGGC-AGUUUCUUUGAGC-GUUGAUU-U--U--CAAUGUCU 46
MBFV-NC_012534.1 CCGU---AAACACAGUUUGAAC--GUUUUUUGGAGAGAG--AU-U--A--UUUUGUUU 46
MBFV-NC_015843.2 ACGU---AAACACAGUUUGAAC--GUUUUUUGGAGAUAG--AC-A-A--CUAUGUCU 46
MBFV-NC_009026.2 AGAA---AAGUUUUC---GAGG-AAAGACGAG-AAGCGAAUUCU-C--U--GAAUGGCU 46
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAAGUUUUUUUGAACAG-----CAAUGACC 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAGUUUUUUUGAUCAGUGG-----A--UUAUGACU 46
MBFV-NC_002640.1 -----UCUAAACAGUUUGUUUGAAUAGAGAGCAG-AUCUCU--GGAA--AAAUG--- 43
MBFV-NC_001474.2 -CGU---AGUUCUAACAGUUU--UUUAAUUAGAGAGCAG-AUCUCU-G-----AUGAAU 46
MBFV-NC_001477.1 ACGU---AGUUCUAACAGUUU--UUU--AUUAGAGAGCAG-AUCUCU-G-----AUGAA 46
MBFV-NC_001475.2 ACGU---AGUGCUGACAGUUU--UUU--AUUAGAGAGCAG-AUCUCU-G-----AUGAAC 46
MBFV-NC_012532.1 AACA---GGUUU-----AAUUUGGAUUUGGAAACGAGAGUUUCU--GG-----UCAUGAAA 46
MBFV-NC_035889.1 ACAG---GUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCU--GG-----UCAUGAAA 46
MBFV-NC_012533.1 AACA---GUUUU-----CGUUGAGAAAAGAAAACAGAUUUCU--GG-----AAAUGAAA 46
MBFV-NC_002031.1 -----UAAUCGUUCGUUGAGC--GAUUAGCAGAGAACUGACCAGA--ACAUG--- 43
MBFV-NC_012735.1 -----UUUCUACUG-AAAGGAGUAGAAGAAAGGAGAUCAUUC--CAAUGGCA 46
MBFV-NC_008719.1 -----AUUUCUACUGAAAGGGUAGAGAA--AAGGAGUUUUGCUUC--CAAUGGCA 46
MBFV-NC_001809.1 -----GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGGC 46
MBFV-NC_005062.1 -----GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCC 46
MBFV-NC_027709.1 -----GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGGC 46
MBFV-NC_001672.1 -----GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGUC 46
MBFV-NC_003690.1 -----GUGUUGAGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGG-----AUGGCC 46
MBFV-NC_003687.1 -----AGAAAAAGACAGCUUAGGAGAAACAAGAGCUGGGAGUGGUUAUG--- 43
MBFV-NC_018705.3 CCGU---AAACACAGUUUGAACGUUUUUUGGAGAGAGACUACU-----AUGUCA 46
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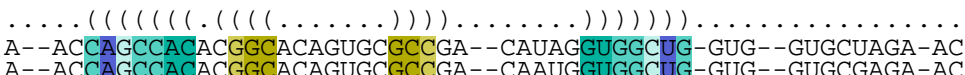
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MBFV-NC_001437.1 A---AAA-AACCAGGAGGGGCC--GGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGG 99
MBFV-NC_007580.2 A---AAA-AACCAGGAAACACC--GGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGG 99
MBFV-NC_034151.1 A---AAA-AACCAGGAGGACCC--GGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGCAC 99
MBFV-NC_009028.2 A---AAA-AACCAGGAAAGUCC--GCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCU 99
MBFV-NC_040776.1 A---AAA-AACCAGGAAAGACC--GGCGGAAGCGGGUUGUCAAUAUGCUAAAACGCCCC 99
MBFV-NC_000943.1 A---AAA-AACCAGGAGGACCC--GGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGG 99
