

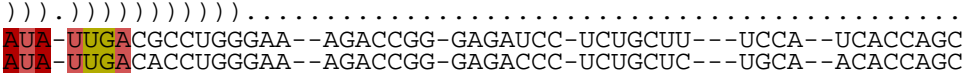
MBFV-NC_009028.2 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCCCAUGUCU---AAA 49
MBFV-NC_040776.1 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUACCAUGUCU---AAA 49
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUUGAAACCAUGUCU---AAA 49
MBFV-NC_001563.2 A--U----UAACACAGUGCGAGCUGUUUCUUGGCAC-GAAGAUCUC-GAUGUCU---AAG 49
MBFV-NC_009942.1 A--U----UAACACAGUGCGAGCUGUUUCUUAAGCAC-GAAGAUCUC-GAUGUCU---AAG 49
MBFV-NC_000943.1 -----UAACGCGGUUUAGAACAGUUUUUUUGAGCUUUUGAUUUUCAAUGUCU---AAA 49
MBFV-NC_006551.1 ---U----UAACACAGUGCCCGCAGUUUCUUUGAGC-GUUGAUUUUUCAAUGUCU---AAG 49
MBFV-NC_001437.1 ---U----UAGUGCAGUUUAAACAGUUUUUUAGAACGGAAGAUAAAC-AUGACU---AAA 49
MBFV-NC_007580.2 ---U----AACUAGCUGCGAACACAGUUUUUUUAGCAGGGAAU-UACCCAAUGUCU---AAA 49
MBFV-NC_012534.1 CCGU----AAACACAGUUUGAAC-GUUUUUUUGGAGAGAGAUUAUU---AUGUUUAACAAA 52
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAAC-GUUUUUUUGGAUAGAGACAACU---AUGUCUAACAAA 52
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAGUUUUUUUUGAACAG---C---AAUGACC---AAA 49
MBFV-NC_032088.1 UC----UAUAACUCAGUGUGAACAGUUUUUUUGAUCAGUGGAU---AAUGACU---AAA 49
MBFV-NC_009026.2 AG-----AAAAGUUUUCGAGGAAAGACGAGAAGCGAAUUCUCUG-AAUGGCU---AGA 49
dISFV-NC_024017.1 UC-----GCGCAUUUUGAGUAGACUAGAUAAACGAAUUAUUCGAUGGUG---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-UUUCUGGUCUAUGAAAAA-CCC 52
MBFV-NC_035889.1 ACAGGUUUU-----AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUCUAUGAAAAA-CCC 51
MBFV-NC_012533.1 AACAGUUUU-----CGUGUUGAGAAAGAAAAACAGAU-UCUUCGGAAAAUGAAAA--CCC 50
dISFV-NC_040610.1 -----GAACAAUUUUAAAGAAAGUCUAAAACUAGAAUUUUAUUUCAAUGGUU---AGA 49
MBFV-NC_002031.1 U-----AAUCGUUCGUUGAGCG-AUUAGCAGAGAACUGACCAGAACAUGC-----45
dISFV-NC_016997.1 A-----AACAUACG-CAUUGAAUAGGUUUGAAAGAGUUGAAUCAAUGAGAUCAAUG 52
MBFV-NC_008719.1 A-----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
MBFV-NC_012735.1 -----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
dISFV-NC_027999.1 G-----GAUUAUUUGUUGUUGAGAGAAACGAGAGAGAGUACUACU-AUGGUUGAGAAA 52
MBFV-NC_018705.3 CCGU----AAACACAGUUUGACAGUUUUUUUGGAGAGAGACUACU---AUGUCAAACAAA 52
.....10.....20.....30.....40.....50.....6



.....(((((((..... 106
MBFV-NC_009028.2 AAACCAGGAAAGUCC--GCUGCAAAGCGGACUGUCAAUJAGGCUAAAACGCCUGGCGUAGC 106
