

dISFV-NC_027999.1 -----GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAAGUACUACU---AUGGUUG 47
MBFV-NC_008719.1 -----UUUCUACUGAAAGGGUAGAGA--A--AAGGAGUUUUGCUUUCUCAUGGCAA 47
MBFV-NC_012735.1 -----UUUCUACUG--AAAGGAGUAGA--AGAAAGGAGAUUCAUUCCCAAUGCAA 47
MBFV-NC_002031.1 -----UAAUCGUUCGUUGAGCGAUUAGCAGAGAAACUGACCAGAACAUC-- 45
dISFV-NC_016997.1 -----A-AACAUAACGCAUUAAGAAUAGGUUUGAAAGAGUUGAAUCA---AUGAGAU 47
dISFV-NC_017086.1 -----AUAACACGUUUGAAUAGGUUUUUAUUGUG-AAUCGAAAUACCG---AU-- 43
MBFV-NC_009028.2 U-----UUAACACAGUCUGAAUAGUUUUUU-AGCA-AGGGAUU-UCC----CAUGUCU- 46
MBFV-NC_040776.1 U-----UUAACACAGUCUGAAUAGUUUUUU-AGCA-GAGGUAC-UAC---CAUGUCU- 46
MBFV-NC_001563.2 A-----UUAACACAGUCGAGCUGUUUCUU-GGCACGAAGAUC-UC----GAUGUCU- 46
MBFV-NC_009942.1 A-----UUAACACAGUGCGAGCUGUUUCUU-AGCACGAAGAUC-UC----GAUGUCU- 46
MBFV-NC_007580.2 -----UAACUUGACUGCGAACAGUUUUUU-AGCA-GGGAAUUACCC---AAUGUCU- 46
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUGAACCA-GAGGUGAAAAAC---CAUGUCU- 46
MBFV-NC_012534.1 CC----GUAAACACAGUUUGAAC-GUUUUUUUGGAGA-GAGAU---UAU---UAUGUUU- 46
MBFV-NC_015843.2 AC----GUAAACACAGUUUGAAC-GUUUUUUUGGAUA-GAGAC---AAC---UAUGUCU- 46
MBFV-NC_000943.1 -----UUAACCGGUUGAAACAGUUUUUUGAGCUUUUUGAUU-UCA---AAUGUCU- 46
MBFV-NC_006551.1 -----UUAACACAGUGCCCGGCAGUUUCUUUGAGC-GUUGAUU-UUC----AAUGUCU- 46
MBFV-NC_001437.1 -----UUAGUGCAGUUUAAACAGUUUUUUGAACCGGAAGAUA-ACC----AUGACU- 46
MBFV-NC_009026.2 -----AGAAAAAGUUUUCGAGGAAAGACGAGAA-GCGAAUUUCUCUG---AAUGUCU- 46
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAGUUUUUUGAACAG---C-----AAUGACC- 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAGUUUUUUGAUCAGUGGAU---AAUGACU- 46
MBFV-NC_012532.1 AA-----CAGGUUUA-----AUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAA- 46
MBFV-NC_035889.1 AC-----AGGUUUUA-----UUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAA- 46
MBFV-NC_012533.1 AA-----CAGUUUUC-----GUGUUGAGAAAGAAACAGAUUCUUCGG---AAUGAAA- 46
MBFV-NC_002640.1 -----UCUAACAGUUUGUUUGAAUAGAGAGAGCAG-AUCUCUGGAAAAUUG- 43
MBFV-NC_001477.1 AC-----GUAGUUCUAACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG---AUGAAC- 46
MBFV-NC_001475.2 AC-----GUAGUGCUGACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG---AUGAAC- 46
MBFV-NC_001474.2 -C-----GUAGUUCUAACAGUUU-UUUAAUAGAGAGCAG-AUCUCUG---AUGAAC- 46
dISFV-NC_024017.1 -----UCGCGCAUUUUGAU-AGAGUUGACUAGAUAAACGAAUUAUC---GAUGUG- 46