MBFV-NC_006551.1 A---AGA-AACCAGGAGGGGCC--GGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGG 99
MBFV-NC_012534.1 AACAAAA-AACCAGGAAACACC--GGGGCAACCGGGUUGUCAAUAUGCUAAAACGCGG 102
MBFV-NC_015843.2 AACAAAA-AACCAGGAAGACCC--GGCUCAGGCCGGGUUGUCAAUAUGCUAAAAGCGCGG 102
MBFV-NC_009026.2 A---GAA-AACCAGGAAGACCC--GGAGGAAACCGGGUCGUCAAUAUGCUAAAAGCGUAC 99
MBFV-NC_009029.2 A---AAA-AACCAGGAAGGCC--GGUAGAAAUCCGGGCCUGUCAAUAUGCUAAAAGCGCG 99
MBFV-NC_032088.1 A---AAA-AACCAGGAAGUCC--GGCAAUCGUCGGACUGUCAAUAUGCUAAAAGCGCGG 99
MBFV-NC_002640.1 AA---CCA-ACGAAAAAAG--GU---GG-UUAGACCACCUU-UCAAUAUGCUGAAACGCGA 99
MBFV-NC_001474.2 AA---CCA-ACGAAAAAAG--GC---GA-AAACACGCCUU-UCAAUAUGCUGAAACGCGA 96
MBFV-NC_001477.1 AA---CCA-ACGAAAAAAG--AC---GG-GUCGACCGUCUU-UCAAUAUGCUGAAACGCGC 96
MBFV-NC_001475.2 AA---CCA-ACGGAAGAG--AC---GG-GAAACCCGUCUA-UCAAUAUGCUGAAACGCGU 96
MBFV-NC_012532.1 AACCCCA-AAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAAUAUGCUAAAAGCGCG 99
MBFV-NC_035889.1 AA-CCCA-AAAAAGAAAU--CC---GGAGGAUUCGGAUUGUCAAUAUGCUAAAAGCGCG 99
MBFV-NC_012533.1 A---CCCA-AAAA--AGGG--CC---AUCGCGGCCUAUCAAUAUGCUGAAGCGCUU 93
MBFV-NC_002031.1 -----U-CUGGUCGUAAGAGCAGGAAAAACCCUGGCCUUCAAUAUGGUACGACGAG 96
MBFV-NC_012735.1 ACAAAGG-GGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAGC-----G 100
MBFV-NC_008719.1 ACAAGAG-GAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGUGAC-----G 100
MBFV-NC_001809.1 AGGAAGACCAUCCUGAAAGGCAAGGGGGGCGGUCCCCUCGACAAUAUGCUAAAAGAGACC 106
MBFV-NC_005062.1 GGAAGGCCAUUCUGAAAGGAAAGGGGGGCGGUCCCCUCGACAGUGUCGAAAGAGACC 106
MBFV-NC_027709.1 AGGAAGUCCAUCCUGAAAGGUAAGGGGGGCGGUCCCCUCGACAGUGUCGAAAGAGACC 106
MBFV-NC_001672.1 AAGAAGGCCAUCCUGAAAGGUAAGGGGGGCGGUCCCCUCGACAGUGUCGAAAGAGACC 106
MBFV-NC_003690.1 GGAAGGCCGUCUAAAAGGAAAGGGGGGGGUCCCCCUCGACAGCGUCGAAAGUGGCC 106
MBFV-NC_003687.1 ---AUGACCUCUCAAAGGAAAGGGGGGCGGUCCCCUAGGCCCAAGCUAAAAGUGCC 100
MBFV-NC_018705.3 AACAAAA-AACCAGGAAGACCC--GGGGCAACCCGGGUUGCAUAUAUGCUAAAAGCGCG 102
0.....70.....80.....90.....100.....110.....1



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MBFV-NC_009942.1 AAUGCCCCGCG-----NNNNNNNN-----GGAGA-----CCCCGUGCCACAAAACACCAC 143
MBFV-NC_001437.1 CCUACCCCCGCG-----NNNNNNNN-----UAGAGGAGA-----CCCCGCAUUUG----- 137
MBFV-NC_007580.2 CGUAUCCCGCG-----NNNNNNNN-----UUAGAGGAGA-----CCCCUUGCCGUUAACG----- 143
MBFV-NC_034151.1 AUCCCGCGCUA-----NNNNNNNN-----AGA-----CCCUUGGCAGUGAA-CGCAA 140
MBFV-NC_009028.2 GGCUAGCGUGU-----NNNNNNNN-----GUGGAGA-----CCCUCAGCAC-CAAGCGCGA 144
MBFV-NC_040776.1 AGCCAGCGUUU-----NNNNNNNN-----A-GGAGA-----CCCUUGGCACACAAGCGCAA 144
MBFV-NC_000943.1 CAUACCCCGCG-----NNNNNNNN-----GA-----CCCCACUCUCAAAGCAUC-- 138