MBFV-NC_040776.1 AAACCAGGAGGACCC--GCCGGAAGCGGGGUUGUCAAUJAGGCUAAAACGCCAGCCGACGC 106
MBFV-NC_034151.1 AAACCAGGAGGACCC--GGAAAGAACCGGGUCUGUAAUJAGGCUAAGCGCACAUCCCGC 106
MBFV-NC_001563.2 AAACCAGGAGGGGCC--GGUAAAAACCGGGCUGUCAAUJAGGCUAAAACCGGUAUGCCCC 106
MBFV-NC_009942.1 AAACCAGGAGGGGCC--GGCAAGAGCCGGGUCUGCAAUJAGGCUAAAACCGGGAUUGCCC 106
MBFV-NC_000943.1 AAACCAGGAGGACCC--GGGAAGCCCCGGGUCUGUCAAUJAGGCUAAAACCGGGCAUACCC 106
MBFV-NC_006551.1 AAACCAGGAGGGGCC--GGAGAAACCGGGGCCAUCAAUJAGGCUAAAACCGGGCAUACCC 106
MBFV-NC_001437.1 AAACCAGGAGGGGCC--GGUAAAAACCGGGCUAUCAAUJAGGCUAAAACCGGGCUUACCC 106
MBFV-NC_007580.2 AAACCAGGAAAACCC--GGUAGAAAACCGGGUUGUCAAUJAGGCUAAAACCGGGCGUAUCC 106
MBFV-NC_012534.1 AAACCAGGAAAACCC--GGGGCAACCCGGGUUGUCAAUJAGGCUAAAAGCGCGGUGCAUCC 109
MBFV-NC_015843.2 AAACCAGGAAGACCC--GGCUCAGGCCGGGUUGUCAAUJAGGCUAAAAGCGCGGAACGUCC 109
MBFV-NC_009029.2 AAACCAGGAAGGCC--GGUAGAAAACCGGGCCUGUCAAUJAGGCUAAAAGCGCGGUGCGUCC 106
MBFV-NC_032088.1 AAACCAGGAAAGUCC--GGCAUUCGUCGGACUGUCAAUJAGGCUAAAAGCGCGGAGUACCC 106
MBFV-NC_009026.2 AAACCAGGAAGACCC--GGAGGAAAACCGGGUCUGUCAAUJAGGCUAAAAGCGUACGGCGGCC 106
dISFV-NC_024017.1 AAUUCAGAAAGACCC--GCCGCAAGCGGGCUGUAAUJAGGCUAAAAC--GCAUACCA 104
dISFV-NC_017086.1 AAACCAAGAAGC----CCGGAAGACGGGCUAUCGAUJAGGCUAAGAGACGUGCGUUGCCU 103
MBFV-NC_002640.1 AACGAAAAAAG--GU--G--GUUAGACCACCUUCAAUJAGGCUAAAACGCGAGAGAAAC 100
MBFV-NC_001474.2 AACGAAAAAAG--GC--G--GAAACACGCCUUCAAUJAGGCUAAAACGCGAGAGAAAC 103
MBFV-NC_001477.1 AACGAAAAAAG--AC--G--GGUCGACCGUCUUCAAUJAGGCUAAAACGCGCGAGAAAC 103
MBFV-NC_001475.2 AACGGAAGAAG--AC--G--GGAAACCGGUCUAUCAAUJAGGCUAAAACGCGUGAGAAAC 103
MBFV-NC_012532.1 AAAGAAGAAAU--CC--GGAGGA--UCCGGAUUGCAAUJAGGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1 AAAAAAGAAAU--CC--GGAGGAUUCGGAUUGCAAUJAGGCUAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1 AAAAAAGG--G--G--GCCGAUCGCGGCCUAUCAAUJAGGCUAAGCGCUUUGGAUCU 100
dISFV-NC_040610.1 CAACUUGGAAAGACC--GUUGCAAGACGGUCUGUCAAUJAGGCUAAGCGGGAUUAACCAC 106
MBFV-NC_002031.1 --UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUJAGGCUAAGCAGGAGGUUCGC 103
dISFV-NC_016997.1 GUCUUUCGAGCUGAGGAGGCCCGUUAACCGGGCCGUCGAUJAGGCUAAAACGUAAGC---- 107
MBFV-NC_008719.1 GGAGCAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUJAGGCUAAGCGGGAUUAAGC---- 107
MBFV-NC_012735.1 GGAUGAAUAAGUCUCGGGUCGAUCCCGAGGCGUCAAUJAGGCUAAGCGGCUAGAG---- 107
