

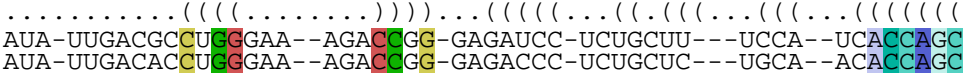
MBFV-NC_009028.2 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCCCAUGUCU---AAA 49
MBFV-NC_040776.1 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUACCAUGUCU---AAA 49
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUUGAAACCAUGUCU---AAA 49
MBFV-NC_001563.2 A--U----UAACACAGUGCGAGCGUUUUCUUGGCAC-GAAGAUUCU-GAUGUCU---AAG 49
MBFV-NC_009942.1 A--U----UAACACAGUGCGAGCGUUUUCUUGGCAC-GAAGAUUCU-GAUGUCU---AAG 49
MBFV-NC_000943.1 -----UAACGCGGUUUUGAACAGUUUUUUUGGAGCUUUUGAUUUUCAAUGUCU---AAA 49
MBFV-NC_006551.1 ---U----UAACACAGUGCCGCGCAGUUUUUUUGAGC-GUUGAUUUUUCAAUGUCU---AAG 49
MBFV-NC_001437.1 ---U----UAGUGCAGUUUAAACAGUUUUUUUAGAACGGAAGAUAAACC-AUGACU---AAA 49
MBFV-NC_007580.2 ---U----AACUUGACUGCGAACAGUUUUUUUAGCAGGGAU-UACCCAAUGUCU---AAA 49
MBFV-NC_012534.1 CCGU----AAACACAGUUUGAAC-GUUUUUUUGGAGAGAGAUUAUU---AUGUUUAACAAA 52
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAAC-GUUUUUUUGGAGAUAGAGACAACU---AUGUCUAACAAA 52
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAAAGUUUUUUUGAACAG----C----AAUGACC---AAA 49
MBFV-NC_032088.1 UC----UAUAACUCAGUGUGAACAAAGUUUUUUUGAUCAGUGGAU---AAUGACU---AAA 49
MBFV-NC_009026.2 AG-----AAAAGUUUUCGAGGAAAGACGAGAAGCGAAUUCUCUG-AAUGGCU---AGA 49
dISFV-NC_024017.1 UC-----GCGCAUUUGUAGAAAGAGUCACAAACUAGAAUUUUAUUGCAUUGGUG---ACU 49
dISFV-NC_017086.1 AU-----AACACGUUUGAAUAGGUUUUAUUGUGAAUCGAAAUACCGAUGGCG---AAU 49
MBFV-NC_002640.1 -----UCUA-ACAGUUUGUUUGAAUAGAGAGCAGAUUCUCUGGAAAAAUG---AA--CC 47
MBFV-NC_001474.2 -CGUAGUUCUA-ACAGUUU-UUUUUAUAGAGAGCAGAUUCUCUG-----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_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAACC 52
MBFV-NC_035889.1 ACAGGUUUU-----AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAACC 51
MBFV-NC_012533.1 AACAGGUUU-----CGUGUUGAGAAAGAAAAACAGAU-UCUUCGGAAAAUGAAAA---CCC 50
dISFV-NC_040610.1 -----GAACAAUUUUAAAGAAAGUCACAAACUAGAAUUUUAUUGCAUUGGUG---AGA 49
MBFV-NC_002031.1 U-----AAUCGUUCGUUGAGCG-AUUAGCAGAGAACUGACCAGAACAUGC-----45
dISFV-NC_016997.1 A-----AACAUACG-CAUUGAAUAGGUUUGAAAGAGUUGAAUCAAUGAGUCAUUG 52
MBFV-NC_008719.1 A-----UUUCUACUGAAAGG-GUAGAGAAAGAGGAGUUUUGCUUUCUACUGGCAACAAG 52
MBFV-NC_012735.1 -----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