MBFV-NC_006551.1 CAUACCCCGCG-----NNNNNNNN-----A-----CCCCGCA-UCAUAAGCAUC-- 136
MBFV-NC_012534.1 UGCAUCC-----NNNNNNNN-----GGAGA-----CCCUUGGCAGUUAACACCAA 143
MBFV-NC_015843.2 AACGUCCC-----NNNNNNNN-----AGGAGA-----CCCUUGGCAGUGAACCACC 144
MBFV-NC_009026.2 GGCGGCCAACG-----NNNNNNNN-----C-----CCCUU--GGCAAAACAGUUAACGCACCA 144
MBFV-NC_009029.2 UGCGUCCCGCG-----NNNNNNNN-----AGAGGAGA-----CCCUCCCAAAAGAGCGCAA 147
MBFV-NC_032088.1 AGCAUCCCGCG-----NNNNNNNN-----AGAGGUUAGAGGAGACCCCCACAU-ACCGCAA 149
MBFV-NC_002640.1 GAGAAACCGCGUAUCAANNNNNNNNACUAGAGGUUAGAGGAGACCCCCCAACA----- 146
MBFV-NC_001474.2 GAGAAACCGCGUGU---NNNNNNNN--CUAGAGGUUAGAGGAGACCCCC--CGAAAC---- 146
MBFV-NC_001477.1 GAGAAACCGCGUGU---NNNNNNNN--UAGAGGUUAGAGGAGACCCCCCGACAAC---- 146
MBFV-NC_001475.2 GAGAAACCGCGUGU---NNNNNNNNACUAGAGGUUAGAGGAGACCCCCCG--CAAA----- 146
MBFV-NC_012532.1 AGUAGCCCGUG-----NNNNNNNN-----GCUUAGAGGAGACCCCCCGGAAAC-- 142
MBFV-NC_035889.1 AGUAGCCCGUG-----NNNNNNNN-----GUUAGAGGAGACCCCCCGGAAAC-- 141
MBFV-NC_012533.1 UGGAUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAAC---- 142
MBFV-NC_002031.1 AGUUCGCUCCUUG---NNNNNNNN-----GAGACCCUCCAGGGAACAA-- 136
MBFV-NC_012735.1 GCUAGAGU-----A---NNNNNNNN-----C-CCAUCCUCCGAGCACAC-- 138
MBFV-NC_008719.1 GCCAAGGCC---A---NNNNNNNN-----GAGGAGACCCUCCGAGCAC-U--- 138
MBFV-NC_001809.1 GCGA-----NNNNNNNN-----UCUCUUCAGGGUUUUUCCU 136
MBFV-NC_005062.1 GCUA-----NNNNNNNN-----CUCUCUUCAGGAAUUUCCU 137
MBFV-NC_027709.1 GCAA-----NNNNNNNN-----CUCUCUUCAGGAAUUUCCU 137
MBFV-NC_001672.1 GCAA-----NNNNNNNN-----CUCUCUUCAGGAAUUUCCU 137
MBFV-NC_003690.1 CCAA-----NNNNNNNN-----CUCUCUUCAGGAAUUUCCU 137
MBFV-NC_003687.1 GCAAUAAGU-----NNNNNNNN-----CU-UCUCCGGGAAUUUCCG 136
MBFV-NC_018705.3 CGCAUCCC-----NNNNNNNNCCCCGCGGAAA-----UCAAGCAAGGCCCAACCU 147
20.....130.....140.....150.....160.....170.....1



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MBFV-NC_009942.1 AACAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGAUUCUUC-U---GCUCUGCAC 197
MBFV-NC_001437.1 CAUCAACAGCAUAUUGACACCUGGGAA-UAGACUGGGAGAUUCUCUG--CUUAUUC 193
MBFV-NC_007580.2 CAAACACAGCAUAUUGACACCUGGGA--AGACAG--GAGAUCCC--UGCUUUUC 194
MBFV-NC_034151.1 AUAAAAACAGCAUAUUGACACCUGGGA--UAGACUGGGAGACCCUCUG--CUUAU-UUC 194
MBFV-NC_009028.2 A-ACAAACAGCAUAUUGACCCUGGGA--AAGACCGGGAGAUCCUCUG--CU--U-UCC 195
MBFV-NC_040776.1 AGACAAACAGCAUAUUGACACCUGGGA--AAGACCGGGAGACCCUCUG--CU--C-UGC 196
MBFV-NC_000943.1 -AAACAACAGCAUAUUGACACCUGGGA--AAGACUAGGAGAUUCUUCUG--CUCUAUUC 193
MBFV-NC_006551.1 AAAAAACAGCAUAUUGACACCUGGGAUUGACUAGGAGAUUCUUCUG--CUCUAUUC 193
MBFV-NC_012534.1 AACAAACAGCAUAUUGACACCUGGGA--AAGACUAGGAGACCCUCUG--UC--CUAAC 196