dISFV-NC_027999.1 ACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUJAGGCUUAGGCGUACUG---- 107
MBFV-NC_018705.3 AAACCAGGAAGACC--GGGCAACCCCGGGUUGUCAAUJAGGCUAAGCGCGGCGCAUCC 109
0.....70.....80.....90.....100.....110.....1



.....)))) 154
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-AAAACACCA-A-AAGAAACAGC 154
MBFV-NC_009942.1 CGCG-----NNNNNNNN--GGAGACCCCGUGCCACAAAACACCACA-ACAAAACAGC 154
MBFV-NC_000943.1 CGCG-----NNNNNNNN-----GACCCACUCUCUAAAAGCAUC-AA-AC--AACAGC 148
MBFV-NC_006551.1 CGCG-----NNNNNNNN-----ACCCCGCA-UCAUAAAGCAUCAA-AA--AACAGC 147
MBFV-NC_001437.1 CGCG-----NNNNNNNN-----UAGAGGAGACCCCGCAUUUGCAUC-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--AGACCCUCCACAAAGAACGCAAAACGAGAGAAAC 158
MBFV-NC_032088.1 CGCG-----NNNNNNNNAGAGGUUAGAGGAGACCCCCACAUACCGCAAA-ACAAAACAGC 159
MBFV-NC_009026.2 AACG-----NNNNNNNNCCCCUUG--GCAAAACAGUUAACGCACCAAAGAAACAGC 155
dISFV-NC_024017.1 AGCGU-G--NNNNNNNN-----CCCCACUGCU-----GAGAGCAAAUAAGC 141
dISFV-NC_017086.1 CGCGUCU--NNNNNNNNAGAGGAGACCCCCCCAA-----AAUGAAGGGGCAU 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-----GUUAGAGGAGACCCCGGAAACGCAAAACAGC 151
MBFV-NC_012533.1 AGCGCUGGCCNNNNNNNN-----GAGACCCCCCGCGCCCAUAACCAACAUAACAGC 152
dISFV-NC_040610.1 CGCU-----NNNNNNNN-----CCCCCAUUAUUUAUAAACAGC 142
MBFV-NC_002031.1 UCCUUGU--NNNNNNNN-----GAG--ACCCUCCAGGGAACAAAUAGUGGGAC 147
dISFV-NC_016997.1 ----UGC--NNNNNNNN-----AGGAAACCCCCCAAAAUGUAUAG-GCUAU 148
MBFV-NC_008719.1 ----CCA--NNNNNNNN-----GAGGAGACCCUCCGAGCAC-UAAUG-CGGA 148
MBFV-NC_012735.1 ----UGA--NNNNNNNN-----C-CCAUCCUCCCGAGCACAUAG-CGGAC 148
dISFV-NC_027999.1 ----GAA--NNNNNNNN-----UGGCGAACCCCAUUGGAAAUUUA--GCUU 145
MBFV-NC_018705.3 C-----NNNNNNNN-----CCCCGCGAAAUCAAGCAAGGCCCAACCCAGC 150
20.....130.....140.....150.....160.....170.....1



)))))))))..... 204
MBFV-NC_009028.2 AUA-UUGACGCCUGGGAA--AGACCGG-GAGAUC-UCUGCUU---UCCA--UCACCAGC 204
MBFV-NC_040776.1 AUA-UUGACACCUGGGAA--AGACCGG-GAGACCC-UCUGCUC---UGCA--ACACCAGC 205
MBFV-NC_034151.1 AUA-UUGACACCUGGGAU--AGACUGG-GAGACCC-UCUGCUUA-UUUA--ACACCAGC 203
MBFV-NC_001563.2 AUA-UUGACACCUGGGGA-UAGACUAG-GGAUUCU-UCUGCUCU---GCA--CAACCAGC 204
MBFV-NC_009942.1 AUA-UUGACACCUGGGGA-UAGACUAG-GGAUUCU-UCUGCUCU---GCA--CAACCAGC 204
MBFV-NC_000943.1 AUA-UUGACACCUGGGGA-AAGACUAG-GAGAUCU-UCUGCUCUAUUCCA--ACAUCAGU 202
MBFV-NC_006551.1 AUA-UUGACACCUGGGGAUUAGACUAG-GAGAUCU-UCUGCUCUAUUCCA--ACAUAAC 202
