

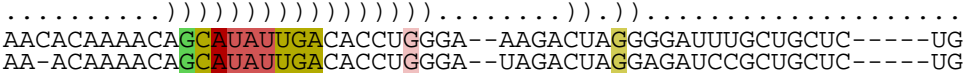
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MBFV-NC_040776.1 U--U-----UAACACAGUCUGAAU--AGUUUUUU--AGCAGAGGU--ACUAC---CAUGUCU-- 46
MBFV-NC_034151.1 -----UAACACAGUGUGAAU--AGUUUUUUUGAACAGAGGUGAAAC---CAUGUCU-- 46
MBFV-NC_001563.2 A--U-----UAACACAGUGCGAGC--UGUUUUUUGGCACGAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_009942.1 A--U-----UAACACAGUGCGAGC--UGUUUUUUGAAGCAGCAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_001437.1 U--U-----AGUCAGUUUAACA--GUUUUUUUGAACGGAAGA--UAAC---CAUGACU-- 46
MBFV-NC_007580.2 U--A-----ACUUGACUGCGAAC--GUUUUUUAGCAGGGAUU--ACCC---AAUGUCU-- 46
MBFV-NC_000943.1 -----UAACGCGGUUUGAAC--AGUUUUUUGGAGCUUUUGAUUUCA--AAUGUCU-- 46
MBFV-NC_006551.1 ----U-----UAACACAGUCCGGC--AGUUUCUUUGAGC--GUUGAUUUUC---AAUGUCU-- 46
MBFV-NC_012534.1 CCGU-----AAACACAGUUUGAAC--GUUUUUUUGGAGAGAG--AUUAU---UAUGUUU-- 46
MBFV-NC_015843.2 ACGU-----AAACACAGUUUGAAC--GUUUUUUUGGAUAGAG--ACAAC---UAUGUCU-- 46
MBFV-NC_009026.2 AGAA-----AAGUUUUC---GAGG--AAAGACGAG--AAGCGAAUUCUCUG---AAUGGCU-- 46
dISFV-NC_017086.1 AUAACACGUUU--GAAUAGGUUUUUAUUGUGAAUCGAAAUACCG-----AUGGCG-- 46
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAGUUUCUGG--U---CAUG----- 43
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCUGG--U---CAUG----- 43
MBFV-NC_012533.1 AACAGUUUU-----CGUGUUGGAAAGAAACAGAUUCUCUGG--A---AAUG----- 43
MBFV-NC_002640.1 -----UCUA--ACAGUUUGUUUGAAUAGAGAGCAG--AUCUCUGGAAA---AAUG----- 43
MBFV-NC_001474.2 --CGUAGUUCUA--ACAGUUU--UUUAAUUAGAGAGCAG--AUCUCUG-----AUG----- 43
ACGUAGUUCUA--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUG----- 43
MBFV-NC_001475.2 ACGUAGUGCUG--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUG----- 43
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AUUUCUACUGAAAGGGU---AG-----AGA--A--AAGGAGUUUGCUUCUCAUGGC-- 45
MBFV-NC_008719.1 -UUUCUACUG--AAAGGA---GU-----AGA--AGAAAGGAGAUUCAUCCCAAUGGC--- 45
MBFV-NC_002031.1 UAAUCGUUUC-----UUGAG-----CGAUUAGCAGAGAACUGACCAGAACAUG-- 43
dISFV-NC_040610.1 --GAACAAUUUUUAAAGAAGACAAACUAAGAAAUU--AAUUUC-----AUGGUU-- 46
dISFV-NC_024017.1 UCGCGCAUUUUUGAUAGAGUU---GACUAGAUAAACGAUUUAUC-----GAUGGUG-- 46
dISFV-NC_016997.1 AAACA---UAACGCAUUUGAAUAGGUUUUGAAAGAGAUUGAAUC-----AAUGAGAUC 48
MBFV-NC_018705.3 CCGUA-----AACACAGUUUGAAGGUUUUUGGAGAGAGACUACU-----AUGUCA-- 46
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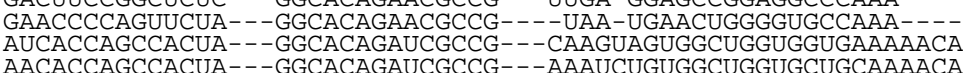
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MBFV-NC_009028.2 -A-----AAA-AACCAGGAAGUCGC--UGCAAAGCGGACUGCAAAUUGCUAAAGCGC 97
MBFV-NC_040776.1 -A-----AAA-AACCAGGAGGACCG--CGGAAGCGGGGUGUCAAUUGCGUAAAGCGC 97
MBFV-NC_034151.1 -A-----AAA-AACCAGGAGGACCGG--AAAGAACCGGGCUUAAUUAUGCUGAAGCGC 97
MBFV-NC_001563.2 -A-----AGA-AACCAGGAGGGGCCGG--UAAAACCGGGCUGCAAAUUGCUAAAGCGC 97
