

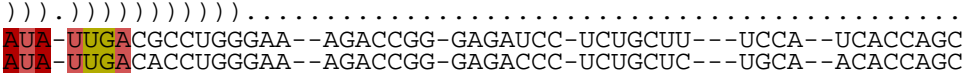
MBFV-NC_009028.2 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCCCAUGUCU---AAA 49
MBFV-NC_040776.1 U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUACCAUGUCU---AAA 49
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUUGAAACCAUGUCU---AAA 49
MBFV-NC_001563.2 A--U----UAACACAGUGCGAGCUGUUUUUUUGGCAC-GAAGAUUCU- GAUGUCU---AAG 49
MBFV-NC_009942.1 A--U----UAACACAGUGCGAGCUGUUUUUUUAGCAC-GAAGAUUCU- GAUGUCU---AAG 49
MBFV-NC_000943.1 -----UAACGCGGUUUUAACACAGUUUUUUUAGAGCUUUUGAUUUUCAAUGUCU---AAA 49
MBFV-NC_006551.1 ---U----UAACACAGUGCCCGCAGUUUUUUUUGAGC-GUUGAUUUUUCAAUGUCU---AAG 49
MBFV-NC_001437.1 ---U----UAGUGCAGUUUAAACAGUUUUUUUAGAACGGAAGAUAAAC-AUGACU---AAA 49
MBFV-NC_007580.2 ---U----AACUUGACUGCGAAACAGUUUUUUUAGCAGGGAU-UACCCAAUGUCU---AAA 49
MBFV-NC_012534.1 CCGU----AAACACAGUUUGAAC-GUUUUUUUGGAGAGAGAUUUU---AUGUUUAACAAA 52
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAAC-GUUUUUUUGGAUAGAGACAACU---AUGUCUAACAAA 52
MBFV-NC_009029.2 UGAUUUACAUAACACAGUGUGAAACAGUUUUUUUUGAACAG---C---AAUGACC---AAA 49
MBFV-NC_032088.1 UC----UAUAACUCAGUGUGAACAGUUUUUUUUGAUCAGUGGAU---AAUGACU---AAA 49
MBFV-NC_009026.2 AG-----AAAAGUUUUUCGAGGAAAGACGAGAAGCGAAUUCUCUG-AAUGGCU---AGA 49
dISFV-NC_024017.1 UC-----GCGCAUUUUUGAGUAAGCUAAGAUAAACGAAUUAUCGAUGGUG---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-UUUUAAUAGAGAGCAGAUUCUCUG-----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 AACAGGUUU-----CGUGUUGAGAAAGAAAAACAGAU-UCUUCGGAAAAUGAAAA--CCC 50
dISFV-NC_040610.1 -----GAACAAUUUUAAAGAAAGUCUAAAACUAGAAUUUUAUUUCAAUGGUU---AGA 49
MBFV-NC_002031.1 U-----AAUCGUUCGUUGAGCG-AUUAGCAGAGAACUGACCAGAACAUGU-----45
dISFV-NC_016997.1 A-----AACAUACG-CAUUGAAUAGGUUUUGAAAGAGUUGAAUCAAUGAGUCAUUG 52
MBFV-NC_008719.1 A-----UUUCUACUGAAAGGAGUAAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
MBFV-NC_012735.1 -----UUUCUACUGAAAGGAGUAAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
dISFV-NC_027999.1 G-----GAUUAUUUGUUGUUUGAGAGAAACGCGAGAGAGUACUACU-AUGGUUGAGAAA 52
MBFV-NC_018705.3 CCGU----AAACACAGUUUGAACAGUUUUUUUGGAGAGAGACUACU---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--GGAAAGAACCGGGGUCGUCAAUJAGGCUAAAACGCCAGCAUCCCGC 106
MBFV-NC_001563.2 AAACCAGGAGGGGCC--GGUAAAAACCGGGGCGUCAAUJAGGCUAAAACGCCGGAUUGCCC 106
MBFV-NC_009942.1 AAACCAGGAGGGGCC--GGCAAGAGCCGGGCGUCAAUJAGGCUAAAACGCCGGAUUGCCC 106