dISFV-NC_040610.1 -----GAACAAAUUUUAAAGACGUCAAACUAGAGAAUUUAUU-UC---AAUGUU- 46
MBFV-NC_018705.3 CC----GUAAACACAGUUUGAACAGUUUUUUGAGAGAGACUACU---AUGCA- 46
.....10.....20.....30.....40.....50.....6



.....((((.....((((.....((((.....((((..... 105
dISFV-NC_027999.1 AGAAACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUACGAU--UAGGCGUAC 105
MBFV-NC_008719.1 CAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUGGU--GACGGCCAA 105
MBFV-NC_012735.1 CAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUUGGU--AGCGGCUAG 105
MBFV-NC_002031.1 -----UGGUCGUAAGAGCAGGGAUAAACCCUGGGCGUCAAUUGGUUACGACGAGGAG 98
dISFV-NC_016997.1 CA AUGGUCUUUCGAGCUAGGAGGCGCGUUAACCGGGCGCUGCAUUAUAAAUAAACGUAAGC 107
dISFV-NC_017086.1 -----GCGAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUUAAGUGAGACGUGCGU 98
MBFV-NC_009028.2 --A-----AAAAACAGGAAAGUCGCGUCAAAGCCGAGCAGUCAAUUGGUUAAAACCGCUGG 101
MBFV-NC_040776.1 --A-----AAAAACAGGAGACCCGCGGAAGGCGGGUUGUCAAUUGGUUAAAACCGCCAG 101
MBFV-NC_001563.2 --A-----AAAAACAGGAGGGGCCCGGUAUAAACCGGGCUGUCAAUUGGUUAAAACCGCGUA 101
MBFV-NC_009942.1 --A-----AGAAACAGGAGGGGCCCGCAAGAGCCGGGCGUUCAAUUGGUUAAAACCGCGAA 101
MBFV-NC_007580.2 --A-----AAAAACAGGAAACCCGCUAGAAACCGGGUUGUCAAUUGGUUAAAACCGCGCG 101
MBFV-NC_034151.1 --A-----AAAAACAGGAGAGACCCCGGAAGACCCGGGUGUCAAUUGGUUAAAACCGGCAU 101
MBFV-NC_012534.1 --AACAAAAAAACAGGAAACCCCGGGCAACCCGGGUUGUCAAUUGGUUAAAGCGCGGUG 104
MBFV-NC_015843.2 --AACAAAAAAAACAGGAAGACCCCGCUCAGGCCGGGUUGUCAAUUGGUUAAAGCGCGGAA 104
MBFV-NC_000943.1 --A-----AAAAACAGGAAGACCCCGGAGAACCCGGGUGCUCAAUUGGUUAAAACCGGCA 101
MBFV-NC_006551.1 --A-----AGAAACAGGAGGGGCCCGGAAGAAACCGGGCCUCAAUUGGUUAAAACCGGCA 101
MBFV-NC_001437.1 --A-----AAAAACAGGAGGGGCCCGUAUAAACCGGGCUAUCAAUUGGUUAAAACCGCGCC 101
MBFV-NC_009026.2 --A-----GAAACACAGGAAGACCCCGGAGAAACCGGGUGCUCAAUUGGUUAAAACCGGCA 101
MBFV-NC_009029.2 --A-----AAAAACAGGAAGGGCCCGUAGAAAUCCGGGCCUCAAUUGGUUAAAGCGCGGUG 101
MBFV-NC_032088.1 --A-----AAAAACAGGAAAGUCGCGCAAUCGUCGAGCUGUCAAUUGGUUAAAGCGCGGAG 101
MBFV-NC_012532.1 --A-----ACCCCAAGAGAAGAAUCCCGAGGA-UCCGGAUUGUCAAUUGGUUAAAACGC---G 98
MBFV-NC_035889.1 --A--A-CCCAAAAAAAGAAUCCCGAGGAUCCCGGAUUGUCAAUUGGUUAAAACGC---G 98
MBFV-NC_012533.1 --A-----CCCAAAAAAAGG-----GCCCAUCGCGGCCUAUCAAUUGGUUGAAGCGCUUUG 95
MBFV-NC_002640.1 --A--A-CCAACGAAAAAAAGGUG-GUUAGACCACCUUCAAUUGGUUGAAACCGCGGA 95