MBFV-NC_001437.1 AUA-UUGACACCUGGGGAU-AGACUGG-GAGAUCU-UCUGCUCUAUCCA--ACAUCAGC 202
MBFV-NC_007580.2 AUA-UUGACACCUGGAAAGA---CAG-GAGAUC-CCUGCUUU--UUUA--ACACCAGC 203
MBFV-NC_012534.1 AUA-UUGACACCUGGGGA-GAGACUAG-GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_015843.2 AUA-UUGACACCUGGGGA-UAGACUAG-GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_009029.2 AUA-UUGACACCUGGGGA--AGACUAG-GGGAUUU-GCUGCUC---UGG--ACUUCGG 207
MBFV-NC_032088.1 AUA-UUGACACCUGGGGAU--AGACUAG-GAGAUC-GCUGCUC---UGG--AACCCAG 208
MBFV-NC_009026.2 AUA-UUGACACCUGGGGA-UAGACCGG--AGAAUU--UGCUGCUCGA--ACACCUC 205
dISFV-NC_024017.1 AUA-UUGACACUUGGGGA-GAGACCAG-AGAUAUGAUGUUGUUUGUCAUUGCAACA 198
dISFV-NC_017086.1 AUA-UUGACACUUGGGGA--AGACCAG-AGGUACUCGUGAUUACCGGCC--ACCAGC 201
MBFV-NC_002640.1 AUA-UUGACG-CUGGGAA--AGACCAGAG-UCCUGCUGUC---UGCA--ACAACAA 205
MBFV-NC_001474.2 AUA-UUGACG-CUGGGAA--AGACCAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_001477.1 AUA-UUGACG-CUGGGAG--AGACCAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_001475.2 AUA-UUGACG-CUGGGAG--AGACCAGAG-UCCUGCUGUC---CUCA--GCAUCAU 205
MBFV-NC_012532.1 AUA-UUGACG--UGGGAA--AGACCAGAGACU-CCAUGAUUUCCACCAC--GCUGGCC 203
MBFV-NC_035889.1 AUA-UUGACG-CUGGGAA--AGACCAGAGACU-CCAUGAUUUCCACCAC--GCUGGCC 203
MBFV-NC_012533.1 AUA-UUGACACCUGGGA--AGACCGGAGACUCUGUUGGUUUUCC--CCAC--ACCAGCC 205
dISFV-NC_040610.1 AUA-UUGACACCAGGGA-UAGACCGG--AGAAUAACACAAUUUCCGGGCACUAAACAA 197
MBFV-NC_002031.1 AUA-UUGACCGCCAGGG-A-AAGACCGGAGUGGUUCUCUGCUUUUCCUC-C--AGAGGUC 200
dISFV-NC_016997.1 AUA-UUGACACCUAGG-A-AAGACCGGAGAUACCUUUGGUUUCUCGAC-A--UACGUC 201
MBFV-NC_008719.1 AUA-UUGACCGCCGGG-A-AAGACCGGAGACACUCCUUGAUUCCACC-U---UUCUCAC 201
MBFV-NC_012735.1 AUA-UUGACCGCCAGGG-A-AAGACCGGAGACACUCCUUGAUUCCACC-U---UUCUCGC 201
dISFV-NC_027999.1 AUA-UUGACACCAGGGGA-AAGACCGGAGAAUACUUGGCUUACCCAG--CAACCAC 200
MBFV-NC_018705.3 AUA-GCUGUAAACAGGGGAAGACUAGAGGUUAGGACUCCUUGCGAGUGAGAACC--U 208
80.....190.....200.....210.....220.....230.....2



..... 257
MBFV-NC_009028.2 CACUAGGCACAGAUCCCGC--AAGUAGUGGCUGGUG--GUGAA---AAACACAUGGAUC 257
MBFV-NC_040776.1 CACUAGGCACAGAUCCCGA--AAUCUGUGGCUGGUG--CUGCA---AAACACA-GGAUC 257
MBFV-NC_034151.1 CACUCGGCACAGAACGCCGU--AAGU-GUGGCUGGUG--GUGAAAAUAAACACA-GAGUC 257
MBFV-NC_001563.2 CACACGGCACAGUGCCCGA--CAUAGGUGGCUGGUG--GUGC--UAGAACACA-GGAUC 257
