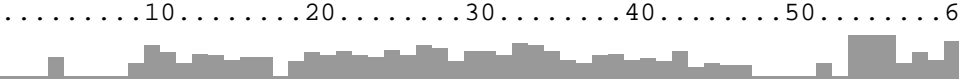
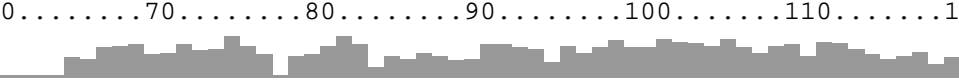


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MBFV-NC_008719.1 -----AUUUCUACUGAAAGG-GUAGAGAAAAGGAGUUUUGCUU--CUCAUGGCCAA 47
MBFV-NC_002031.1 -----UAAUCGUUCGUUGAGCG-AUUAGACGAGAAACUGACCG--AACAUGUC-- 45
MBFV-NC_012532.1 AAC-----AGGUUU-----AAUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA 47
MBFV-NC_035889.1 ACA-----GGUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAA 47
MBFV-NC_012533.1 AAC-----AGUUUU-----CGUGUUGAGAAGAAACAGAUUUUCUGG--AAUAGAAA 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_012534.1 CCGU-----AAACACAGUUUGAAC-GUUUUUUGGAGA-GA---GAUUA---UUAUGUUUA 47
MBFV-NC_015843.2 ACGU-----AAACACAGUUUGAAC-GUUUUUUGGAUA-GA---GACAA---CUAUGUCUA 47
MBFV-NC_034151.1 U-----AACACAGUGUGAAUAGUUUUUUGAACA-GAGGUGAAAA---CCAUGUCUA 47
MBFV-NC_009028.2 U--U---UAACACAGUCUGAAUAGUUUUUUAGCA--AGGGA-UUUC---CCAUGUCUA 47
MBFV-NC_040776.1 U--U---UAACACAGUCUGAAUAGUUUUUUAGCA--GAGGU-ACUA---CCAUGUCUA 47
MBFV-NC_001563.2 A--U---UAACACAGUGCAGCUGUUUCUUGGCAC-GAAGA-UCU---CGAUGUCUA 47
MBFV-NC_009942.1 A--U---UAACACAGUGCAGCUGUUUCUUAGCAC-GAAGA-UCU---CGAUGUCUA 47
MBFV-NC_000943.1 -----UAACGCGGUUGAACAGUUUUUUGGAGCUUUUGAUUU-C---AAUUGUCUA 47
MBFV-NC_006551.1 ---U---UAACACAGUGCCCGGCAGUUUUUUGGAGC-GUUGAUUU-U---CAAUGUCUA 47
MBFV-NC_001437.1 ---U---UAGUGCAGUUUAACAGUUUUUUGAACCGGAAGUAAC-C---C-AUGACUA 47
MBFV-NC_007580.2 ---U---AACUUGACUGCGAACAGUUUUUU--AGCAGGGAAUUAAC---CAAUGUCUA 47
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAAAGUUUUUGAACAG-----CAAUGACCA 47
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUAACAAGUUUUUGAUCAGUGGA-----UAUAGACUA 47
MBFV-NC_009026.2 -----AGAAAAGUUUUCGAGGAAGACGAGAAGCGAAUUCUCU---GAAUGGCUA 47
MBFV-NC_018705.3 CCGU-----AAACACAGUUUGAACGUUUUUUGGAGAGAGACUA-----CUAUGUCA 47
TBFV-NC_003690.1 GUGUUGAGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGA-----AUGGCC 47
TBFV-NC_005062.1 GUGUUGAGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGG-----AUGGCC 47
TBFV-NC_027709.1 GUGUUGAGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGG-----AUGGCCA 47
TBFV-NC_001672.1 GUGUUGAGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGG-----AUGGUA 47
TBFV-NC_001809.1 GUGUUGAGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGG-----AUGGCA 47
TBFV-NC_003687.1 -----AGAAAAA-----GACAGCUUAGGAAACAAGAGCUGGGAGUGGUUAUG--- 43



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MBFV-NC_012735.1 CAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAGCGGCUAGA- 106