MBFV-NC_000943.1 AAACCAGGAGGACCC--GGGAAGCCCCGGGUCGUCAAUJAGGCUAAAACGCCGGCAUACCC 106
MBFV-NC_006551.1 AAACCAGGAGGGGCC--GGAGAAACCGGGGCCUCAAUJAGGCUAAAACGCCGGCAUACCC 106
MBFV-NC_001437.1 AAACCAGGAGGGGCC--GGUAAAAACCGGGGCUAUAJAGGCUAAAACGCCGGCCUACCC 106
MBFV-NC_007580.2 AAACCAGGAAAACCC--GGUAGAAACCGGGGUUGUCAAUJAGGCUAAAACGCCGGCGUAUCC 106
MBFV-NC_012534.1 AAACCAGGAAAACCC--GGGGCAACCCGGGUUGUCAAUJAGGCUAAAAGCGCGGUGCAUCC 109
MBFV-NC_015843.2 AAACCAGGAAGACCC--GGCUCAGGCCGGGUUGUCAAUJAGGCUAAAAGCGCGGAACGUCC 109
MBFV-NC_009029.2 AAACCAGGAAGGCC--GGUAGAAUUGCGGGCGUCAAUJAGGCUAAAAGCGCGGUGCGUCC 106
MBFV-NC_032088.1 AAACCAGGAAAGUCC--GGCAUUCGUCGGACUGUCAAUJAGGCUAAAAGCGCGGAGUACCC 106
MBFV-NC_009026.2 AAACCAGGAAGACCC--GGAGGAAACCGGGGUCGUCAAUJAGGCUAAAAGCGUACGGCGGCC 106
dISFV-NC_024017.1 AAUUCAGAAAGGCC--GCCGCAAGCGGGGCGUCAAUJAGGCUAAAAC--GCAUACCA 104
dISFV-NC_017086.1 AAACCAAGAAGC----CCGGAAGACGGGCUAUAJAGGCUAAAAGCGUGCGUUGCCU 103
MBFV-NC_002640.1 AACGAAAAAAG--GU--G--GUUAGACCACCUUCAAUJAGGCUAAAACGCGAGAGAAAC 100
MBFV-NC_001474.2 AACGAAAAAAG--GC--G--GAAACACGCCUCAAUJAGGCUAAAACGCGAGAGAAAC 103
MBFV-NC_001477.1 AACGAAAAAAG--AC--G--GGUCGACCGUCUCAAUJAGGCUAAAACGCGCGAGAAAC 103
MBFV-NC_001475.2 AACGGAAGAAG--AC--G--GGAAACCGGUCUCAAUJAGGCUAAAACGCGUGAGAAAC 103
MBFV-NC_012532.1 AAAGAAGAAAU--CC--GGAGGA--UCCGGAUUGUCAAUJAGGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1 AAAAAAGAAAU--CC--GGAGGAUUCGGAUUGUCAAUJAGGCUAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1 AAAAAAGG--G--G--GCCGAUCGCGGCCUCAAUJAGGCUAAAAGCGCUUUGGAUCU 100
dISFV-NC_040610.1 CAACUUGGAAAGACC--GUUGCAAGACGGUCUGUCAAUJAGGCUAAAAGCGGAUUAACCC 106
MBFV-NC_002031.1 --UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUJAGGCUAACGACGAGGAGUUCGC 103
dISFV-NC_016997.1 GUCUUUCGAGCUGAGGAGGCCCGGUUAAACGGGCGCGUCAAUJAGGCUAAAACGUAAGC----107
MBFV-NC_008719.1 GGAGCAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUJAGGCUAAAACGCGGAGUAGCC----107
MBFV-NC_012735.1 GGAUGAAUAAGUCUCGGGUCGUAUCCGAGGCGUCAAUJAGGCUAACGCGGCUAGAG----107
dISFV-NC_027999.1 ACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUJAGGCUUAGGCGUACUG----107
MBFV-NC_018705.3 AAACCAGGAAGACC--GGGCAACCCGGGUUGUCAAUJAGGCUAAAAGCGCGGCGCAUCC 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--GGAGACCCCGUGCCACAAACACCAACA-ACAAACAGC 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-----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--AGACCCUCCACAAAGAGCGCAAAACGCAAAACAGC 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-----GAGACCCCGCGCCCAUAACCAACAUAAAACAGC 152