MBFV-NC_009942.1 CACACGGCACAGUGCCCGA--CAUAGGUGGCUGGUG--GUGC--GAGAACACA-GGAUC 257
MBFV-NC_000943.1 CACAAGGCACCGAGCGCCGA--ACACUGUGACUGAUG--GGGGAGAAGACCACA-GGAUC 257
MBFV-NC_006551.1 CACAAGGCACAGAGCGCCGA--AAAUUGUGGCUGGUG--GGGAACUAGACCACA-GGAUC 257
MBFV-NC_001437.1 UACUAGGCACAGAGCGCCGA--AGUAGUAGCUGGUG--GUGAGGAAGAACACA-GGAUC 257
MBFV-NC_007580.2 CACAAGGCACAGAGCGCCGU--AAAGUGUGGCUGGUG--GUGAA-AAAAUCACA-GGAUC 257
MBFV-NC_012534.1 CAUCUCCGCACAGAGCGCCGU--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GGUC 257
MBFV-NC_015843.2 CACUUGGCACAGAUCCCGA--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GGAUC 257
MBFV-NC_009029.2 CUCUCGGCACAGAACGCCGU--UGA-GGAGCCGGAGG--CCCAA---AACAC-CAGAU 257
MBFV-NC_032088.1 UUCUCGGCACAGAACGCCG--UAA-UGUAGUGGGU--GCCAA---AACAC-CGAAU 257
MBFV-NC_009026.2 CACCCGGCACAGAACGCCGA--CA-UGGUGGGAGGGG--UCCUAAGACACCA--GAUUC 257
dISFV-NC_024017.1 GCCAUGGCACAGAGCGCCUG--AAGAGAGCUGUUGUUAUAAUUUACAAAAAUAACAU 256
dISFV-NC_017086.1 UACACGGCACAGCGCGCCGG--AAAAGGUAGUUUG--AGUUUUAUAAACAAAGUAU 256
MBFV-NC_002640.1 UCCA-GGCACAGAGCGCCG--AAGAUGAUUGGUGU--UGU---UGAUCAACAGGU 256
MBFV-NC_001474.2 UCCA-GGCACAGAGCGCCAG--AAAUUGGAAUGGUGC--UGU---UGAAUACAGGU 256
MBFV-NC_001477.1 UCCA-GGCACAGAACGCCAA--AAAUUGGAAUGGUGC--UGU---UGAAUACAGGU 256
MBFV-NC_001475.2 UCCA-GGCACAGAACGCCAG--AAAUUGGAAUGGUGC--UGU---UGAAUACAGGU 256
MBFV-NC_012532.1 GCCA-GGCACAGAUCCCGA--ACUUCGGCGGCGCGU--GUGG--GGAAUCCAUUGGU 256
MBFV-NC_035889.1 GCCA-GGCACAGAUCCCGA--AUAGCGGCGGCGCGU--GUGG--GGAAUCCAUUGGU 256
MBFV-NC_012533.1 CCCU-GGCACAGAGCGCC--GAGUAGGUGGUUGGU--GUGG--AGCAACCAACAGGA 256
dISFV-NC_040610.1 UAC-AAGCACAGAGCGCUU--AGAAGGUUGUUGUUCUUGACGAGAGAAUUGUAU 254
MBFV-NC_002031.1 UGU-GAGCACAGUUUGCUCAAGAAUAGACAGACCUUUGGAU---ACAAACACAAAACC 255
dISFV-NC_016997.1 CUAAUGGCACAGAGCGCCAU--AAGAAACGGAGUAGGAUGG---AAAAACAUAAAGUA 255
MBFV-NC_008719.1 CCUUAAGCACAGAUUGCUUG--AAGUACGGGUGGGAUGUUG---GGAACCAACAGUG 255
MBFV-NC_012735.1 CCUAAAGCACAGAUUGCUUG--AAGUACGGGCGGGAUGUUG---AGAACCAACUAGUG 255
dISFV-NC_027999.1 CGAAAGGCACAGAUUGCCUGA--AACAUUCGGUGGUUGAUGU---GUAACCAACAGUG 255
MBFV-NC_018705.3 -----CACACAAAACAGCAUUAUAGACUCCGGGAAGAGACUAGGAGACCC--U 254
40.....250.....260.....270.....280.....290.....3



..... 258
MBFV-NC_009028.2 -U-- 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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