

MBFV-NC_002031.1	-----UAAUCGUUCGUUGAGC--GAUUAGCAGAGAACUGACCAG-AACAUGUC-----UGGUC--GUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUAACGACGAGGAGUUCGCUCCUUGU--NNNNNNNN-----GAGACCCUCCAGGGAAACAAAUAGUGGGACCAUAUGACGCCAGGGA--AAGACCGGAGUGGUUCUCUGCUUUUC-CUCCAGAGGUUCUGUGAGCACAGUUGUCUCAAGAAUAAGCAGACCUCUUGGAUGACAAACACAAACCACU	258
MBFV-NC_012735.1	-----UUUCUACUG-AAAGGAGUAGAAGAAAGGAGAUUCAUUC-CCAUGGCAACAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCGAGGGCGUCAAUAUGGU--AGCGGCUAGAGU-----GA--NNNNNNNN-----C-CCAUUCCCUCGAGACACACAUAGC-GGACCAUAUGACGCCAGGGA--AAGACCGGAGACACUCCUUGAUUCUCACCUUUCUCGCCCUAAAGCACAGAUUGCUU--GAAUGCAGGGCGGGGAAGUUGAGAACCAACUAGUGUCU	258
MBFV-NC_008719.1	-----AUUUCUACUGAAAGGGUAGAGAA--AAGGAGUUUUGCUU-CUCAUGGCAACAAGAGGAGCAAGCAAGUCUCGGGUCACGACCAGGGGCGUCAAUAUGGU--GACGGCCAAGGC-----CA--NNNNNNNN-----GAGGAGACCCUCCGAGCAC-UAUAGC-GGACCAUAUGACGCCUGGGA--AAGACCGGAGACACUCCUUGAUUCUCACCUUUCUCACCCUUAAGCACAGAUUGCUU--GAAUGCAGGGUGGGGAAGUUGGGAACCAACUAGUGUCU	258
MBFV-NC_012532.1	AA-----CAGGUUA-----AUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA--ACCCCAAAGAAGAAAUCC--GGAGGA-UCGGAUUGUCAAUAUGCUAAAACGC--GGAG-UAGCCC--GUGNNNNNNNN-----GGUUAGAGGAGACCCCCCGGAAAACGCAAAACAGCAUAUGACG--UGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCACGCUGGCCGCCAGGCACAGAUUCGCC--GAACUUCGGCGGCCGGUGUGGGGAAAUCCAUGGUUUUCU	258
MBFV-NC_035889.1	AC-----AGGUUUUA-----UUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA--A-CCCAAAAAAGAAAUCC--GGAGGAUUCGGAUUGUCAAUAUGCUAAAACGC--GGAG-UAGCCC--GUGNNNNNNNN-----GUUAGAGGAGACCCCCCGGAAAACGCAAAACAGCAUAUGACG-CUGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCACGCUGGCCGCCAGGCACAGAUUCGCC--GAAUAGCGGCGGCCGGUGUGGGGAAAUCCAUGGGUCU	258
MBFV-NC_012533.1	AA-----CAGUUUUC-----GUGUUGAGAAAGAAAAACAGAUUCUUCGG--AAAUGAAAA--CCCAAAAAAGG-----GCCGCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGAUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAACAUAAAACAGCAUAUGACACCUGGGA--AAGAC-CGGAGACUCUGUUGGCUUUCGCCACACCAGCCCCUGGCACAGAGCGCC--AGAUCGGGGGUUGGUUGGAGCAACCCAACAGGACU	258
MBFV-NC_001477.1	AC-----GUAGUUCUACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACA-A-CCAACGGAAAAAGAC--G--GGUCGACCGUCUUCAAUAUGCUGAAACGCGCGAGAAACCGCGUGU--NNNNNNNN--UAGAGGUUAGAGGAGACCCCCCGCACAAACAAACAGCAUAUGACG-CUGGGA--GAGAC-CAGAGAUCCUGCU-GUCUCUACAGCAUCA--UCCAGGCACAGAACGCC--AAAAAUG--GAAUGGUGCUGUUGAAUCAACAGGUUCU	258
