```
                   ............................................................
MBFV-NC_012532.1   AAC-----AGGUUU------AAUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA 47
MBFV-NC_035889.1   ACA-----GGUUUU------AUUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA 47
MBFV-NC_012533.1   AAC-----AGUUUU------CGUGUUGAGAAAGAAAACAGAUUCUUCGG--AAAUGAAAA 47
MBFV-NC_002640.1   ------------UCUAACAGUUUGUUUGAAUAGAGAGCAG-AUCUCUGGAAAAAUG---A 44
MBFV-NC_001477.1   ACG-----UAGUUCUAACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACA 47
MBFV-NC_001475.2   ACG-----UAGUGCUGACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACA 47
MBFV-NC_001474.2   -CG-----UAGUUCUAACAGUUU-UUUAAUUAGAGAGCAG-AUCUCUG-----AUGAAUA 47
MBFV-NC_034151.1   --------UAACACAGUUGAAUAGUUUUUUGAACAGAGGUGAAAACC-----AUGUCU- 46
MBFV-NC_001563.2   A--U----UAACACAGUGCGAGCUGUUUCUUGGCACGAAGA-UCUC-G-----AUGUCU- 46
MBFV-NC_009942.1   A--U----UAACACAGUGCGAGCUGUUUCUUAGCACGAAGA-UCUC-G-----AUGUCU- 46
MBFV-NC_009028.2   U--U----UAACACAGUCUGAAUAGUUUUUUAGCA-AGGGA-UUUCCC-----AUGUCU- 46
MBFV-NC_040776.1   U--U----UAACACAGUCUGAAUAGUUUUUUAGCA-GAGGU-ACUACC-----AUGUCU- 46
MBFV-NC_007580.2   ---U----AACUUGACUGCGAACAGUUUUUUAGCA-GGGAAUUACCCA-----AUGUCU- 46
MBFV-NC_012534.1   CCGU----AAACACAGUUUGAACGUUUUUUGGAGA----GAGAUUAUU-----AUGUUUA 47
MBFV-NC_015843.2   ACGU----AAACACAGUUUGAACGUUUUUUGGAUA----GAGACAACU-----AUGUCUA 47
MBFV-NC_000943.1   --------UAACGCGGUUUGAACAGUUUUUUGGAGCUUUUGAUUUCAA-----AUGUCU- 46
MBFV-NC_006551.1   ---U----UAACACAGUGCCGGCAGUUUCUUUGAGC-GUUGAUUUUCA-----AUGUCU- 46
MBFV-NC_001437.1   ---U----UAGUGCAGUUUAAACAGUUUUUUAGAACGGAAGAUAACC------AUGACU- 46
MBFV-NC_009026.2   ---------AGAAAAGUUUUCGAGGAAAGACGAGAAGCGAA-UUCUCUG---AAUGGCU- 46
MBFV-NC_009029.2   UGAUUACAUAACACAGUGUGAACAAGUUUUUUGAACAG----CA---------AUGACC- 46
MBFV-NC_032088.1   UC----UAUAACUCAGUGUGAACAAGUUUUUUGAUCAGUGGAUA---------AUGACU- 46
MBFV-NC_002031.1   -----------UAAUCGUUCGUUGAGC--GAUUAGCAGAGAACUGACCAGAACAUGUC-- 45
MBFV-NC_008719.1   -----------AUUUCUACUGAAAGGGUAGAGAA--AAGGAGUUUUGCUUCUCAUGGCAA 47
MBFV-NC_012735.1   -----------UUUCUACUG-AAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAAUGGCAA 47
MBFV-NC_018705.3   CCGU----AAACACAGUUUGAACGUUUUUUGGAGAGAGACUACU---------AUGUCAA 47
                   .........10........20........30........40........50........6
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                   ..........((((...((....((......((((((.(((((((((((...........