dISFV-NC_027999.1 G-----GAUUAUUUGUUGUUUGAGAGAAACGAGAGAGUACUACU-AUGGUUGAGAAA 52
MBFV-NC_018705.3 CCGU----AAACACAGUUUGAACAGUUUUUUUGGAGAGAGACUACU---AUGUCAAACAAA 52
.....10.....20.....30.....40.....50.....6



MBFV-NC_009028.2 AAACCAGGAAAGUCC---GCUGCAAAGCGGGACUGUCAAUUAGCUAAAACGCCUGGCGUAGC 106
MBFV-NC_040776.1 AAACCAGGAGGACCC---GCCGGAAGCGGGGUGUCAAUUAGCUGAAACGCCAGCCGACGC 106
MBFV-NC_034151.1 AAACCAGGAGGACCC---GGAAAGAACCGGGUCGUUAAUAGCUGAAGCGCACAUCCCGC 106
MBFV-NC_001563.2 AAACCAGGAGGGGCC---GGUAAAAACCGGGCUGUCAAUUAGCUAAAACCGGUAUGCCCC 106
MBFV-NC_009942.1 AAACCAGGAGGGGCC---GGCAAGAGCGGGGUGUCAAUUAGCUAAAACCGGGAUUGCCC 106
MBFV-NC_000943.1 AAACCAGGAGGACCC---GGGAAGCCCCGGGUCGUCAAUUAGCUGAAAACCGGGCAUACCC 106
MBFV-NC_006551.1 AAACCAGGAGGGGCC---GGAGAAACCGGGCCAUCAAUUAGCUGAAACCGCGGCAUACCC 106
MBFV-NC_001437.1 AAACCAGGAGGGGCC---GGUAAAAACCGGGCUGUCAAUUAGCUGAAAACCGGGCUACCC 106
MBFV-NC_007580.2 AAACCAGGAAACCC---GGUAGAAACCGGGUUGUCAAUUAGCUGAAAACCGGGCGUATCC 106
MBFV-NC_012534.1 AAACCAGGAAACCC---GGGGCAACCCGGGUUGUCAAUUAGCUGAAAGCGCGGUGCAUCC 109
MBFV-NC_015843.2 AAACCAGGAAGACCC---GGCUCAGGCCGGGUUGUCAAUUAGCUGAAAGCGCGGAACGUCC 109
MBFV-NC_009029.2 AAACCAGGAAGGCC---GGUAGAAUUGCGGGCGUCAAUUAGCUGAAAGCGCGGUGCGUCC 106
MBFV-NC_032088.1 AAACCAGGAAAGUCC---GGCAAUCGUCGGACUGUCAAUUAGCUGAAAGCGCGGAGUACCC 106
MBFV-NC_009026.2 AAACCAGGAAGACCC---GGAGGAAACCGGGUCGUCAAUUAGCUGAAAGCGUACGGCGGCC 106
dISFV-NC_024017.1 AAUUCAGAAAGGCC---GCCGCAAGCGGGGUGUAAUAGCUUUAACAC---GCAUACCA 104
dISFV-NC_017086.1 AAACCAAGAAGC-----CCGGAAGACGGGCUAUCGAUUAUGUGAGACGUGCGUUGCCU 103
MBFV-NC_002640.1 AACGAAAAAG--GU---G--GUUAGACCACCUUUCAAUAGCUGAAACGCGAGAGAAAC 100
MBFV-NC_001474.2 AACGGAAGAAAG--GC---G--AAAAACACGCCUUUCAAUAGCUGAAACGCGAGAGAAAC 103
MBFV-NC_001477.1 AACGGAAGAAAG--AC---G--GGUCGACCGUCUUUCAAUUAGCUGAAACGCGGAGAAAC 103
MBFV-NC_001475.2 AACGGAAGAAAG--AC---G--GGAAACCGUCUAUCAAUUAGCUGAAACGCGGAGAGAAAC 103
MBFV-NC_012532.1 AAAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAAUUAGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1 AAAAAAGAAAU--CC---GGAGGAUUCGGAUUGUCAAUUAGCUGAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1 AAAAAAGG-----GCCGCAUCGCGGCCUAUCAAUUAGCUGAAGCGCUUUGGAUCU 100
dISFV-NC_040610.1 CAACUUGGAAAGACC---GUUGCAAGACGGGUCUGUCAAUUAGCUGAAGCGGAUUAACCAC 106
