```
                                                   (((.(((((..............((.(((((.......((.((((((.((((((((((((..
MBFV-NC_012532.1  AACAGGU-U-------UAAUUUGGAUUUGGAAA--CGA-GAGUUUCUGGUCAUGAAAA-ACCCCAAAGAAGAAAAUC--CGGAGGA-UCCGGAUUGUCAAUAUGCUAA 92
MBFV-NC_035889.1  ACAGGUU-U-------UAUUUUGGAUUUGGAAA--CGA-GAGUUUCUGGUCAUGAAAA-CCCAAAAAAGAAAAUC--CGGAGGAUUCCGGAUUGUCAAUAUGCUAA 92
MBFV-NC_002640.1  -------UCUAACAGUUUUGUUUGAAUAGAGAG--CAG-AUCUCUGGAAAAAUG---A-A--CCAACGAAAAAAGG--UGGUUAGACC--ACCUUUCAAUAUGCUGA 86
MBFV-NC_001477.1  ACGUAGU-UCUAACAGUUU-UUU-AUUAGAGAG--CAG-AUCUCUG-----AUGAACA-A--CCAACGGAAAAAGA--CGGGUCGACC--GUCUUUCAAUAUGCUGA 89
MBFV-NC_001475.2  ACGUAGU-GCUGACAGUUU-UUU-AUUAGAGAG--CAG-AUCUCUG-----AUGAACA-A--CCAACGGAAGAAGA--CGGGAAAACA--GUCUUUCAAUAUGCUGA 89
MBFV-NC_001474.2  -CGUAGU-UCUAACAGUUU-UUUAAUUAGAGAG--CAG-AUCUCUG-----AUGAAUA-A--CCAACGGAAAAAGG--CGAAAAACAC--GCCUUUCAAUAUGCUGA 89
MBFV-NC_012533.1  A-----------------ACAGUUUCGUGUUGAGAAAGAAAACAGAUUCUUCGGAAAUGAA---------AACCCAAAAAGG--GGCCGCAUCGCGGCCUAUCAAUAUGCUGA 86
MBFV-NC_009028.2  U--U----UAACACAGUCUGAAUAGUUUUUU--GGCAAGGA-UUUC--CCAUGUCUA----AAAAACCAGGGAAAGUCCGCUGCAAAGCGGACUGUCAAUAUGCUAA 92
MBFV-NC_040776.1  U--U----UAACACAGUCUGAAUAGUUUUUU--GACAGAGGU-ACUA--CCAUGUCUA----AAAAACCAGGGAAGACCCGCCGGAAGCGGGGUUGUCAAUAUGCUAA 92
MBFV-NC_034151.1  ------UAACACAGUGUGAAUAGUUUUUU--GGACAGAGGUGUGAA---CCAUGUCUA----AAAAACCAGGAGGACCCGGGAAGAACCGGGUCGUUAAUAUGCUGA 92
MBFV-NC_000943.1  -------UAACGCGGUUUGAACAGUUUUUUUGGAGCUUUUGA-UUUC--AAAUGUCUA----AAAAACCAGGAGGACCCGGGAAGCCCCGGGUCGUCAAUAUGCUGA 92
MBFV-NC_006551.1  ---U----UAACACAGUGCCGGCAGUUUCUUUGAGC--GUUGA-UUUU--CAAUGUCUA----AAAAACCAGGAGGGCCCGGAAGAAACCGGGCCAUCAAUAUGCUGA 92
MBFV-NC_001563.2  A--U----UAACACAGUGCGAGCUGUUUCUUUGGCAC-GAAGA-UCUC--G-AUGUCUA----AGAAACCAGGAGGGCCCGGUAAAAAGCGGGCUGUCAAUAUGCUAA 92
MBFV-NC_009942.1  A--U----UAACACAGUGCGAGCUGUUUCUUUGGCAC-GAAGA-UCUC--G-AUGUCUA----AGAAACCAGGGCCCGGCAAGAGCCGGGCUGUCAAUAUGCUAA 92
MBFV-NC_001437.1  ---U----UAGUGCAGUUUAAACAGUUUUUUUAGAACGGAAGA-UAAC--C-AUGACUA----AAAAACCAGGAGGGCCCGGUAAAAACCGGGCUAUCAAUAUGCUGA 92
MBFV-NC_007580.2  ---U----UAACUUGACAGUGUUUGAUAGUUUUU-ACCC--A-AUGUCUA----AAAAACCAGGAGGACCCGGUAAAUGGUUGUCAAUAUGCUGA 92
MBFV-NC_012534.1  CCGU----AAACAGUUUUGAAC-GUUUUUUGGAG---AGAGAUUA--UUAUGUUUA-ACAAAAACCAGGGGCAACCCGGGCAAUCAAUAUGCUAA 95
MBFV-NC_015843.2  ACGU----AAACACAGUUUUGAAC-GUUUUUUGGAU----AGAGACAA--CUAUGUCUA-ACAAAAACCAGGAGACCCGGCUCAGGCCGGGUUGUCAAUAUGCUAA 95