MBFV-NC_009942.1 -A-----AGA-AACCAGGAGGGGCCGG--CAAGAGCGGGGCUUCAAUUAUGCUAAAGCGC 97
MBFV-NC_001437.1 -A-----AAA-AACCAGGAGGGGCCGG--UAAAACCGGGCUGCAAAUUGCUAAGCGC 97
MBFV-NC_007580.2 -A-----AAA-AACCAGGAAGACCCGG--UAGAAACCGGGUUAUCAAUUGCUAAAGCGC 97
MBFV-NC_000943.1 -A-----AAA-AACCAGGAGGACCGG--GAAGCCCCGGGUGUCAAUUAUGCUAAAGCGC 97
MBFV-NC_006551.1 -A-----AGA-AACCAGGAGGGGCCGG--AAGAACCAGGGCCAUAUUAUGCUAAGCGC 97
MBFV-NC_012534.1 -AA--CAAAA-AACCAGGAAGACCGG--GGCAACCCGGGUUUAUCAAUUGCUAAAGCGC 100
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dISFV-NC_017086.1 -AA--UAAACCCAA---GAAGCCCGG--AAGA---CGGGCUUCAAUUAUGUAGAGACGU 94
MBFV-NC_012532.1 -AA--AAACCCCAAGAAGAAAUCCGG--AGGA-UCCGGAUUGUCAAUUAUGCUAAAGCGC 97
MBFV-NC_035889.1 -AA--AAA-CCCCAAAAAGAAAUCCGG--AGGAUUCGGAUUGUCAAUUAUGCUAAAGCGC 97
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MBFV-NC_002640.1 -----AA--CCAACGAAAAAAGGUGG--UUAGACCACCUU--UCAAUUAUGCUGAAACGC 91
MBFV-NC_001474.2 -AA--UAA--CCAACGAAAAAAGGCGA--AAAACACGCCUU--UCAAUUAUGCUGAAACGC 94
MBFV-NC_001477.1 -AA--CAA--CCAACGAAAAAAGAGCGG--GUCGACGUCUU--UCAAUUAUGCUGAAACGC 94
MBFV-NC_001475.2 -AA--CAA--CCAACGGAAGAAGACGG--GAAAACCGUCUA--UCAAUUAUGCUGAAACGC 94
dISFV-NC_027999.1 GAA--AACUAGAGUAGG---AUCUCGGGUUAAAACCCGAGGUGUCAAUUAUGAU-----UA 98
AAC--AAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUAUGG-----GA 98
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0.....70.....80.....90.....100.....110.....1



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MBFV-NC_032088.1 GGAGCAUCCCGCG-----NNNNNNNN--AGAGGUUAGAGGAGACCCCCACAU--ACCGCA 147
MBFV-NC_009028.2 CUGGCUAGCGUGU-----NNNNNNNN--GUGGAGA---CCCUCAGCAC--CAAGCGC 142
MBFV-NC_040776.1 CCAGCCGCGGUU-----NNNNNNNN--A--GGAGA---CCCUGGCACACAAAGCGC 142
MBFV-NC_034151.1 ACAUCCCGCGCUA-----NNNNNNNN--AGAGGAGA---CCCUGGCAGUGAA--CGC 138
MBFV-NC_001563.2 GGUUAGCCCCCGCG-----NNNNNNNN--GAGGAGA---CCCCGUGCCA--AAAACACC 142
MBFV-NC_009942.1 GGAAUAGCCCCCGCG-----NNNNNNNN--GGAGA---CCCCGUGCCACAAACACC 141
MBFV-NC_001437.1 GGCCUACCCCGCG-----NNNNNNNN--UAGAGGAGA---CCCGCAUUUG-----137
MBFV-NC_007580.2 GCGGUUACCCCGCG-----NNNNNNNN--UUAGAGGAGA---CCCUCUGCCGUUACG---143
MBFV-NC_000943.1 GGCAUACCCCGCG-----NNNNNNNN--GA---CCCCACUCUCAAAGAGCAUC 138
MBFV-NC_006551.1 GGCAUACCCCGCG-----NNNNNNNN--A---CCCCGCA--UCAUAAGCAUC 136
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MBFV-NC_015843.2 GGAACGCUCC-----NNNNNNNN--AGGAGA---CCCUGCGAGUGAACACC 142
MBFV-NC_009026.2 ACGGCGGCCAACG-----NNNNNNNN--C---CCCUU--GGCAAAACAGUUAACGCAC 142
dISFV-NC_017086.1 GCGUUGCCUCGCGUCU-----NNNNNNNN--AGAGGAGACCCCCCAAAAU---138
MBFV-NC_012532.1 GGAGUAGCCCGUG-----NNNNNNNN--GGUAGAGGAGACCCCCCGGAAAC---142
MBFV-NC_035889.1 GGAGUAGCCCGUG-----NNNNNNNN--GUUAGAGGAGACCCCCCGGAAAC---141