MBFV-NC_012532.1   ACCCCAAAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAG 101
MBFV-NC_035889.1   A-CCCAAAAAAGAAAU--CC---GGAGGAUUCCGGAUUGUCAAUAUGCUAAAACGCGGAG 101
MBFV-NC_012533.1   --CCCAAAAAAGG----------GCCGCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUG 95
MBFV-NC_002640.1   A--CCAACGAAAAAAG--GU---G--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGA 95
MBFV-NC_001477.1   A--CCAACGGAAAAAG--AC---G--GGUCGACCGUCUUUCAAUAUGCUGAAACGCGCGA 98
MBFV-NC_001475.2   A--CCAACGGAAGAAG--AC---G--GGAAAACCGUCUAUCAAUAUGCUGAAACGCGUGA 98
MBFV-NC_001474.2   A--CCAACGGAAAAAG--GC---G--AAAAACACGCCUUUCAAUAUGCUGAAACGCGAGA 98
MBFV-NC_034151.1   --AAAAAACCAGGAGGACCC---GGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGCACAU 101
MBFV-NC_001563.2   --AAGAAACCAGGAGGGCCC---GGUAAAAACCGGGCUGUCAAUAUGCUAAAACGCGGUA 101
MBFV-NC_009942.1   --AAGAAACCAGGAGGGCCC---GGCAAGAGCCGGGCUGUCAAUAUGCUAAAACGCGGAA 101
MBFV-NC_009028.2   --AAAAAACCAGGAAAGUCC---GCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGG 101
MBFV-NC_040776.1   --AAAAAACCAGGAGGACCC---GCCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAG 101
MBFV-NC_007580.2   --AAAAAACCAGGAAAACCC---GGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGGCG 101
MBFV-NC_012534.1   ACAAAAAACCAGGAAAACCC---GGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUG 104
MBFV-NC_015843.2   ACAAAAAACCAGGAAGACCC---GGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAA 104
MBFV-NC_000943.1   --AAAAAACCAGGAGGACCC---GGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGGCA 101
MBFV-NC_006551.1   --AAGAAACCAGGAGGGCCC---GGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGGCA 101
MBFV-NC_001437.1   --AAAAAACCAGGAGGGCCC---GGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCC 101
MBFV-NC_009026.2   --AGAAAACCAGGAAGACCC---GGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGG 101
MBFV-NC_009029.2   --AAAAAACCAGGAAGGCCC---GGUAGAAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUG 101
MBFV-NC_032088.1   --AAAAAACCAGGAAAGUCC---GGCAAUCGUCGGACUGUCAAUAUGCUAAAGCGCGGAG 101
MBFV-NC_002031.1   -------UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUA---CGACGAG 95
MBFV-NC_008719.1   CAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGUG--ACGGCCAA 105
MBFV-NC_012735.1   CAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUA--GCGGCUAG 105
MBFV-NC_018705.3   ACAAAAAACCAGGAAGACCC---GGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGCG 104
                   0........70........80........90........100.......110.......1
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                   ............................................................
MBFV-NC_012532.1   UAGCCCGUG------NNNNNNN--------GUUAGAGGAGACCCCCCGGAAAACGCAAAA 147