MBFV-NC_008719.1 CAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCCGGGGCGUCAAUAUGGUGACGGCCAAG- 106
MBFV-NC_002031.1 -----UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGU-ACGACGAGGA 97
MBFV-NC_012532.1 -ACCCCAAAGAAGAAAU--CCGGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAGUA 103
MBFV-NC_035889.1 -A-CCCAAAAAGAAAU--CCGGAGGAUCCGGAUUGUCAAUAUGCUAAAACGCGGAGUA 103
MBFV-NC_012533.1 -----CCCAAAAAGG-----GCCCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGA 97
MBFV-NC_002640.1 -A--CCAACGAAAAAAG--GUG--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGAGA 97
MBFV-NC_001477.1 -A--CCAACGAAAAAAG--ACG--GGUCGACCGUCUUUCAAUAUGCUGAAACGCGCGAGA 100
MBFV-NC_001475.2 -A--CCAACGGAAGAAG--ACG--GGAAAACCGUCUAUCAAUAUGCUGAAACGCGUGAGA 100
MBFV-NC_001474.2 -A--CCAACGGAAAAAG--GCG--AAAAACACGCCUUUCAAUAUGCUGAAACGCGAGAGA 100
MBFV-NC_012534.1 -ACAAAAAACCCAGGAAACCCGGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUGCA 106
MBFV-NC_015843.2 -ACAAAAAACCCAGGAAGACCCGGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAACG 106
MBFV-NC_034151.1 ---AAAAACAGGAGGACCCGGAAAGAACCGGGUCGUAAAUAUGCUGAAAGCGCACAUCC 103
MBFV-NC_009028.2 ----AAAAACCAGGAAAGUCCGUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCU 103
MBFV-NC_040776.1 ----AAAAACCAGGAGGACCCGCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAUG 103
MBFV-NC_001563.2 ----AGAAACCAGGAGGGCCGGUAAAAACCGGCUGUCAAUAUGCUAAAACCGGUAUG 103
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MBFV-NC_006551.1 ----AGAAACCAGGAGGGCCGGGAAGAAACCGGGCAUCAAUAUGCUGAAACCGGGCAU 103
MBFV-NC_001437.1 ----AAAAACAGGAGGGCCGGUAAAAACCGGCUAUCAAUAUGCUGAAACCGGCCUA 103
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MBFV-NC_009029.2 ----AAAAACCAGGAAGGCGCGGUAAGAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUGCG 103
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MBFV-NC_009026.2 ----GAAAACCAGGAAGACCCGGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGGCG 103
MBFV-NC_018705.3 -ACAAAAAACCCAGGAAGACCCGGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGCGCA 106
TBFV-NC_003690.1 GGAAGGCCUUCUAAAA---GGAAAGGGGGGGGGUCCCCUCACGAGCGUCGAAAGUG 103
TBFV-NC_005062.1 GGAAGGCCUUCUGAAA---GGAAAGGGGGGGGGUCCCCUCACGAGUGUCGAAAGAG 103
TBFV-NC_027709.1 GGAAGUCCAUCCUGAAA---GGUAAGGGGGGGGGUCCCCUCACGAGUGUCGAAAGAG 103
TBFV-NC_001672.1 AGAAGGCCAUCCUGAAA---GGUAAGGGGGGGGGUCCCCUCACGAGUGUCGAAAGAG 103
TBFV-NC_001809.1 GGAAGACCAUCCUGAAA---GGCAAGGGGGGGGGUCCCCUCACGAGUGUCGAAAGAG 103