MBFV-NC_002031.1 --UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUUAGGUUACGACGAGGAGUUCGC 103
dISFV-NC_016997.1 GUCUUUCGAGCUGAGGAGGCCCGUUAACCGGGCCGUCGAUUAUAAUAAACGUAAGC-----107
MBFV-NC_008719.1 GGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUAGGUGACGGGCAAGG-----107
MBFV-NC_012735.1 GGAUGAAUUAAGUCUCGGGUCGUAUCCGAGGCGUCAAUUAGGUUAGCGGCUAGAG-----107
dISFV-NC_027999.1 ACUAGAGUAGGAUCUCGGGUUAAACCCGAGGUGUCAAUUAGUUAAGGCGUACUG-----107
MBFV-NC_018705.3 AAACCAGGAAGACC---GGGGCAACCCGGGUUGUCAAUUAGCUAAAGCGCGGCGCAUCC 109
0.....70.....80.....90.....100.....110.....1



MBFV-NC_009028.2 GUGU-----NNNNNNNN---GUGGAGACCCUCAGCAC-CAAGCGCGAA--ACAAACAGC 154
MBFV-NC_040776.1 GUUU-----NNNNNNNN---A-GGAGACCCCUUGGCACACACAGCGCAAA-GACAAACAGC 155
MBFV-NC_034151.1 GCUA-----NNNNNNNN-----AGACCCCUUGGCAGUGAA-CGCAAA-UAAAAACAGC 151
MBFV-NC_001563.2 CGCG-----NNNNNNNN-----GAGGAGACCCCGUGCCA-AAAAACCCA-A-AAGAAACAGC 154
MBFV-NC_009942.1 CGCG-----NNNNNNNN-----GGAGACCCCGUGCCACAAACACCACA-ACAAAACAGC 154
MBFV-NC_000943.1 CGCG-----NNNNNNNN-----GACCCACUCUCUAAAAGCAUC-AA-AC--AACAGC 148
MBFV-NC_006551.1 CGCG-----NNNNNNNN-----ACCCCGCA-UCAUAAAGCUAACA-AA--AACAGC 147
MBFV-NC_001437.1 CGCG-----NNNNNNNN-----UAGAGGAGACCCCGCAUUAUGCAUC-A--AACAGC 148
MBFV-NC_007580.2 CGCG-----NNNNNNNNUUAGAGGAGACCCCUUGCCGUUAAAC---GC-AAACAACAGC 154
MBFV-NC_012534.1 C-----NNNNNNNN-----GGAGACCCCUUGCGAGUUAAACACCAAA-ACAAAACAGC 154
MBFV-NC_015843.2 C-----NNNNNNNN-----AGGAGACCCCUUGCGAGUGAACACCACA-A-GAAACAGC 154
MBFV-NC_009029.2 CGCG-----NNNNNNNNAGAGG---AGACCCUCCACAAAGAACGCGAAACGCGAGAGAAAC 158
MBFV-NC_032088.1 CGCG-----NNNNNNNNAGAGGUUAGAGGAGACCCCCACAUAACCGCAAA-ACAAAACAGC 159
MBFV-NC_009026.2 AACG-----NNNNNNNNCCCCUUG---GCAAAACAGUUAACGCACCAAAAAGAAACAGC 155
dISFV-NC_024017.1 AGCGU-G-----NNNNNNNN-----CCCCACUGCU-----GAGAGCAAAAUAGC 141
dISFV-NC_017086.1 CGCGUCU---NNNNNNNNAGAGGAGACCCCCCCCA-----AAUUGAAGGGGCACU 148
MBFV-NC_002640.1 CGCGUAUCAANNNNNNNN---ACUAGAGGUUAGAGGAGACCCCCCAACA--CAAAAACAGC 156
MBFV-NC_001474.2 CGCGUGU---NNNNNNNN---CUAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAACAGC 156
MBFV-NC_001477.1 CGCGUGU---NNNNNNNN---UAGAGGUUAGAGGAGACCCCCCGCACAAACAACAAACAGC 156
MBFV-NC_001475.2 CGUGUGU---NNNNNNNN---ACUAGAGGUUAGAGGAGACCCCCCG--CAAAUAAAAACAGC 156
MBFV-NC_012532.1 CGUG-----NNNNNNNN-----GGUAGAGGAGACCCCGGAAACGCAAAACAGC 152