MBFV-NC_009026.2  ---------AGAAAAGUUUUCGAGGAAAGACGAGAAGCGAAUUCUCU--GAAUGGCUA----GAAAACCAGGAGACCCGGAGGAAACCGGGUCGUCAAUAUGCUAA 92
MBFV-NC_009029.2  UGAUUACAAUAUCAGUGUGAACAAGUUUUUUGAUCAGUGGA----------CAAUGACCA----AAAAACCAGGAGGACUGGCAAUCGUCGGGCCGUCAAUAUGCUAA 92
MBFV-NC_032088.1  UC----UAUAACUCAGUGUGAACAAGUUUUUUUGAUCAGUGGA------UAAUGACUA----AAAAACCAGGGAAAGUCCGGCAAUCGUCGGGACUGUCAAUAUGCUAA 92
MBFV-NC_012735.1  --------------UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAUGGCAACAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGU 96
MBFV-NC_008719.1  -----------AUUUCUACUGAAAGG-GUAGAGAAAAGGAGUAUUUUGCUUCUCAUGGCAACAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGU 96
MBFV-NC_002031.1  ---------UAAUCGUUCGUUGAGCG-AUUAGACAGAGGAGACUA-------CUAUGUC---AACAAAAAACAAAUAGUGGUCGUAAAGCUCAAGAAAAAACCCGGGUUGUCAAUAUGCUAA 87
MBFV-NC_018705.3  CCGUA----AACACAGUUUGAACGUUUUUUUUGGAGAGAGACUA-------CUAUGUCAA----AAAAAACCAGGAGACCCGGCAAUCGUCGGGUUGUCAAUAUGCUAA 95
          .........10........20........30........40........50........60........70........80........90........100.....

                       ..((((((......))))))...............((((......))))..........))))))))))))))))))))).)))))))...)))
MBFV-NC_012532.1  AACGCGGAGUAGCCCGUG------NNNNNNN--------GGUUAGAGGAGACCCCCCGGAAAACGCAAAACAGCAUAUUGACG--UGGGA--AAGACCA-GAGACUC 180
MBFV-NC_035889.1  AACGCGGAGUAGCCCGUG----NNNNNNN--------GUUAGAGGAGACCCCCCGGAAAACGCAAAACAGCAUAUUGACG-CUGGGA--AAGACCA-GAGACUC 180
MBFV-NC_002640.1  AACGCGAGAGAAACCGCGUAUCAANNNNNNNN---ACUAGAGGUUAGAGGAGACCCCCCCAACA--CAAAACAGCAUAUUGACG-CUGGGA---AAGAC---CAGAGAU 183
MBFV-NC_001477.1  AACGCGUGAGAAACCGCGUGUG--NNNNNNN----UAGAGGUUAGAGGAGACCCCCGGAAAACGCAAAACAGCAUAUUGACG-CUGGGA---GAGAC---CAGAGAU 183
MBFV-NC_001475.2  AACGCGUGAGAAACCGUGUGUG---NNNNNNN----ACUAGAGGUUAGAGGAGACCCCCG--CAAAUAAAAACAGCAUAUUGACG-CUGGGA---GAGAC---CAGAGAU 183
MBFV-NC_001474.2  AACGCGAGAGAAACCGCGUGU---NNNNNNN----CUUAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAACAGCAUAUUGACG-CUGGGA---AAGAC---CAGAGAU 183
MBFV-NC_012533.1  AGCGCUUUGGAUCUAGCGCUGGCCNNNNNNNN--GAGACCCCCC------GCGUCCAUAACCAUAAAACAGCAUAUUGACACCUGGGA--AAGACCG-GAGACCU 182
MBFV-NC_009028.2  AACGCCAGCCUAGCGUGU----NNNNNNN---GUGGACACCCUCAGCAC-CAAGCGCGAA--ACAAACAGCAUAUUGACGCGCUGGGA--AAGACCGGGGAGAUCC 185