TBFV-NC_003687.1 --AUGACCACUUCUAAA---GGAAAGGGGGGGGGUCCCCUAGCGCAAGCUUAAAGUG 97



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MBFV-NC_002031.1 GUUCGCUCCUUGUNNNNNNN-----GAGACCCUCCAGGGAACAAUAGUGGGAC 146
MBFV-NC_012532.1 GCCCGUG-----NNNNNNNN-----GGUAGAGGAGACCCCCCGGAAACGCAAAAC 149
MBFV-NC_035889.1 GCCCGUG-----NNNNNNNN-----GUUAGAGGAGACCCCCCGGAAACGCAAAAC 148
MBFV-NC_012533.1 UCUAGCGCUGGCCNNNNNN-----GAGACCCCGCGCCCAUAACCAACUAAAAAC 149
MBFV-NC_002640.1 AACCGCGUAUCAANNNNNNNNACUAG--AGGUUAGAGGAGACCCCCCAACA--CAAAAAC 153
MBFV-NC_001477.1 AACCGCGUGU--NNNNNNNN--UAG--AGGUUAGAGGAGACCCCCCGCACAAACAACAAC 153
MBFV-NC_001475.2 AACCGUGUGU--NNNNNNNNACUAG--AGGUUAGAGGAGACCCCCCG--CAAAUAAAAAC 153
MBFV-NC_001474.2 AACCGCGUGU--NNNNNNNN-CUAG--AGGUUAGAGGAGACCCCCCG--CGAAUAAAAAC 153
MBFV-NC_012534.1 U---CCC-----NNNNNNNN--GG--AGACCCCUUGCGAGUUAACACCACA-ACAAAAC 151
MBFV-NC_015843.2 U---CCC-----NNNNNNNNA-GG--AGACCCCUUGCGAGUGAACACCACA-A-GAAAC 151
MBFV-NC_034151.1 CGCGCUA-----NNNNNNNN-----AGACCCCUUGCGAGUGAACG-CAAA-UAAAAAC 148
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MBFV-NC_040776.1 AGCGUUU-----NNNNNNNNA-GG--AGACCCCUGGCACACAGCGCAAA-GACAAAC 152
MBFV-NC_001563.2 CCCCGU-----NNNNNNNNGAGG--AGACCCCGUCCA-AAAACCA-A-AAGAAC 151
MBFV-NC_009942.1 CCCCGG-----NNNNNNNN--GG--AGACCCCGUGCCACAAAACACCACA-ACAAAAC 151
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MBFV-NC_006551.1 CCCCGCG-----NNNNNNNN-----ACCCCGCA-UCAUAAGCAUCA-----AAAAAC 144
MBFV-NC_001437.1 CCCCGCG-----NNNNNNNN--U--AGAGGAGACCCCGAUUGCA-----UCA-AAC 145
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MBFV-NC_009029.2 UCCCGCG-----NNNNNNNNAGAGG--AGACCCUCCCAAGAAAGCGCAAAACACAAAAC 155
MBFV-NC_032088.1 UCCCGCG-----NNNNNNNAGAGGUUAGAGGAGACCCCAUAGCGCAAA-ACAAAAC 156
MBFV-NC_009026.2 GCCAACG-----NNNNNNNN-----CCCCUUGGCAAAACAGUUAACGCACCAAAAGAAAC 152
MBFV-NC_018705.3 UCCC-----NNNNNNNN-----CCCCCGGAAAUCAAAGCAAGGCCCAACCU 147
TBFV-NC_003690.1 GCCCCAA-----NNNNNNNNCUC-----UCUUCAGGAUAUUCCUCCUCCUAUACCAA 151
TBFV-NC_005062.1 ACCGCUA-----NNNNNNNNCUC-----UCUUCAGGAUAUUCCUCCUCCUAUACCAA 151
TBFV-NC_027709.1 ACCGCAA-----NNNNNNNNCUC-----UCUUCAGGAUAUUCCUCCUCCUAUACCAA 151
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TBFV-NC_001809.1 ACCGCGA-----NNNNNNNN-UC-----UCUUCAGGGUAUUCCUCCUCCUAUACCAA 151
TBFV-NC_003687.1 ACCGCAAAUAAGUNNNNNNNNCU-----UCCUCGGGAUAUUCCGCCUCCUAUACCAA 151



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MBFV-NC_002031.1 CAUAUU-GACGCCAGGGA---AAGACC-GGAGUGGUUUCCUGCUUUC-CUCCAGGAGU 199