MBFV-NC_001477.1 --A--A-CCAACGGAAAAAAGACG-GGUCGACCGUCUUUCAAUUGGUUGAAACCGCGGA 98
MBFV-NC_001475.2 --A--A-CCAACGGAAGAAGACG-GGAAACCGCUCUAUCAAUUGGUUGAAACCGCGGA 98
MBFV-NC_001474.2 --A--A-CCAACGGAAAAAAGGCG-AAAAACACGCCUUCAAUUGGUUGAAACCGCGGA 98
dISFV-NC_024017.1 -----ACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUAAUUAUGGUAACACGCAUCA 101
dISFV-NC_040610.1 -----AGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUUGGUCGAAAGCGGAUUA 101
MBFV-NC_018705.3 --AACAAAAAAACAGGAAGACCCCGGGCAACCCGGGUUGUCAAUUGGUUAAAGCGCGGCG 104
0.....70.....80.....90.....100.....110.....1



.....NNNNNNNN-----UGGCGAACC--CCAUGGAAAUUUAGC- 142
dISFV-NC_027999.1 UGG-----AA--NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UAUAGC- 143
MBFV-NC_008719.1 GGC-----CA--NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UAUAGC- 143
MBFV-NC_012735.1 AGU-----GA--NNNNNNNN-----C-CCAUUCCUCCCGAGCACACAUAGC- 143
MBFV-NC_002031.1 UUCGCUCCUUGU-----NNNNNNNN-----G--AGACCCUCCAGGGAACAAUAGUG 142
dISFV-NC_016997.1 UGC-----NNNNNNNN-----A-GGAACCCCCCAAAAUAGUAGG- 143
dISFV-NC_017086.1 UGCCUCGCGUCU--NNNNNNNN-----A-GAGGAGACCCCCCAAAAUAGAGG- 143
MBFV-NC_009028.2 CUAGCGUGU-----NNNNNNNN-----GUGGAGACCCCUAGCAC-CAAGCGCGAA-ACAA 149
MBFV-NC_040776.1 CCAGCGUUU-----NNNNNNNN-----A-GGAGACCCUUGGCACAAAGCGCAA-GACAA 150
MBFV-NC_001563.2 UGCCCCGCG-----NNNNNNNN-----GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAA 149
MBFV-NC_009942.1 UGCCCGCG-----NNNNNNNN-----GGAGACCCCGUGCCCAAAACACCA-ACAAA 149
MBFV-NC_007580.2 UAUCCCGCG-----NNNNNNNNUAGAGGAGACCCUUGCCGUAA--CGCAA--CA 149
MBFV-NC_034151.1 CCCGCGCUA-----NNNNNNNN-----AGACCCCUUGGCAGUGAA-CGCAA-UAAA 146
MBFV-NC_012534.1 CAUCCC-----NNNNNNNN-----GGAGACCCCUUGCGAGUUAACACCAA-ACAAA 149
MBFV-NC_015843.2 CGUCCC-----NNNNNNNN-----AGGAGACCCCUUGCGAGUGAACACCAC-A-GAA 149
MBFV-NC_000943.1 UACCCCGCG-----NNNNNNNN-----GACCCACUCUCAAAAGCAUC-AA---ACA 143
MBFV-NC_006551.1 UACCCCGCG-----NNNNNNNN-----ACCCCGCA-UCAUAAGCAUCAAA---AAA 142
MBFV-NC_001437.1 UACCCCGCG-----NNNNNNNN-----UAGAGGAGACCCCGCAUUUGC-AU---CAA 143
MBFV-NC_009026.2 CGGCCAACG-----NNNNNNNN-----CCCUUGGCAAAACAGUUAACCGACCAA-AAGAA 150
MBFV-NC_009029.2 CGUCCCGCG-----NNNNNNNNAGAGG--AGACCCUCCACAAAGAAGCGCAAACACAAA 153
MBFV-NC_032088.1 CAUCCCGCG-----NNNNNNNNAGAGGUUAGAGGAGACCCCAUACCGCAA-ACAAA 154
MBFV-NC_012532.1 GAG-UAGCCC--GUGNNNNNNNN-----GGUUAGAGGAGACCCCGGAAAACGCAAA 147