MBFV-NC_035889.1 CGUG-----NNNNNNNN-----GUUAGAGGAGACCCCCCGGAAACGCAAAACAGC 151
MBFV-NC_012533.1 AGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAACAUAACAGC 152
dISFV-NC_040610.1 CGCU-----NNNNNNNN-----NNNNNNNA-----CCCCUCAUUUUUAUUAACAGC 142
MBFV-NC_002031.1 UCCUUGU---NNNNNNNN-----GAG---ACCCUCCAGGGAACAAAUAGUGGGACC 147
dISFV-NC_016997.1 ----UGC-----NNNNNNNN-----AGGAAACCCCCCAAAAUGUAUAG-GCUAUU 148
MBFV-NC_008719.1 ----CCA-----NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UAAUAG-CGGACC 148
MBFV-NC_012735.1 ----UGA-----NNNNNNNN-----C-CCAUCCUCCCGAGCACACAUAG-CGGACC 148
dISFV-NC_027999.1 ----GAA-----NNNNNNNN-----UGGCGAACCCCAUUGGAAAUUUA---GCUUC 145
MBFV-NC_018705.3 C-----NNNNNNNN-----CCCCGCGAAAUCAAGACAAAGGCCCAACCCUGGC 150
20.....130.....140.....150.....160.....170.....1



MBFV-NC_009028.2 AUA-UUGACGCUGGGAA--AGACGG--GAGAUC-UCUGCUU---UCCA--UACCCAGC 204
MBFV-NC_040776.1 AUA-UUGACACUUGGAA--AGACGG--GAGACCC-UCUGCUU---UGCA--ACACCAGC 205
MBFV-NC_034151.1 AUA-UUGACACUUGGAU--AGACGG--GAGACCC-UCUGCUU--UUUCA--ACACCAGC 203
MBFV-NC_001563.2 AUA-UUGACACUUGGA--UAGACUAG--GGGAUCU-UCUGCUU---GCA--CAACCAGC 204
MBFV-NC_009942.1 AUA-UUGACACUUGGA--UAGACUAG--GAGAUCU-UCUGCUU---GCA--CAACCAGC 204
MBFV-NC_000943.1 AUA-UUGACACUUGGAA--AAGACUAG--GAGAUCU-UCUGCUUUAUCCA--ACAUCAGU 202
MBFV-NC_006551.1 AUA-UUGACACUUGGAAUUAUAGACUAG--GAGAUCU-UCUGCUUUAUCCA--ACAUCAC 202
MBFV-NC_001437.1 AUA-UUGACACUUGGAAU--AGACUAG--GAGAUCU-UCUGCUUUAUCCA--ACAUCAGC 202
MBFV-NC_007580.2 AUA-UUGACACUUGGAAAGA---GAG--GAGAUC-CCUGCUUU---UUCA--ACACCAGC 203
MBFV-NC_012534.1 AUA-UUGACACUUGGA--GAGACUAG--GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_015843.2 AUA-UUGACACUUGGA--UAGACUAG--GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_009029.2 AUA-UUGACACUUGGAA--AGACUAG--GGGAUUU-GCUGCUC---UGG--ACUCCGG 207
MBFV-NC_032088.1 AUA-UUGACACUUGGAU--AGACUAG--GAGAUC-GCUGCUC---UGG--AAACCCAG 208
MBFV-NC_009026.2 AUA-UUGACACUUGGA--UAGACUAG--AGAAUU---UGCUGCUCGA--ACACUCC 205
dISFV-NC_024017.1 AUA-UUGACACUUGGAA--GAGACUAG--AGAUAUGAUGUUGUUUGUCAUUGCAACA 198
dISFV-NC_017086.1 AUA-UCGACACUUGGAA--AGACUAG--AGUAUCUGCUGAUUACCCGCC---ACCAGAC 201
MBFV-NC_002640.1 AUA-UUGACG-CUGGAA--AGACUAGAGAG-UCCUGCUGUC---UGCA--ACAUCAA 205