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MBFV-NC_001474.2 GAGAGAAACCGCGUGU---NNNNNNNN--CUAGAGGUUAGAGGAGACCCCC--CGAAC---146
MBFV-NC_001477.1 GCGAGAAACCGCGUGU---NNNNNNNN--UAGAGGUUAGAGGAGACCCCCCGCAAC---146
MBFV-NC_001475.2 GUGAGAAACCGUGUGU---NNNNNNNNACUAGAGGUUAGAGGAGACCCCCCG--CAAA---146
dISFV-NC_027999.1 GCGUACUGGA---A---NNNNNNNN-----UGGCGAACCCCAUUGGAAAU---138
MBFV-NC_008719.1 CGGCCAAGGCC---A---NNNNNNNN-----GAGGAGACCCUCCGAGCAC--U-138
MBFV-NC_012735.1 CGGCUAGAGUG---A---NNNNNNNN-----C--CAUUCUCCUCCGAGCACAC--138
MBFV-NC_002031.1 GGAGUUCGCUCCUUGU---NNNNNNNN-----GAG---ACCCUCCAGGGAACAA--136
dISFV-NC_040610.1 AUUACCAACCGCU---NNNNNNNN-----ACCCCUCAUUUAU---131
dISFV-NC_024017.1 AUCACCAAGCGUG---NNNNNNNN-----CCCCACU--GCUGAG---130
dISFV-NC_016997.1 AAGCGUC---NNNNNNNN-----AGGAAACCCCCCAAAAU---138
MBFV-NC_018705.3 GCGCAUCC-----NNNNNNNN-----CCCCCGGAUAUCA--131
20.....130.....140.....150.....160.....170.....1



.....))))))))).....)).....
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MBFV-NC_032088.1 AA--ACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGUCCCGCUGCUC----UG 199
MBFV-NC_009028.2 GAA--ACAAAACAGCAUAUUGACGCCUGGGA--AAGACCGGAGAUCCUCUGCUC--U-UCC 195
MBFV-NC_040776.1 AAAGACAACACAGCAUAUUGACACCUGGGA--AAGACCGGAGACCCUCUGCUC--C-UGC 196
MBFV-NC_034151.1 AAUAUAAAACAGCAUAUUGACACCUGGGA--UAGACUGGAGACCCUCUGCUCU--U-UUC 194
MBFV-NC_001563.2 A--AAAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGGAUUCUUC--UGCUC--UGCAC 197
MBFV-NC_009942.1 ACAACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGAUUCUUC--UGCUC--UGCAC 197
MBFV-NC_001437.1 --CAUCAAACAGCAUAUUGACACCUGGGA--UAGACUGGAGAUUCUUCUGCUCU--AUCUC 193
MBFV-NC_007580.2 --CAACAACAGCAUAUUGACACCUGGGA--AGACAG--GAGAUCCCCUGCUCU--UUUUC 194
MBFV-NC_000943.1 ---AAACAACAGCAUAUUGACACCUGGGA--AAGACUAGGAGAUUCUUCUGCUCU--AUUCC 193
MBFV-NC_006551.1 ---AAAAAACAGCAUAUUGACACCUGGGAUAGACUAGGAGAUUCUUCUGCUCU--AUUCC 193
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MBFV-NC_009026.2 CAAAGAAACAGCAUAUUGACACCUGGGA--UAGACC--GGAGAAUUGCGUCCU--CG--C 196
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MBFV-NC_012533.1 ---AUAACAGCAUAUUGACACCUGGGA--AAGACCGGAGACUCCUUGGCUU--UCCCC 196
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MBFV-NC_001474.2 ---AAAAACAGCAUAUUGACG--CUGGGA--AAGACCAAGAUCCUG--CUGU--CUCUG 196
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MBFV-NC_008719.1 --AUAGC--GGACCAUAUUGACGCGUGG--A--AAGACCGAGACACUCCUUGAUU--CUCAC 192
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dISFV-NC_040610.1 --UAUAUAACAGCAUAUUGACACCAAGG--A--UAGACCGAGAAA--U--ACACAUUUCUGG 185
dISFV-NC_024017.1 --AGCAAAUAGCAUAUUGACACUUGGGA--GAGACCAAGAUU--UGAUGUUGUUUGUCA 186
dISFV-NC_016997.1 --AUAGGCUUUAUAUUGACACCAUGGA--AAGACCGAGAUACCUUUGGUUUCUGGAG 193
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.....