MBFV-NC_012532.1 AGCAUA-UUGACG--UGGGA--AAGACCA-GAGACUCCAUGAGUUU-CCACCACGCUGGC 202
MBFV-NC_035889.1 AGCAUA-UUGACG-CUGGGA--AAGACCA-GAGACUCCAUGAGUUU-CCACCACGCUGGC 202
MBFV-NC_012533.1 AGCAUA-UUGACACCUGGGA--AAGACCG-GAGACUGUGUGGUU-UCCCCACACCAGC 204
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MBFV-NC_009029.2 AGCAUA-UUGACACCUGGGA--AAGACUAGGGGAUUGC-UGCUC--UGGCAUCCCGG 208
MBFV-NC_032088.1 AGCAUA-UUGACACCUGGGA--UAGACUAGGAGAUCCG-UGCUC--UGGACCCAGU 209
MBFV-NC_009026.2 AGCAUA-UUGACACCUGGGA--UAGACC-GGAGAAUUGCUGCCUC--GCAACACCUCC 205
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TBFV-NC_027709.1 UUCCCCUCUCAGAGGGG-----GGGCGUCUUGUUCCUCGAGCCACCAUC 201
TBFV-NC_001672.1 UUCCCCUCUCAGAGGGG-----GGGCGUCUUGUUCCUCGAGCCACCAUC 201
TBFV-NC_001809.1 UUCCCCUCUCAGAGGGG-----GGGCGUCUUGUUCCUCGAGCCACCAUC 201
TBFV-NC_003687.1 UUCCCCCAGAACUGGGG-----GGGCGUCUUGUUCCUCGAGCCACCACC 201



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MBFV-NC_012735.1 CCUAAAGCACAGAUUGCUU--GAAUGCAGGGCGGGGAAGUUGAGAACCAACUAGUGUCU 258
MBFV-NC_008719.1 CCUUAAGCACAGAUUGCUU--GAAUGCAGGGUGGGGAAGUUGGGAACCAACUAGUGUCU 258
MBFV-NC_002031.1 CUGUGAGCACAGUUUGCUAAGAAUAAGCAGACCUUUGGAUGACAACACAAAACCACU 258
MBFV-NC_012532.1 CGCCAGGCACAGAUCGCC--GAACUUCGGCGGCGGUGUGGGGAAAUCCAUGGUUUCU 258
MBFV-NC_035889.1 CGCCAGGCACAGAUCGCC--GAAUAGCGGCGGCGGUGUGGGGAAAUCCAUGGUUUCU 258
MBFV-NC_012533.1 CCCCUGGCACAGAGCGCC--AGAUCGGGGUGUGGUGGAGCAACCAACAGGACU 258
MBFV-NC_002640.1 AUCCAGGCACAGAGCGCC--GCAAGAUG--GAUUGGUUGUUGAUCCAACAGGUUCU 258
MBFV-NC_001477.1 UUCCAGGCACAGAGCGCC--AAAAAUG--GAAUGGUCGUUGAUCCAACAGGUUCU 258
MBFV-NC_001475.2 UUCCAGGCACAGAGCGCC--AGAAAUG--GAAUGGUCGUUGAUCCAACAGGUUCU 258
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MBFV-NC_034151.1 CACUUGGCACAGAGCGCC--GUAAGU-GUGGCUGGUGGUGGUA--GAACACAGAGUCU 258
MBFV-NC_009028.2 CACUAGGCACAGAUCGCC--GCAAGUAGUGGCUGGUGGUGA--AACACAUGGAUCU 258
MBFV-NC_040776.1 CACUAGGCACAGAUCGCC--GAAUUGUGGCUGGUGGUGGAGCA--AACACA-GGAUCU 258
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MBFV-NC_009942.1 CACACGGCACAGUCGCC--GACAAUGGUGGCUGGUGGUGCGA--GAACACAGGAUCU 258
MBFV-NC_000943.1 CACAAGGCACCGAGCGCC--GAACACUGAGCUGAUGGGGGAGAAGACCACAGGAUCU 258
MBFV-NC_006551.1 CACAAGGCACAGAGCGCC--GAAAUUGUGGCUGGUGGGAACUAGACCACAGGAUCU 258
MBFV-NC_001437.1 UACUAGGCACAGAGCGCC--GAAGUAGUAGCUGGUGGUGGAGGAACACAGGAUCU 258
MBFV-NC_007580.2 CACAAGGCACAGAGCGCC--GUAAGUAGUGGCUGGUGGUGA--AAAUCACAGGAUCU 258
MBFV-NC_009029.2 U-CUGGCACAGAGCGCC--GUUGAGGA-----GCCGGAGGCCAAAACACCAGAUCU 258
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