

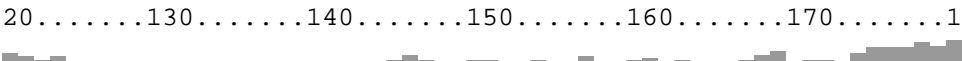
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MBFV-NC_008719.1 AUUUCUACUGAAAGG-GUAGAGAAA-----AGGAGUUUGCUUCUACUGGCCAACAAAG 52
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAACCCC 52
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAA-CCC 51
MBFV-NC_012533.1 AACAGUUUU-----CGGUUGAGAAAGAAAACAGAU-UCUUCGGAUAUGAAAA--CCC 50
MBFV-NC_001474.2 -CGUAGUUCUA-ACAGUUU-UUUAAUUAGAGAGCAGAUUCUCUG-----AUGAAUAA--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_002640.1 -----UCUA-ACAGUUUGUUUGAAUAGAGAGCAGAUUCUCUGGAAAAAUG-----AA--CC 47
MBFV-NC_009028.2 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCUCCAUUGUCU---AAA 49
MBFV-NC_040776.1 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUACCAUGUCU---AAA 49
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MBFV-NC_001563.2 A--U----UAACACAGUGCGAGCUGUUUCUUGGCAC-GAAGAUCUC-GAUGUCU---AAG 49
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MBFV-NC_008719.1 GGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUUGGUGA--CGGCCAAG---- 106
MBFV-NC_012532.1 AAAGAAGAAAU--CC--GGAGGA-UCCGGAUUGUCAAUUAGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1 AAAAAAGAAU--CC--GGAGGAUUCGGGAUUGUCAAUUAGCUAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1 AAAAAAGG-----GCCGCAUCGCGGCCUAUCAAUUAGCUGAAGCGCUUUGGAUCU 100
MBFV-NC_001474.2 AACGGAAAAAG--GC--G--AAAAACACGCCUUUCAAUUAGCUGAAACGCGAGAGAAAC 103
MBFV-NC_001477.1 AACGGAAAAAG--AC--G--GGUCGACCGUCUUUCAAUUAGCUGAAACGCGGAGAAAC 103
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MBFV-NC_009028.2 AAACCAGGAAAGUCC--GCUGCAAAGCGGACUGUCAAUUAGCUAAAACGCCUGGCUAGC 106
MBFV-NC_040776.1 AAACCAGGAGGACCC--GCCGGAAGGCGGGUUGUCAAUUAGCUGAAACGCCAGGCCAGC 106
MBFV-NC_034151.1 AAACCAGGAGGACCC--GGAAAGAACC GGCGUCGUUAAUAGCUGAAGCGCACAUCCCGC 106
MBFV-NC_001563.2 AAACCAGGAGGGCCC--GGUAAAAACCGGGCUGUCAAUUAGCUAAAACGCGGUAUGCCC 106
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MBFV-NC_001437.1 AAACCAGGAGGGCCC--GGUAAAAACCGGGCUUCAAUUAGCUGAAACGCGGCCUACCC 106
MBFV-NC_007580.2 AAACCAGGAAAAACCC--GGUAGAAACCGGGUUGUCAAUUAGCUAAAACGCGGGCUAUCC 106
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MBFV-NC_015843.2 AAACCAGGAAGACCC--GGCUCAGGCCGGGUUGUCAAUUAGCUAAAGCGCGGAACGUCC 109
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MBFV-NC_032088.1 AAACCAGGAAAGUCC--GGCAAUCGUCGGACUGUCAAUUAGCUAAAGCGCGGAGCAUCC 106
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MBFV-NC_035889.1 CGUG-----NNNNNNNN-----GUUAGAGGAGACCCCCCGGAAAACGCAAAACAGC 151
MBFV-NC_012533.1 AGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAACAUAAAACAGC 152
MBFV-NC_001474.2 CGCGUGU-----NNNNNNNN-----CUAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAACAGC 156
MBFV-NC_001477.1 CGCGUGU-----NNNNNNNN-----UAGAGGUUAGAGGAGACCCCCCGCACAAACAACAGC 156
MBFV-NC_001475.2 CGUGUGU-----NNNNNNNN-----ACUAGAGGUUAGAGGAGACCCCCCG-CAAAUAAAAACAGC 156
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MBFV-NC_012534.1 CCGGCACAGAGCGCCGU--AAG-UGUGCGUGGUGUG--GUAGAACACA-GUGUCU 258
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MBFV-NC_009029.2 UCGGCACAGAACCCGU--UGA-GAGCCGAGGCCCAA---AACAC-CAGAUUCU 258
MBFV-NC_032088.1 UAGGCACAGAGCGCCG--UAA-UGAUCUGGGUGGCCAA---AACAC-CAGAUUCU 258
MBFV-NC_009026.2 CCGGCACAGAACCCGA--CA-UGUGGGGGGUGCCUAGAACACCA--GAAUCU 258
MBFV-NC_018705.3 CACACAAAAACAGC---AUUUUAGACAUUGGGAAGACUAGGAGACCC-UCUGU 258