MBFV-NC_035889.1   UAGCCCGUG------NNNNNNN---------UUAGAGGAGACCCCCCGGAAAACGCAAAA 146
MBFV-NC_012533.1   GAUCUAGCGCUGGCCNNNNNNN--------AGACCCCCCGCGCCCAUAACCAACAUAAAA 147
MBFV-NC_002640.1   GAAACCGCGUAUCAANNNNNNNC--UAGAGGUUAGAGGAGACCCCCCCAACA--CAAAAA 151
MBFV-NC_001477.1   GAAACCGCGUGU---NNNNNNN----AGAGGUUAGAGGAGACCCCCCGCACAACAACAAA 151
MBFV-NC_001475.2   GAAACCGUGUGU---NNNNNNNC--UAGAGGUUAGAGGAGACCCCCCG--CAAAUAAAAA 151
MBFV-NC_001474.2   GAAACCGCGUGU---NNNNNNN---UAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAAA 151
MBFV-NC_034151.1   CCCGCGCUA------NNNNNNN-------GACCCCUUGGCAGUGAAC-GCAAA-UAAAAA 146
MBFV-NC_001563.2   UGCCCCGCG------NNNNNNNA---GGAGACCCCGUGCCA-AAAACACCA-A-AAGAAA 149
MBFV-NC_009942.1   UGCCCCGCG------NNNNNNN-----GAGACCCCGUGCCACAAAACACCACA-ACAAAA 149
MBFV-NC_009028.2   CUAGCGUGU------NNNNNNNU---GGAGACCCCUCAGCAC-CAAGCGCGAA--ACAAA 149
MBFV-NC_040776.1   CCAGCGUUU------NNNNNNN----GGAGACCCCUUGGCACACAAGCGCAAA-GACAAA 150
MBFV-NC_007580.2   UAUCCCGCG------NNNNNNNUAGAGGAGACCCCUUGCCGUUAACGC------AAACAA 149
MBFV-NC_012534.1   CAUCCC---------NNNNNNN-----GAGACCCCUUGCGAGUUAACACCAAA-ACAAAA 149
MBFV-NC_015843.2   CGUCCC---------NNNNNNN----GGAGACCCCUUGCGAGUGAACACCACA-A-GAAA 149
MBFV-NC_000943.1   UACCCCGCG------NNNNNNN--------ACCCCACUCUCAAAAGCAUC-AA-A--CAA 143
MBFV-NC_006551.1   UACCCCGCG------NNNNNNN----------CCCCGCAUCAUAAGCAUCAAA-A--AAA 142
MBFV-NC_001437.1   UACCCCGCG------NNNNNNN--------AGAGGAGACCCCGCAUUUGCAUC-A---AA 143
MBFV-NC_009026.2   CGGCCAACG------NNNNNNNCCCUUGGCAAAACAGUUAACGCA----CCAA-AAGAAA 150
MBFV-NC_009029.2   CGUCCCGCG------NNNNNNNGAGG--AGACCCUCCCACAAAGAAGCGCAAACACAAAA 153
MBFV-NC_032088.1   CAUCCCGCG------NNNNNNNGAGGUUAGAGGAGACCCCCACAUACCGCAAA-ACAAAA 154
MBFV-NC_002031.1   GAGUUCGCUCCUUGUNNNNNNN-------------AGACCCUCCAGGGAACAAAUAGUGG 142
MBFV-NC_008719.1   G----------GCCANNNNNNN----------AGGAGACCCUCCCGAGCAC-UAUAGC-G 143
MBFV-NC_012735.1   A----------GUGANNNNNNN-----------CCAUUCCCUCCCGAGCACACAUAGC-G 143
MBFV-NC_018705.3   CAUCCC---------NNNNNNNCCCGCGGAAAUCAAAGCAAGGCCCAACCUG--GCAAAG 153
                   20.......130.......140.......150.......160.......170.......1
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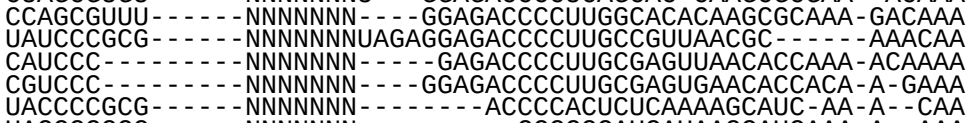

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                   .))))))))))))))))).......)).))...)))).....................
MBFV-NC_012532.1   CAGCAUAUUGACG--UGGGAA--AGACCAG-AGACUCCAUGAGUUUCCACCACGCUGGCC 202
MBFV-NC_035889.1   CAGCAUAUUGACG-CUGGGAA--AGACCAG-AGACUCCAUGAGUUUCCACCACGCUGGCC 202
MBFV-NC_012533.1   CAGCAUAUUGACACCUGGGAA--AGACCGG-AGACUCUGUUGGCUUUCCCCACACCAGCC 204
MBFV-NC_002640.1   CAGCAUAUUGACG-CUGGGAA--AGACCAG-AGAUCCUGCUG-UCUCUGCAACAUCAAUC 206