MBFV-NC_001475.2	AC-----GUAGUCUACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACA-A-CCAACGGAAAAAGAC--G--GGAAAAACGUCUAUCAAUAUGCUGAAACGCGUGAGAAACCGUGU--NNNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCGCA-AAUA-AAACAGCAUAUGACG-CUGGGA--GAGAC-CAGAGAUCCUGCU-GUCUCCUACGCAUCA--UCCAGGCACAGAACGCC--AGAAAAUG--GAAUGGUGCUGUUGAAUCAACAGGUUCU	258
MBFV-NC_001474.2	-C-----GUAGUUCUACAGUUU-UUUAAUAGAGAGCAG-AUCUCUG-----AUGAAUA-A-CCAACGGAAAAAGGC--G--AAAAACACGCCUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUGU--NNNNNNNN--CUAGAGGUUAGAGGAGACCCCCCGCA-AACAAAAAACAGCAUAUGACG-CUGGGA--AAGAC-CAGAGAUCCUGCU-GUCUCCUACGCAUCA--UCCAGGCACAGAACGCC--AGAAAAUG--GAAUGGUGCUGUUGAAUCAACAGGUUCU	258
MBFV-NC_002640.1	-----UCUAAACAGUUUGUUUGAAUAGAGAGCAG-AUCUCUGGAAAAAUG--A-A-CCAACGAAAAAAGGU--G--GUUAGAACCACCUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUAUCAANNNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCGCA-CACAA-AAACAGCAUAUGACG-CUGGGA--AAGAC-CAGAGAUCCUGCU-GUCUCUGCAACAUCA--AUCCAGGCACAGAGCGCC--GCAAGAUG--GAUUGGUGUUGUUGAUCCAACAGGUUCU	258
MBFV-NC_012534.1	CC-----GUAAACACAGUUUGAAC-GUUUUUUGGAGA-GAGAU--UAU--UAUGUUUAACAAAAAACAGGAAAAACCC--GGGGCAACCAGGUUGUCAAUAUGCUAAAGCGCGGUGCAUCC--NNNNNNNN--GGAGACCCCUUGCGAGUUAAACACCACA-ACAAAACAGCAUAUGACACCUGGGA--GAGACUAGGAGACCCUCUGUCC--UAACAACACCAGCCAUCCGGGCACAGAGCGCC--GUAAGU-GUGGCUGGUGGUGGUAGAACACAGUG--UCU	258
MBFV-NC_015843.2	AC-----GUAAACACAGUUUGAAC-GUUUUUUGGAUA-GAGAC--AAC--UAUGUCUAAACAAAAAACAGGAAAGACCC--GGCUCAGGCAGGUUGUCAAUAUGCUAAAGCGCGGAACGUCC--NNNNNNNN--AGGAGACCCCUUGCGAGUGAACACCACA-A-GAAACAGCAUAUGACACCUGGGA--UAGACUAGGAGACCCUCUGUCC--UAACAACACCAGCCAUCCGGGCACAGAUUCGCC--GAAAGU-GUGGCUGGUGGUGGUAGAACACAGGA--UCU	258
MBFV-NC_001563.2	A-----UUAACACAGUGCGAGCUGUUUCU-GGCACGAAGAUC-UC-----GAUGUCUA--AGAAACCAGGAGGGCCC--GGUAAAAACAGGGCUGUCAAUAUGCUAAAACGCGUAUGCCCCGCG--NNNNNNNN--GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAAACAGCAUAUGACACCUGGGA--UAGACUAGGGGAUCUUCUGCU--CUGCACAACCAGCCACACGGGCACAGUGCGCC--GACAUAGGUGGCUGGUGGUGCUAGAACACA-GGA-UCU	258
MBFV-NC_009942.1	A-----UUAACACAGUGCGAGCUGUUUCU-AGCACGAAGAUC-UC-----GAUGUCUA--AGAAACCAGGAGGGCCC--GGCAAGAGCAGGGCUGUCAAUAUGCUAAAACGCGGAAUGCCCCGCG--NNNNNNNN--GGAGACCCCGUGCCACAAAACACCACA-ACAAAACAGCAUAUGACACCUGGGA--UAGACUAGGAGAUUCUUCUGCU--CUGCACAACCAGCCACACGGGCACAGUGCGCC--GACAUAGGUGGCUGGUGGUGCGGAGAACACA-GGA-UCU	258
MBFV-NC_009028.2	U-----UUAACACAGUCUGAAUAGUUUUU-AGCA-AGGGAUU-UCC--CAUGUCUA--AAAAACCAGGAAAGUCC--GCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCUAGCGUGU--NNNNNNNN--GUGGAGACCCCUAGCAC-CAAGCGCGAA--ACAAACAGCAUAUGACGCCUGGGA--AAGACCGGAGAUCCUCUGCU--UCCAUACACCAGCCACUAGGCACAGAUUCGCC--GCAAGUAGUGGCUGGUGGUGA-AAAACACAUGGA-UCU	258