MBFV-NC_015843.2 AA-GAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGACCCUCUG--UC--CUAAC 196
MBFV-NC_009026.2 AAAGAAACAGCAUAUUGACACCUGGGA--UAGACC-GGAGAAUUGCU--GCCUC--C 196
MBFV-NC_009029.2 CACAAACAGCAUAUUGACACCUGGGA--AAGACUAGGGGAUUGUCUG--CUC---UG 198
MBFV-NC_032088.1 -ACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGAUCCGUG--CUC---UG 199
MBFV-NC_002640.1 -CAAAAACAGCAUAUUGACG-CUGGGA--AAGACC--AGAGAUCCUG--CUGUCUG 196
MBFV-NC_001474.2 -AAAAACAGCAUAUUGACG-CUGGGA--AAGACC--AGAGAUCCUG--CUGUCUCCU 196
MBFV-NC_001477.1 -AACAAACAGCAUAUUGACG-CUGGGA--GAGACC--AGAGAUCCUG--CUGUCUCUA 196
MBFV-NC_001475.2 -UAAAAACAGCAUAUUGACG-CUGGGA--GAGACC--AGAGAUCCUG--CUGUCUCCU 196
MBFV-NC_012532.1 -GAAAACAGCAUAUUGACG--UGGGA--AAGACC--AGAGAUCCAGAGUUUCCACC 194
MBFV-NC_035889.1 -GAAAACAGCAUAUUGACG-CUGGGA--AAGACC--AGAGAUCCAGAGUUUCCACC 194
MBFV-NC_012533.1 -AUAAAACAGCAUAUUGACACCUGGGA--AAGACC--GGAGACUCUGUGGCCUUUCCC 196
MBFV-NC_002031.1 AUAGUGGGACCAUAUUGACGCCAGGGA--AAGACCGGAGGUUCUUCU--GCUUUUCCU 191
MBFV-NC_012735.1 AUA-GCGGACCAUAUUGACGCCAGGGA--AAGACCGGAGACACUCCUU--GAUUCUCAC 192
MBFV-NC_008719.1 AUA-GCGGACCAUAUUGACGCCUGGGA--AAGACCGGAGACACUCCUU--GAUUCUCAC 192
MBFV-NC_001809.1 CCUCCUAUACCAAA-UUCCCCCUCG-ACAGAGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_005062.1 CCUCCUAUACCAAA-UUCCCCCUCU-UCUGAGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_027709.1 CCUCCUAUACCAAA-UUCCCCCUCG-ACAGAGGGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_001672.1 CCUCCUAUACCAAA-UUCCCCCUCG-UCAGAGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_003690.1 CCUCCUAUACCAAA-UUCCCCCUCG-UCAGAGGGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_003687.1 CCUCCUAUACCAAA-UUCCCCCAGG-AAACUGGGGGGGGGCGGUUCUUGU--UCUCCUGA 192
MBFV-NC_018705.3 GGCAACAGCUGUAACCAAGGCAAGGACUAGAGGUAGAGGACCC----- 193
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MBFV-NC_009942.1 A---ACAGCCACACGGACAGUCGCGCA--CAAUGGUGGUG-GUG--GUGCGAGA-AC 249
MBFV-NC_001437.1 AACAUAGCUACUAGGACAGAGCGCGCA--AGUAUUGAGCUG-GUGGUGAGGAAGA-AC 249
MBFV-NC_007580.2 AACACAGCCACAGGGACAGAGCGCGU--AAAGUGGUGUG-GUGGUGAAAAA-A-UC 249
MBFV-NC_034151.1 AACACAGCCACUCCGACAGAACGCCGU--AAGUGUGGUG-GUGGUGAA-AAUAAAC 249
MBFV-NC_009028.2 AUCACAGCCACUAGGACAGAUCCGCGC--AAGUAUGGUGUG-GUGGUGAAAAACACAU 252
MBFV-NC_040776.1 AACACAGCCACUAGGACAGAUCCGCGA--AAUUGUGGUG-GUGGUGCAAAACACA- 252
MBFV-NC_000943.1 AACAUAGCUACUAGGACAGAGCGCGA--ACACUGAGCUG-AUGGGGGAAGAGAC-C 249
MBFV-NC_006551.1 AACAUACCCACAAGGACAGAGCGCGCA--AAAUUGUGGUG-GUGGGGAACUAGAC-C 249
MBFV-NC_012534.1 AACACAGCCAUCCGGACAGAGCGCGU--AAG-UGUGGUG-GUGGUGGUAGA--AC 249
MBFV-NC_015843.2 AACACAGCCAUUGGACAGAGCGCGA--AAG-UUGGUG-GUGGUGGUAGA--AC 249