MBFV-NC_001477.1   CAGCAUAUUGACG-CUGGGAG--AGACCAG-AGAUCCUGCUG-UCUCUACAGCAUCAUUC 206
MBFV-NC_001475.2   CAGCAUAUUGACG-CUGGGAG--AGACCAG-AGAUCCUGCUG-UCUCCUCAGCAUCAUUC 206
MBFV-NC_001474.2   CAGCAUAUUGACG-CUGGGAA--AGACCAG-AGAUCCUGCUG-UCUCCUCAGCAUCAUUC 206
MBFV-NC_034151.1   CAGCAUAUUGACACCUGGGAU--AGACUGGGAGACCCUCUGCUU-AUUUCAACACCAGCC 203
MBFV-NC_001563.2   CAGCAUAUUGACACCUGGGAU--AGACUAGGGGAUCUUCUGCU---CUGCACAACCAGCC 204
MBFV-NC_009942.1   CAGCAUAUUGACACCUGGGAU--AGACUAGGAGAUCUUCUGCU---CUGCACAACCAGCC 204
MBFV-NC_009028.2   CAGCAUAUUGACGCCUGGGAA--AGACCGGGAGAUCCUCUGCU---UUCCAUCACCAGCC 204
MBFV-NC_040776.1   CAGCAUAUUGACACCUGGGAA--AGACCGGGAGACCCUCUGCU---CUGCAACACCAGCC 205
MBFV-NC_007580.2   CAGCAUAUUGACACCUGGAAA----GACAGGAGAUCCCCUGCUU--UUUCAACACCAGCC 203
MBFV-NC_012534.1   CAGCAUAUUGACACCUGGGAG--AGACUAGGAGACCCUCUGUCC--UAACAACACCAGCC 205
MBFV-NC_015843.2   CAGCAUAUUGACACCUGGGAU--AGACUAGGAGACCCUCUGUCC--UAACAACACCAGCC 205
MBFV-NC_000943.1   CAGCAUAUUGACACCUGGGAA-AAGACUAGGAGAUCUUCUGCUCUAUUCCAACAUCAGUC 202
MBFV-NC_006551.1   CAGCAUAUUGACACCUGGGAAUUAGACUAGGAGAUCUUCUGCUCUAUUCCAACAUCAACC 202
MBFV-NC_001437.1   CAGCAUAUUGACACCUGGGAAU-AGACUGGGAGAUCUUCUGCUCUAUCUCAACAUCAGCU 202
MBFV-NC_009026.2   CAGCAUAUUGACACCUGGGAU--AGACCGGAGAAUUUGCUGCC---UCGCAACACCUCCC 205
MBFV-NC_009029.2   CAGCAUAUUGACACCUGGGAA--AGACUAGGGGAUUUGCUGCU---CUGGACUUCCGGCU 208
MBFV-NC_032088.1   CAGCAUAUUGACACCUGGGAU--AGACUAGGAGAUCCGCUGCU---CUGGAACCCCAG-U 208
MBFV-NC_002031.1   GACCAUAUUGACGCCAGGGAAAGACCGGAGUGGUUCUCUGCUUUUCCUCCAGAGGUCUGU 202
MBFV-NC_008719.1   GACCAUAUUGACGCCUGGGAAAGACCGGAGACACUCCUUGAUUCUCACCUUUCUCACCCU 203
MBFV-NC_012735.1   GACCAUAUUGACGCCAGGGAAAGACCGGAGACACUCCUUGAUUCUCACCUUUCUCGCCCU 203
MBFV-NC_018705.3   CUGUAAACCAGGGGAAGGACUAGAGGUUAGAGGAGACCCCUUG----------CGAGUG 202
                   80.......190.......200.......210.......220.......230.......2
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                   ......
MBFV-NC_012532.1   GCCAGG 208
MBFV-NC_035889.1   GCCAGG 208
MBFV-NC_012533.1   CCC--U 208
MBFV-NC_002640.1   C----A 208
MBFV-NC_001477.1   C----A 208
MBFV-NC_001475.2   C----A 208
MBFV-NC_001474.2   C----A 208
MBFV-NC_034151.1   ACUC-G 208
MBFV-NC_001563.2   ACAC-- 208
MBFV-NC_009942.1   ACAC-- 208
MBFV-NC_009028.2   ACUA-- 208
MBFV-NC_040776.1   ACU--- 208
MBFV-NC_007580.2   ACAA-G 208
MBFV-NC_012534.1   AUC--- 208
MBFV-NC_015843.2   ACU--- 208
MBFV-NC_000943.1   ACAAGG 208
MBFV-NC_006551.1   ACAAGG 208
MBFV-NC_001437.1   ACUAGG 208
MBFV-NC_009026.2   ACC--- 208
MBFV-NC_009029.2   ------ 208
MBFV-NC_032088.1   ------ 208
MBFV-NC_002031.1   GAGCAC 208
MBFV-NC_008719.1   UAA-GC 208
MBFV-NC_012735.1   AAA-GC 208
MBFV-NC_018705.3   AACACC 208
                   40....
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