dISFV-NC_040610.1 CGCU-----NNNNNNNN-----CCCCCAUUAUUUAUAAAACAGC 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-UAUAG-CGGAC 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-----CCCCGCGAAAUCAAGCAAGGCCCAACCCUGC 150
20.....130.....140.....150.....160.....170.....1



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MBFV-NC_009028.2 AUA-UUGACGCCUGGGAA--AGACCGG-GAGAUGC-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-GAGAUGC-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--ACUUCGGG 207
MBFV-NC_032088.1 AUA-UUGACACCUGGGGAU--AGACUAG-GAGAUCG-GCUGCUC---UGG--AACCCGAG 208
MBFV-NC_009026.2 AUA-UUGACACCUGGGGA-UAGACCGG--AGAAUU--UGCUGCUCGCA--ACACCUC 205
dISFV-NC_024017.1 AUA-UUGACACUUGGGGA-GAGACCAG-AGAUAUGAUGUUGUUUGUCAUUGCACAACA 198
dISFV-NC_017086.1 AUA-UUGACACUUGGGGA--AGACCAG-AGGUACUCGUGAUUACCCGCC--ACCAGAC 201
MBFV-NC_002640.1 AUA-UUGACG-CUGGGAA--AGACCAGAG-UCCUGCUGUCU---UGCA--ACAACAA 205
MBFV-NC_001474.2 AUA-UUGACG-CUGGGGA--AGACCAGAG-UCCUGCUGUCU---CUCA--GCAUCAU 205
MBFV-NC_001477.1 AUA-UUGACG-CUGGGAG--AGACCAGAG-UCCUGCUGUCU---CUCA--GCAUCAU 205
MBFV-NC_001475.2 AUA-UUGACG-CUGGGAG--AGACCAGAG-UCCUGCUGUCU---CUCA--GCAUCAU 205
MBFV-NC_012532.1 AUA-UUGACG--UGGGAA--AGACCAGAGACU-CCAUGAGUUUCCACCAC--GCUGGCC 203
MBFV-NC_035889.1 AUA-UUGACG-CUGGGGA--AGACCAGAGACU-CCAUGAGUUUCCACCAC--GCUGGCC 203
MBFV-NC_012533.1 AUA-UUGACACCUGGGA--AGACCGGAGACUCUGUUGGUUUUCC--CCAC--ACCAGCC 205
dISFV-NC_040610.1 AUA-UUGACACCAGGGA-UAGACCGG--AGAAUAACACAAUUUCCUGGGCACUAAACAA 197
MBFV-NC_002031.1 AUA-UUGACCGCCAGGG-A-AAGACCGGAGUGGUUUCUCUGCUUUUCCUC-C--AGAGGUC 200
dISFV-NC_016997.1 AUA-UUGACACCUAGG-A-AAGACCGGAGAUACCUUUGGUUUCUCCGAC-A--GAAUCUC 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-AAGACCGGAGAAUACUUGGCUUACCCGAG--CAACCAC 200
MBFV-NC_018705.3 AUA-UUGACUUAACAGGGGAAGACUAGAGGUUAGGACUCCUUGCGAGUGAGAACC--U 208
80.....190.....200.....210.....220.....230.....2



..... 257
MBFV-NC_009028.2 CACUAGGCACAGAUCCGCCG--AAGUAGUGGCUGGUG--GUGAA---AAACACAUGGAUC 257
MBFV-NC_040776.1 CACUAGGCACAGAUCCGCCG--AAUCUGUGGCUGGUG--CUGCA---AAACACA-GGAUC 257
MBFV-NC_034151.1 CACUCGGCACAGAACGCCGU--AAGU-GUGGCUGGUG--GUGAAAAUAAACACA-GAGUC 257