MBFV-NC_040776.1	U-----UUAACACAGUCUGAAUAGUUUUU-AGCA-GAGGUAC-UAC--CAUGUCUA--AAAAACCAGGAGGACCC--GCCGGAAGGAGGUUGUCAAUAUGCUGAAACGCCCAGCCAGCGUU--NNNNNNNN--A-GGAGACCCCUUGGCACACAAGCGCAA-GACAAACAGCAUAUGACACCUGGGA--AAGACCGGAGACCCUCUGCU--CUGCAACACCAGCCACUAGGCACAGAUUCGCC--GAAAUUCUGUGGCUGGUGGUGCUGC-AAAACACA-GGA-UCU	258
MBFV-NC_007580.2	-----UAACUUGACUGGCAACAGUUUUU-AGCA-GGGAUUUACC--AAUGUCUA--AAAAACCAGGAAACCC--GGUAGAAACAGGUUGUCAAUAUGCUGAAACCGCGCUAUCCCGCG--NNNNNNNN--UAGAGGAGACCCCUUGCGUUA--CGCAA--CAACAGCAUAUGACACCUGG-A--AAGAC-AGGAGAUCCCUUGCU--UUUCAACACCAGCCACAAGGCACAGAGCGCC--GUAAAGUUGGCUGGUGGUGGUGA-AAAACACAUGGA-UCU	258
MBFV-NC_034151.1	-----UAACACAGUGGAAUAGUUUUUAGACA-GAGGUGAAAC--CAUGUCUA--AAAAACCAGGAGGACCC--GGAAAGAACAGGUCGUCAAUAUGCUGAAGCGCAUCCCGCGUA--NNNNNNNN--AGACCCCUUGGCAGUGAA-CGCAA-UAAAAACAGCAUAUGACACCUGGGA--UAGACUGGAGACCCUCUGCU--AUUCAAACACCAGCCACUCGGGCACAGAACGCC--GUAAGU-GUGGCUGGUGGUGGUGA-AAAACACAAGAGUUCU	258
MBFV-NC_000943.1	-----UAACGCGGUUUGAACAGUUUUUGGAGCUUUUGAU-UCA--AAUGUCUA--AAAAACCAGGAGGACCC--GGGAAGCCCAGGUCGUCAAUAUGCUAAAACGCGCAUACCCCGCG--NNNNNNNN--GACCCACUCUCAAAGCAUC-AA--ACAACAGCAUAUGACACCUGGGA--AAGACUAGGAGAUUCUUCUGCU--UUAACAACUAGUACAAGGCACAGAGCGCC--GAACACUGUGACUGAUGGGGAGAGAACACAGGAUCU	258
MBFV-NC_006551.1	-----UUAACACAGUGCCGGCAGUUUCUUGAGC-GUUGAUU-UUC--AAUGUCUA--AGAAACCAGGAGGGCCC--GGAAGAAACAGGGCCAUAUAUGCUGAAACGCGGCAUACCCCGCG--NNNNNNNN--ACCCGCA-UCAUAAGCAUCAAA--AAAACAGCAUAUGACACCUGGGAUUAGACUAGGAGAUUCUUCUGCUUAUUCCAACAUAACCAACAAGGCACAGAGCGCC--GAAAUUUGUGGCUGGUGGGGAAUAGACCACAGGAUCU	258
MBFV-NC_001437.1	-----UUAGUGCAGUUUAAACAGUUUUUAGAACGGAAGAU-ACC--AUGACUA--AAAAACCAGGAGGGCCC--GGUAAAAACAGGGCUAUAUAUGCUGAAACGCGGCCUACCCCGCG--NNNNNNNN--UAGAGGAGACCCCGCAUUUGC-AU--CAAACAGCAUAUGACACCUGGGA-UAGACUGGAGAUUCUUCUGCUUAUCUACAACUAGCUACUAGGCACAGAGCGCC--GAAGUAGUAGCUGGUGGUGAGGAAGAACACAGGAUCU	258
MBFV-NC_009026.2	-----AGAAAAGUUUCGAGGAAAGACGAGAA-GCGAAUUCUCUG--AAUGGCUA--GAAAACCAGGAAAGACCC--GGAGGAAACAGGUCGUCAAUAUGCUAAAGCGUACGGCGGCCAACG--NNNNNNNN--CCCUUGGCAAAACAGUUAAACGCACCAA-AAGAAACAGCAUAUGACACCUGGGA--UAGACCGGAGAAUUUGCUGCC--UCGCAACACCUCACCACCCGGGCACAGAACGCC--GACA-UGGUGGGAGGGGUCGUAGAGACCAA--GAUUCU	258
MBFV-NC_032088.1	UC--UUAUAUCUAGUGUGAACAGUUUUUGAUAGUGGAU-----AAUGACUA--AAAAACCAGGAAAGUCC--GGCAAUUCGUGGACUGUCAAUAUGCUAAAGCGCGGAGCAUCCCGCG--NNNNNNNN--AGAGGUUAGAGGAGACCCCAUAUACCGCAA-ACAAAACAGCAUAUGACACCUGGGA--UAGACUAG	