MBFV-NC_009026.2 AACACUCCACCCGGACAGAACCGCGA--CA-UGUGGGGAG-GGUUCGUAAGACAC-- 250
MBFV-NC_009029.2 GACUUCGGGUCUCGGACAGAACCGCGU--UGA-GAGCCGG-AGGCCAAA-----AC 249
MBFV-NC_032088.1 GAACCCCAUUCUAGGACAGAACCGCG--UAA-UGAUCGG-GGUGCCAAA-----AC 249
MBFV-NC_002640.1 CAACAUCAAUCC-AGGACAGAGCGCGCG--AAGAUUGAUUGG-UGUUGUUG--A-UCCA 249
MBFV-NC_001474.2 CAGCAUCAUUC--AGGACAGAACCGCGA--AAAAUGGAUUGG-UGUGUUG--A-AUCA 249
MBFV-NC_001477.1 CAGCAUCAUUC--AGGACAGAACCGCGA--AAAAUGGAUUGG-UGUGUUG--A-AUCA 249
MBFV-NC_001475.2 CAGCAUCAUUC--AGGACAGAACCGCGA--AAAAUGGAUUGG-UGUGUUG--A-AUCA 249
MBFV-NC_012532.1 ACGCUGGCGCG--AGGACAGAUCCGCGA--ACUUCGGCGCG--GGUGUGGGGA--AUCC 249
MBFV-NC_035889.1 ACGCUGGCGCG--AGGACAGAUCCGCGA--AUAGCGCGCG--GGUGUGGGGA--AUCC 249
MBFV-NC_012533.1 ACACCGAGCCCG--UGGACAGAGCGCG--AGAUCCGGGGGUG--GGUGUGGAGCA-ACCC 249
MBFV-NC_002031.1 CCAGAGGUGUGU--GAGACAGUUUGCU--GAAUGAUAAGAGAC--UUUGAUGACAAAC 249
MBFV-NC_012735.1 CUUUCUCCCUUAAAGACAGAUUGCUU--GAAUGCAGGGUGG--GGAAGUUGGGAACCAA 249
MBFV-NC_008719.1 GGCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_001809.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_005062.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_027709.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_001672.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_003690.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_003687.1 GCCACCAUACCCAGACAGAGUAGUCUGACAAGGAGGUGAUGUGUGACUCGGAAAA-- 249
MBFV-NC_018705.3 -----CUUGCGAUGAACACCACACAAAAACAGCAUAUUGACACCUUGGGAAGACUA 245
40.....250.....260.....270.....280.....290.....3



MBFV-NC_001563.2 ACAGGAUC---U 258
MBFV-NC_009942.1 ACAGGAUC---U 258
MBFV-NC_001437.1 ACAGGAUC---U 258
MBFV-NC_007580.2 ACAGGAUC---U 258
MBFV-NC_034151.1 GGA---UC---U 258
MBFV-NC_009028.2 GGA---UC---U 258
MBFV-NC_040776.1 GGA---UC---U 258
MBFV-NC_000943.1 ACAGGAUC---U 258
MBFV-NC_006551.1 ACAGGAUC---U 258
MBFV-NC_012534.1 ACAGGUC---U 258
MBFV-NC_015843.2 ACAGGAUC---U 258
MBFV-NC_009026.2 -CAGAUUC---U 258
MBFV-NC_009029.2 ACCGAUC---U 258
MBFV-NC_032088.1 ACCGAAUC---U 258
MBFV-NC_002640.1 ACAGGUUC---U 258
MBFV-NC_001474.2 ACAGGUUC---U 258
MBFV-NC_001477.1 ACAGGUUC---U 258
MBFV-NC_001475.2 ACAGGUUC---U 258
MBFV-NC_012532.1 AUGGUUC---U 258
MBFV-NC_035889.1 AACGAGUC---U 258
MBFV-NC_012533.1 AACAGGAC---U 258
MBFV-NC_002031.1 AACAGGAC---U 258
MBFV-NC_012735.1 CUAGGUC---U 258
MBFV-NC_008719.1 CUAGGUC---U 258
MBFV-NC_001809.1 ACACCCGC---U 258
MBFV-NC_005062.1 ACACCCGC---U 258
MBFV-NC_027709.1 ACACCCGC---U 258
MBFV-NC_001672.1 ACACCCGC---U 258
MBFV-NC_003690.1 ACACCCGC---U 258
MBFV-NC_003687.1 ACACCCGC---U 258
MBFV-NC_018705.3 GCAGACCCUCUGU 258
00.....310.