MBFV-NC_001563.2 CACACGGCACAGUGCGCCG--CAUAGGUGGCUGGUG--GUGC--UAGAACACA-GGAUC 257
MBFV-NC_009942.1 CACACGGCACAGUGCGCCG--CAUAGGUGGCUGGUG--GUGC--GAGAACACA-GGAUC 257
MBFV-NC_000943.1 CACAAGGCACCGAGCGCCG--ACACUGUGACUGAUG--GGGGAGAAGACCACA-GGAUC 257
MBFV-NC_006551.1 CACAAGGCACAGAGCGCCG--AAUUGUGGCUGGUG--GGGAACUAGACCACA-GGAUC 257
MBFV-NC_001437.1 UACUAGGCACAGAGCGCCG--AGUAGUAGCUGGUG--GUGAGGAAGAACACA-GGAUC 257
MBFV-NC_007580.2 CACAAGGCACAGAGCGCCG--AAAGUGUGGCUGGUG--GUGAA-AAAAUCACA-GGAUC 257
MBFV-NC_012534.1 CAUCUGGCACAGAGCGCCG--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GGUC 257
MBFV-NC_015843.2 CACUUGGCACAGAUCCGCCG--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GGAUC 257
MBFV-NC_009029.2 CUCUCGGCACAGAACGCCG--UGA-GGAGCCGGAGG--CCCAA---AACAC-CAGAU 257
MBFV-NC_032088.1 UUCUCGGCACAGAACGCCG--UAA-UGUAGUGGGU--GCCAA---AACAC-CGAAU 257
MBFV-NC_009026.2 CACCCGGCACAGAACGCCG--CA-UGGUGGGAGGGG--UCCUAAGACACCA--GAUUC 257
dISFV-NC_024017.1 GCCAUGGCACAGAGCGCCG--AAGAGAGCUGUUGUUAUAAUUUACAAAAAUAACAU 256
dISFV-NC_017086.1 UACACGGCACAGCGCGCCG--AAAAGGUAGUUUG--AGUUUUAUAAACAAAGUAU 256
MBFV-NC_002640.1 UCCA-GGCACAGAGCGCCG--AAGAUGAUUGGUGU--UGU---UGAUCAACAGGUU 256
MBFV-NC_001474.2 UCCA-GGCACAGAGCGCCG--AAAUGGAAUGGUGC--UGU---UGAAUCAAAGGUU 256
MBFV-NC_001477.1 UCCA-GGCACAGAACGCCA--AAAUGGAAUGGUGC--UGU---UGAAUCAAAGGUU 256
MBFV-NC_001475.2 UCCA-GGCACAGAACGCCG--AAAUGGAAUGGUGC--UGU---UGAAUCAAAGGUU 256
MBFV-NC_012532.1 GCCA-GGCACAGAUCCGCCG--ACUUCGGCGGCGCGU--GUGG--GGAAUCCAUUGGUU 256
MBFV-NC_035889.1 GCCA-GGCACAGAUCCGCCG--AUAGCGGCGGCGCGU--GUGG--GGAAUCCAUUGGUU 256
MBFV-NC_012533.1 CCCU-GGCACAGAGCGCC--GAGAAGGUGGUUGGU--GUGG--AGCAACCCAACAGGA 256
dISFV-NC_040610.1 UAC-AAGCACAGAGCGCUU--GAGUAGGUUGUUGUUCUUGACGAGAGAAUUGUAU 254
MBFV-NC_002031.1 UGU-GAGCACAGUUUGCUCAAGAAUAGACAGACCUUUGGAU---ACAAACCAAAAACC 255
dISFV-NC_016997.1 CUAAUGGCACAGAGCGCCG--AAGAAACGGAGUAGGAUUG---AAAAACAUAAGUA 255
MBFV-NC_008719.1 CCUUAAGCACAGAUUGCUUG--AAGUACGGGUGGGAUGU---GGAACCAACUAGU 255
MBFV-NC_012735.1 CCUAAAGCACAGAUUGCUUG--AAGUACGGGCGGGAUGU---AGAACCAACUAGU 255
dISFV-NC_027999.1 CGAAAGGCACAGAUUGCCUG--AACAUUCGGUGGUUGUAUG---GUAACCAACAGU 255
MBFV-NC_018705.3 -----CACACAAAACAGCAUUAUGACUCCUGGGAAGAGCAUAGGACCC--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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