MBFV-NC_001474.2 AUA-UUGACG-CUGGAA--AGACUAGAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_001477.1 AUA-UUGACG-CUGGAG--AGACUAGAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_001475.2 AUA-UUGACG-CUGGAG--AGACUAGAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_012532.1 AUA-UUGACG-UUGGAA--AGACUAGAGACU-CCAUGAGUUUCCACCAC---GCUUGCC 203
MBFV-NC_035889.1 AUA-UUGACG-CUGGAA--AGACUAGAGACU-CCAUGAGUUUCCACCAC---GCUUGCC 203
MBFV-NC_012533.1 AUA-UUGACACUUGGAA--AGACUAGAGACUCUGUUGGUUUUCC---CCAC---ACCAGCC 205
dISFV-NC_040610.1 AUA-UUGACACUUGGA--UAGACUAG--AGAAAUACACAAUUUUCGGGCACUAAACAC 197
MBFV-NC_002031.1 AUA-UUGACGCCAGG-A-AAGACUAGAGUGGUUUCUGCUUUUCCUC-C---AGAGGUC 200
dISFV-NC_016997.1 AUA-UCGACACUAGG-A-AAGACUAGAGAUACCUUUGGUUUCUCGAC-A---GUAUCC 201
MBFV-NC_008719.1 AUA-UUGACGCCUGG-A-AAGACUAGAGACACUUCUUGGUUCCACC-U---UUUCAC 201
MBFV-NC_012735.1 AUA-UUGACGCCAGG-A-AAGACUAGAGACACUCCUUGAUUUCUACC-U---UUUCAC 201
dISFV-NC_027999.1 AUA-UUGACACUUGGAA--AAGACUAGAGAAUACUGGUUACUUGCAG---CAACCAG 200
MBFV-NC_018705.3 AAAGCUGAACAACAGGGGAAGGACUAGAGGUUAUUGGACUCCUUGCGAGUGAACAC--208
80.....190.....200.....210.....220.....230.....2



MBFV-NC_009028.2 (((((.....))).....)))))).....)))).....)))).....)) 257
MBFV-NC_040776.1 CACUAGGCACAGAUCCGC--AAGUAGUGGCGUGG--GUGAA---AAACACAUGGAUC 257
MBFV-NC_034151.1 CACUAGGCACAGAUCCGC--AAUCUGUGGCGUGG--CUGCA---AAACACA-GGAUC 257
MBFV-NC_001563.2 CACUCGGCACAGAACGCCGU--AAGU-GUGGCGUGG--GUGA---UAGAACACA-GAGUC 257
MBFV-NC_009942.1 CACACGGCAGAGUGCCGA--CAUAGGUGGCGUGG--GUGC---UAGAACACA-GGAUC 257
MBFV-NC_000943.1 CACACGGCAGAGUGCCGA--CAUAGGUGGCGUGG--GUGC---GAGAACACA-GGAUC 257
MBFV-NC_000943.1 CACAAGGCACCGAGCGCCGA--ACACUGUACUGAUG--GGGAGAAGACCACA-GGAUC 257
MBFV-NC_006551.1 CACAAGGCACAGAGCGCCGA--AAUUGUGGCGUGG--GGGAACUAGACCACA-GGAUC 257
MBFV-NC_001437.1 UACUAGGCACAGAGCGCCGA--AGUAGUGGCGUGG--GUGAAGGAAGAACACA-GGAUC 257
MBFV-NC_007580.2 CACAAGGCACAGAGCGCCGU--AAAGUGUGGCGUGG--GUGAA-AAAAUCACA-GGAUC 257
MBFV-NC_012534.1 CAUCCGGCAGAGAGCGCCGU--AAG-UGUGGCGUGG--GUG--GUAGAACACA-GUGUC 257
MBFV-NC_015843.2 CACUUGGCACAGAUCCGCCGA--AAG-UGUGGCGUGG--GUG--GUAGAACACA-GGAUC 257
MBFV-NC_009029.2 CUCUCGGCACAGAACGCCGU--UGA-GGAGCCGGAGG--CCCAA---AACAC-CAGAU 257
MBFV-NC_032088.1 UUCUAGGCACAGAACGCCG--UAA-UGAACUGGGG--GCCAA---AACAC-CGAUC 257