MBFV-NC_040776.1  AACGCCCAGCCAGCGUUU----NNNNNNN---A-GGAGACCCCUUGGCACACAAGCGCAAA-GACAAACAGCAUAUUGACACCUGGGA--AAGACCGGGGAGAUCC 186
MBFV-NC_034151.1  AGCGCCACAUCCCGCGCUA----NNNNNNN----AGACCCCACUCUCAAAAGCAUC---AAACAACAGCAUAUUGACACCUGGGA---UAGACUGGGGAGACCC 182
MBFV-NC_000943.1  AACGCGGCAUACCCCGCG-----NNNNNNN-----GACCCCACUCUCAAAAGCAUC---AAACAACAGCAUAUUGACACCUGGGA--AAGACUAGGGAGAUCU 180
MBFV-NC_006551.1  AACGCGGCAUACCCCGCG-----NNNNNNN-----ACCCCGCA-UCAUAAGCAUC---A-AAAAAACAGCAUAUUGACACCUGGGAAUUAGACUAGGGAGAUCU 180
MBFV-NC_001563.2  AACGCGGUAUGCCCCGCG-----NNNNNNN--GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGGAUCU 185
MBFV-NC_009942.1  AACGCGGUAUGCCCCGCG-----NNNNNNN--GGAGAACCCCAAAACCACA-AUGUCUAGCAUAUUGACACCUGGGA--UAGACUAGGGGAUCU 185
MBFV-NC_001437.1  AACGCGGCCUAGCCCGCG-----NNNNNNN-UAGAGGAGACCCC--GCAUUU-------GC-AUCAACAGCAUAUUGACACCUGGGAA--UAGACUGGGGAGAUCU 180
MBFV-NC_007580.2  AACGCGGCGUAUCCCGCG-----NNNNNNNUUAGAGGAGACCCCUUGCCGUUAA-----CGC-AACAACAGCAUAUUGACACCUGGGAA---AGACAG---GAGAU 181
MBFV-NC_012534.1  AGCGCGGUGCAUCCC------NNNNNNN------GGAGACCCCUUGCGAGUUAACACCAAA-ACAAAACAGCAUAUUGACACCUGGGA--GAGACUAGGGGAGACCC 185
MBFV-NC_015843.2  AGCGCGAACGUCCC-------NNNNNNN------GGAGACCCCUUGCGAGUGACACCACCA-A-GAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGAGACCC 185
MBFV-NC_009026.2  AGCGUACGGCGGCCAACG-----NNNNNNNCCCCUUGG----CAAAACAGUUAACGCACCAA-AAGAAACAGCAUAUUGACACCUGGGA--UAGACC-GGAGAAUU 185
MBFV-NC_009029.2  AGCGCGGUGCGGUCCCGCG----NNNNNNNAGAGG--AGACCCUCCCACAAAGAAGCGCAAACAAAACAGCAUAUUGACACCUGGGA--AAGACUAGGGGAUUU 189
MBFV-NC_032088.1  AGCGCGGAGCAUCCCGCG-----NNNNNNNAGAGGUUAGAGGAGACCCCCACAUACCGCAAA-ACAAAACAGCAUAUUGACACCUGGGA---UAGACUAGGGAGAUCC 190
MBFV-NC_012735.1  AGCGGCUAGA----------GUGANNNNNNN--------C--CCAUUCCCUCCCGAGCACACAUAGC-GGACCAUAUUGACGCGCCUGGGA----AAGACC-GGAGACAC 178
MBFV-NC_008719.1  GACGGCCAAG----------GCCANNNNNNN----------GAGGAGACCCUCCCGAGCAC-UAUAGC-GGACCAUAUUGACGCGCCUGGGA----AAGACC-GGAGACAC 178
MBFV-NC_002031.1  -ACGACGAGGAGUUCGCUCCCUUGUNNNNNNN----------GAGACCUCCAGGGAACAAAUAGUGGGCACCAUAUUGACGCGCCAGGGA----AAGACC-GGAGUGGU 177
MBFV-NC_018705.3  AGCGCGGCGCAUCCC---------NNNNNNNCCCCGCGGAAAUCAAAGCAAGGCCCAACCUG--GCAAAGCUGUAAACGGGAAGGACUAGGAGGUUAGAGGAGAC 191
          ..........110.......120.......130.......140.......150.......160.......170.......180.......190.......200.......210..

                        .)))).................((((((((.((((.......)))...))))))))...........