MBFV-NC_035889.1 GAG-UAGCCC--GUGNNNNNNNN-----GUUAGAGGAGACCCCGGAAAACGCAAA 146
MBFV-NC_012533.1 GAUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCGCGCCCAUAAACCAUAUAA 147
MBFV-NC_002640.1 GAAACCGCUAUCANNNNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCAA-CACAA-AA 151
MBFV-NC_001477.1 GAAACCGCGUGU--NNNNNNNN-----UAGAGGUUAGAGGAGACCCCGCACAACAACA 151
MBFV-NC_001475.2 GAAACCGUGUGU--NNNNNNNN--ACUAGAGGUUAGAGGAGACCCCGCA-AAUAA-AA 151
MBFV-NC_001474.2 GAAACCGUGUGU--NNNNNNNN--CUAGAGGUUAGAGGAGACCCCCGCA-AACAAAA 151
dISFV-NC_024017.1 CCAAGCGUG-----NNNNNNNN-----CCCACUGCUGAGUG-CAA 136
dISFV-NC_040610.1 CCACCCGCU-----NNNNNNNN-----ACCCCUCAUUAUUUAUUA 137
MBFV-NC_018705.3 CAUCCC-----NNNNNNNN-----CCCGCGGAAUCAAGCAAGGCCAAC 145
20.....130.....140.....150.....160.....170.....1



..))))))))).....))))))..... 198
dISFV-NC_027999.1 U--UCAAUUAUGACACCAGGGAAAGACCGGAGAAACUUCUUGCUUACUCGCAGC--AACC 198
MBFV-NC_008719.1 GGACCAUUAUAGACCGCCAGGG-AAAGACCGGAGACACUCCUUGAUU-CUCACCUU-UCUC 199
MBFV-NC_012735.1 GGACCAUUAUAGACGCCAGGG-AAAGACCGGAGACACUCCUUGAUU-CUCACCUU-UCUC 199
MBFV-NC_002031.1 GGACCAUUAUAGACGCCAGGG-AAAGACCGGAGUGGUUCUCUGCUU-UUCCUCC-AGAG 197
dISFV-NC_016997.1 CUUAUUAUAGCAGACCCUAGG-AAAGACUGGAGAUAC-CUUUGUUCUUCGCACAC--UAAC 199
dISFV-NC_017086.1 GCACUUAUAGCAGACUUGGG-AAAGACAGAGGUACUCGCUGAUUACCGCCAC--CAG- 199
MBFV-NC_009028.2 ACAGCAUUAUAGACCGCCUGGGA--AAGACCGGGAGAUCCUUGCU--UUCCAUC--ACCA 202
MBFV-NC_040776.1 ACAGCAUUAUAGACACCUGGGA--AAGACCGGAGACCCUUGCU--CUGCAC--ACCA 203
MBFV-NC_001563.2 ACAGCAUUAUAGACACCUGGGA--UAGACUAGGGGAUCUUCUGCU--CUGCACA--ACCA 202
MBFV-NC_009942.1 ACAGCAUUAUAGACACCUGGGA--UAGACUAGGAGAUCCUUGCU--CUGCACA--ACCA 202
MBFV-NC_007580.2 ACAGCAUUAUAGACACCUGG-A--AAGAC-AGGAGAUCCCUUGCU--UUUCAAC--ACCA 201
MBFV-NC_034151.1 ACAGCAUUAUAGACACCUGGGA--UAGACUGGGAGACCCUUGCUU-AUUUCAAC--ACCA 201
MBFV-NC_012534.1 ACAGCAUUAUAGACACCUGGGA--GAGACUAGGAGACCCUUGUCC--UAACAAC--ACCA 203
MBFV-NC_015843.2 ACAGCAUUAUAGACACCUGGGA--UAGACUAGGAGACCCUUGCU--UAACAAC--ACCA 203
MBFV-NC_000943.1 ACAGCAUUAUAGACACCUGGGAA-AAGACUAGGAGAUCCUUGCUCUAUCCAAC--AUCA 200
MBFV-NC_006551.1 ACAGCAUUAUAGACACCUGGGAAUAGACUAGGAGAUCCUUGCUCUAUCCAAC--AUCA 200
MBFV-NC_001437.1 ACAGCAUUAUAGACACCUGGGAA-UAGACUGGGAGAUCCUUGCUCUAUCCAAC--AUCA 200
MBFV-NC_009026.2 ACAGCAUUAUAGACACCUGGGA--UAGACCGGAGAAUUGCUGCC--UCGCAAC--ACCU 203