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MBFV-NC_032088.1 GAACCCAGUUCUA---GGCACAGAACGCCG---UAA--UGAACUGGGGUGGCCAAA--- 247
MBFV-NC_009028.2 AUCACCAGCCACUA---GGCACAGAUCCGCC---CAAGUAGUGGCUGGUGGUGAAAAACA 249
MBFV-NC_040776.1 AACACAGCCACUA---GGCACAGAUCCGCC---AAAUUGUGGCUGGUGGUGCAAAACA 250
MBFV-NC_034151.1 AACACAGCCACUC---GGCACAGAACGCCG---UAAGU--GUGCGUGGUGGUGAA--AUUA 246
MBFV-NC_001563.2 A--ACCAGCCACAC---GGCACAGUGCGCCG---ACAUAGGUGGCUGGUG--GUGCUAGA 247
MBFV-NC_009942.1 A--ACCAGCCACAC---GGCACAGUGCGCCG---ACAUUGUGGCUGGUG--GUGCGAGA 247
MBFV-NC_001437.1 AACAUAGCUACUA---GGCACAGAGCGCCG---AAGUAGUAGCUGGUGGUGGAGGAAGA 247
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MBFV-NC_035889.1 ACGCUGGCCGCCA---GGCACAGAUCCGCC---AAUAGCGCGCGCCGGUGUGGGGAAAU 247
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MBFV-NC_001475.2 CAGCAUCAUCCA---GGCACAGAACGCCA---GAAAUGGAAUUGGUGCUGUUG--AAU 247
dISFV-NC_027999.1 AGCAUCCACCCGAAA---GGCACAGUAUGCCU---GAAACUUCGCGUGGUGUAGUAGUAAA 246
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MBFV-NC_012735.1 CUUUCUGGCCCUAA---AGCACAGAUUGCUU--G--AAUGCAGGCGGGGAAUUGGAGAAC 246
MBFV-NC_002031.1 C--AGAGGUCUGA---AGCACAGUUGCUAAGAUAAGUAGCUGUUCUUG--GAUGACAAA 246
dISFV-NC_040610.1 GCACUAAACACUAGA---AGCACAGGCGCUUCGAGAAGUAGUUUUUGACGAGAGAA 244
dISFV-NC_024017.1 AUGCAGCAACAGCCAUGGCACAGAGCGCCUAAGAGAGCGUUGUUUAUAAUUUACAAAA 246
dISFV-NC_016997.1 ACUAACUCUGAAU---GGCACAGAGCGCCAAAGAACCGAGUUGAUUGGAAAAAGAC 249
MBFV-NC_018705.3 ACCCCUCCGAG---UUAACACCAAGCAAAAAACAGCAUAUUGACACCUUGGGAAGACU 244
40.....250.....260.....270.....280.....290.....3



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MBFV-NC_009029.2 -ACACCAGAUUC--- 258
MBFV-NC_032088.1 -ACACCGAAUUC--- 258
MBFV-NC_009028.2 CAUGGA---UCU--- 258
MBFV-NC_040776.1 CA--GA---UCU--- 258
MBFV-NC_034151.1 AACACAGAGUCU--- 258
MBFV-NC_001563.2 -ACACAGGAUCU--- 258
MBFV-NC_009942.1 -ACACAGGAUCU--- 258
MBFV-NC_001437.1 -ACACAGGAUCU--- 258
MBFV-NC_007580.2 -UCACAGGAUCU--- 258
MBFV-NC_000943.1 C--CACAGGAUCU--- 258
MBFV-NC_006551.1 C--CACAGGAUCU--- 258
MBFV-NC_012534.1 -ACACAGUGUCU--- 258
MBFV-NC_015843.2 -ACACAGGAUCU--- 258
MBFV-NC_009026.2 C---CAGAUUCU--- 258
dISFV-NC_017086.1 CAACAAGUAUCU--- 258
MBFV-NC_012532.1 UAG--GUU--UCU--- 258
MBFV-NC_035889.1 CCAUGGGU--CUU--- 258
MBFV-NC_012533.1 CCAACAGG--ACU--- 258
MBFV-NC_002640.1 CAACAGGU--UCU--- 258
MBFV-NC_001474.2 CAACAGGU--UCU--- 258
MBFV-NC_001477.1 CAACAGGU--UCU--- 258
MBFV-NC_001475.2 CAACAGGU--UCU--- 258
dISFV-NC_027999.1 CAACCAGUUUCU--- 258
MBFV-NC_008719.1 CAACUAGUGUCU--- 258
MBFV-NC_012735.1 CAACUAGUGUCU--- 258
MBFV-NC_002031.1 CACAAAACUCU--- 258
dISFV-NC_040610.1 UAU--GUAAUACUUGG 258
dISFV-NC_024017.1 AAUAACAUUUCU--- 258
dISFV-NC_016997.1 UA---AGUAUCU--- 258
MBFV-NC_018705.3 AGGAGACCCUCU--GU 258
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