MBFV-NC_012532.1  CAUGAGUUUCCA-CCACGCUGGCCGCCAGGCACAGAUCGCCGAACUUCGGCGGCGGGUGUGGG-GAAAUCCAUG-GUUUCU 258
MBFV-NC_035889.1  CAUGAGUUUCCA-CCACGCUGGCCGCCAGGCACAGAUCGCCGAAUAGCGGCGGCGGGUGUGGG-GAAAUCCAUG-GGUCUU 258
MBFV-NC_002640.1  CCUG-CUGUCUC-UGCAACACUCAAUCCAGGCACAGAUCGCCGCAAGAUGGGAUGGGUGUUGUG---AUCCAACA-GGUUCU 258
MBFV-NC_001477.1  CCUG-CUGUCUC-UACAGCAUCAAUUCAGGCACAGAACGCCAAAAAAUGGGAAUGGGUGCUGUUG---AAUCAACA-GGUUCU 258
MBFV-NC_001475.2  CCUG-CUGUCUC-UCUCAGCAUCAUUCAGGCACAGAACGCCAGAAAAUGGGAAUGGGUGCUGUUG---AAUCAACA-GGUUCU 258
MBFV-NC_001474.2  CCUG-CUGUCUC-UCUCAGCAUCAUUCAGGCACAGAACGCCAGAAAAUGGGAAUGGGUGCUGUUG---AAUCAACA-GGUUCU 258
MBFV-NC_012533.1  UGUUGGCUUUCC-CCACACCAGCCCCCUGGCACAGAGCGCCAGA----UC-GGGGGUUGGGUGUGGAGCAACCCAAC-AGGACU 258
MBFV-NC_009028.2  UCUCUCUG----CAUCACCAGCCACUAGGCACAGAUCGCCG-CAAGUA-GUGGGUGGGUGUG---AAAAACAUGGAUCU 258
MBFV-NC_040776.1  UCUGCUCUG----CAACACAGCCACUAGGCACAGAUCGCCG-AAAUCU-GUGGCUGGGUGUUG---CAAAACACA-GGAUCU 258
MBFV-NC_034151.1  UCUGCUUAUUU--CAACACAGCCACUCGGCACAGAACGCCG-UAAGU--GUGGCUGGUGGUGAAAAUAAACACAGAG-UCU 258
MBFV-NC_000943.1  UCUGCUCUAUUC-CAACAUCAGUCACAAGGCACCGAGUCCG-AACACU-GUGACUGAUGGGGGAGAAGACCACA-GGAUCU 258
MBFV-NC_006551.1  UCUGCUCUA----CAACAUCAAACCACAAGGCACAGAUCGCCG-AAAAUU-GUGGCUGGGUGUAGACCACCA-GGAUCU 258
MBFV-NC_001563.2  UCUGCUCU---G-CACAACCAGCCACAAGGCACAGUGCGCC-ACAUAG-GUGGGUGGUGGUGCU--AGAACACA-GGAUCU 258
MBFV-NC_009942.1  UCUGCUCU---G-CACAACCAGCCACAAGGCACAGUGCGCC-ACAUAG-GUGGGUGGUGGUGCG--AGAACACA-GGAUCU 258
MBFV-NC_001437.1  UCUGCUCUAUCU-CAACAUCAGCUACUAGGCACAGAACGCCG-AAGUAU-GUAGCUGGUGUGGAGAAGAACACA-GGAUCU 258
MBFV-NC_007580.2  CCCUGCUCUA---CAACACCAGCCACUAGGCACAGAGCGCCG-UAAAGU-GUGGCUGGUGGUGUAAAA-AUCACA-GGAUCU 258
MBFV-NC_012534.1  UCUGUCCUA--A-CAACACCAGCCAUCCGGCACAGAGCGCCG-UAAG-U-GUGGCUGGGUGUG--GUAGACACA-GUGUCU 258
MBFV-NC_015843.2  UCUGUCCUA--A-CAACACCAGCCACUUGGCACAGAUCGCCG-AAAG-U-GUGGCUGGGUGUG--GUAGACACA-GGGUCU 258
MBFV-NC_009026.2  UGCUGCCU--CG-CAACACCUUCCCCCCGGCACAGAACGCCG-ACA-UG-GUGGGAGGGGUCG---UAAGACACCAGAUUCU 258
MBFV-NC_009029.2  GCUGCUCU-----GGAACCCGGCAGUUCUCGGCACAGAACGCCG-UUGA-U-GAACUGGGUGCC----AAAACACC-GAAUCU 258
MBFV-NC_032088.1  GCUGCUCU-----GGAACCCGGCAGUUCUCGGCACAGAACGCCG-UAA-U-GAACUGGGUGCC----AAAACACC-GAAUCU 258
MBFV-NC_012735.1  UCCUUGAUUCUCACCUUUCUCGCCCUAAAGCACAGAUUGCUU--GAAUGCAGGGCGGGGAAGUUGAGAACCAACUAGUGUCU 258
MBFV-NC_008719.1  UCUUGAUUCUUUCCUCCAGAGUCUGU-GAGCACAGUUUCUCAAGAAUAAAGCAGACCUUUGGAUGACAAACACAAAACCACU 258
MBFV-NC_002031.1  UCUCUGCUUUUCCUUCCAGAGUCUGU-GAGCACAGUUUCUCAAGAAUAAAGCAGACCUUUGGAUGACAAACACAAAACCACU 258
MBFV-NC_018705.3  CCCUUGCGAGUGAACACCACACAAAAACAGCAUA-----UUGACACCUGGGAAAGACUAGGAGACCC----------UCUGU 258
          .........220.......230.......240.......250.......260.......270.......280.......290....
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