MBFV-NC_009026.2 CACCCGGCACAGAACGCCGA--CA-UGGUGGGAGGG--UCCUAAGACACAC---GAUUC 257
dISFV-NC_024017.1 GCCAUUGGCACAGAGCGCCUG--AAGAGAGCUGUUGUUAUAAUUUACAAAAAUAACAU 256
dISFV-NC_017086.1 UACACGGCACAGCGCGCCGG--AAAAGGUAGUUUG---AGUUUGUAAACAAACAGUAU 256
MBFV-NC_002640.1 UCCA-GGCACAGAGCGCCGC--AAGAUGAUUGGUGU---UGU---UGAUCAACAGGUU 256
MBFV-NC_001474.2 UCCA-GGCACAGAACGCCAG--AAAUGGAUUGGUG---UGU---UGAAUCAAACAGGU 256
MBFV-NC_001477.1 UCCA-GGCACAGAACGCCAA--AAAUGGAUUGGUG---UGU---UGAAUCAAACAGGU 256
MBFV-NC_001475.2 UCCA-GGCACAGAACGCCAG--AAAUGGAUUGGUG---UGU---UGAAUCAAACAGGU 256
MBFV-NC_012532.1 GCCA-GGCACAGAACGCCGA--ACUUCGGCGGCCGGU---GUGG--GGAAUCCAUUGGU 256
MBFV-NC_035889.1 GCCA-GGCACAGAUCCGCCGA--AUAGCGCGCGCCGGU---GUGG--GGAAUCCAUUGGU 256
MBFV-NC_012533.1 CCCU-GGCACAGAGCGCC---AGAUCGGGUGGUGU---GUGG--AGCAACCCAACAGGA 256
dISFV-NC_040610.1 UAC-AAGCACAGAGCGCUUC--GAGAAGUAGUUGUUCUUGACGAGAGAAUUAUGUAU 254
MBFV-NC_002031.1 UGU-GAGCAGAGUUUGCUAAGAAUAGCAGACCUUUGGAU---ACAAACCAAAAACC 255
dISFV-NC_016997.1 CUAAUGGCACAGAGCGCCAU--AAGAAACGGAGUUAGUAUG---AAAAACAUAAGUA 255
MBFV-NC_008719.1 CCUUAAGGCACAGAUUGGUG--AAGUAGGUGGUGGUAUGU---GGAACCAACUAGU 255
MBFV-NC_012735.1 CCUAAAGGCACAGAUUGGUG--AAGUAGGUGGUGGUAUGU---AGAACCAACUAGU 255
dISFV-NC_027999.1 CGAAAGGCACAGAUUGGUGA--AACAUUCGGGUUAGUAUG---GUAACCAACAGU 255
MBFV-NC_018705.3 -----CACACAAAAACGAUUAUGACACCUGGAAAGACUAGGAGACCC---U 254
40.....250.....260.....270.....280.....290.....3



MBFV-NC_009028.2 258
MBFV-NC_040776.1 -U-- 258
MBFV-NC_034151.1 -U-- 258
MBFV-NC_001563.2 -U-- 258
MBFV-NC_009942.1 -U-- 258
MBFV-NC_000943.1 -U-- 258
MBFV-NC_006551.1 -U-- 258
MBFV-NC_001437.1 -U-- 258
MBFV-NC_007580.2 -U-- 258
MBFV-NC_012534.1 -U-- 258
MBFV-NC_015843.2 -U-- 258
MBFV-NC_009029.2 -U-- 258
MBFV-NC_032088.1 -U-- 258
MBFV-NC_009026.2 -U-- 258
dISFV-NC_024017.1 CU-- 258
dISFV-NC_017086.1 CU-- 258
MBFV-NC_002640.1 CU-- 258
MBFV-NC_001474.2 CU-- 258
MBFV-NC_001477.1 CU-- 258
MBFV-NC_001475.2 CU-- 258
MBFV-NC_012532.1 CU-- 258
MBFV-NC_035889.1 UU-- 258
MBFV-NC_012533.1 CU-- 258
dISFV-NC_040610.1 UUGG 258
MBFV-NC_002031.1 -ACU 258
dISFV-NC_016997.1 -UCU 258
MBFV-NC_008719.1 -UCU 258
MBFV-NC_012735.1 -UCU 258
dISFV-NC_027999.1 -UCU 258
MBFV-NC_018705.3 CUGU 258
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