MBFV-NC_009029.2 ACAGCAUUAUAGACACCUGGGA--AAGACUAGGGGAUUGCUGCU--CUGGACU--UCCG 206
MBFV-NC_032088.1 ACAGCAUUAUAGACACCUGGGA--UAGACUAGGAGAUCCGUGCU--CUGGAAC--CCC 207
MBFV-NC_012532.1 ACAGCAUUAUAGACG--UGGGA--AAGAC-CAGAGACUCCAUGAGUUCCACCAC--GCUG 200
MBFV-NC_035889.1 ACAGCAUUAUAGACG-CUGGGA--AAGAC-CAGAGACUCCAUGAGUUCCACCAC--GCUG 200
MBFV-NC_012533.1 ACAGCAUUAUAGACACCUGGGA--AAGAC-CGGAGACUCUGUUGGCUUCCCCAC--ACCA 202
MBFV-NC_002640.1 ACAGCAUUAUAGACG-CUGGGA--AAGAC-CAGAGAUCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001477.1 ACAGCAUUAUAGACG-CUGGGA--GAGAC-CAGAGAUCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001475.2 ACAGCAUUAUAGACG-CUGGGA--GAGAC-CAGAGAUCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001474.2 ACAGCAUUAUAGACG-CUGGGA--AAGAC-CAGAGAUCUGCU-GUCUCUACAGC--AUCA 204
dISFV-NC_024017.1 AUAGCAUUAUAGACACUUGGGAAGAGACCAGAGAUUAUGAUUGUUGUCAUUGCU-GACA 195
dISFV-NC_040610.1 ACAGCAUUAUAGACACCAGGG-AUAGACGGAGAAU-ACACAAUUGCCUGGACUAUAA 195
MBFV-NC_018705.3 CUGCAUUAUAGCUAAACCAGGGGAAGGACGAGAGGUACAGAAUCCUUGGCGC--AGUG 203
80.....190.....200.....210.....220.....230.....2



ACCGAAAG-GCACAGUAUGCCUGA-AA-CAUUCGGUGGUUGAUGUAGUAAACAACCAGUU 255
dISFV-NC_027999.1 ACCCUUAA-GCACAGAUUGCUUG--AA-UGCAGGGUGGGGAAGUUGGGAACCAACUAGUG 255
MBFV-NC_008719.1 GCCCUAAA-GCACAGAUUGCUUG--AA-UGCAGGGCGGGGAAGUUGAGAACCAACUAGUG 255
MBFV-NC_012735.1 GUCUGAAG-GCACAGUUGCUAAGAA-UAAGCAGACCUUUGGAUGACAAACCAAAAGC 255
dISFV-NC_002031.1 UCCUAAUG-GCACAGAGCGCCAU--AA-GAAACGAGUUAUGAUGGAAAAACAUAAGU 255
dISFV-NC_016997.1 ACUACACG-GCACAGCGCGCCGG--AA-AAGGUAGUUGGAGGUUGUAAAACAACAAGUA 255
dISFV-NC_017086.1 GCCACUAG-GCACAGAUCCGCC--CA-AGUAGUGGCUUGUGUGGA-AAAAACAUGGA- 255
MBFV-NC_009028.2 GCCACUAG-GCACAGAUCCGCC--AA-AUCUGGCGUGGUGUGC-AAAAACA-GGA- 255
MBFV-NC_040776.1 GCCACACG-GCACAGUGCGCCG--AC-AUAGGUGGCUUGGUGGUGCUAGAACACA-GGA- 255
MBFV-NC_001563.2 GCCACACG-GCACAGUGCGCCG--AC-AUAGGUGGCUUGGUGGUGGUGGAGAACACA-GGA- 255
MBFV-NC_009942.1 GCCACACG-GCACAGUGCGCCG--AC-AAUGGUGGCUUGGUGGUGGAGAACACA-GGA- 255
MBFV-NC_007580.2 GCCACAAG-GCACAGAGCGCCG--UA-AAGUUGGCGUGGUGGUGGAAAAACACAGGA- 255
MBFV-NC_034151.1 GCCACUCG-GCACAGAACGCCG--UA-AGU-GUGGCUUGGUGGUGGAAAAUAAACACAGAG 255
MBFV-NC_012534.1 GCCAUCCG-GCACAGAGCGCCG--UA-AGU-GUGGCUUGGUGGUGGUGGAGAACACAGUG- 255
MBFV-NC_015843.2 GCCAUUCG-GCACAGAUCCGCC--AA-AGU-GUGGCUUGGUGGUGGUAAGAACACAGGA- 255
MBFV-NC_000943.1 GUCACAAG-GCACCAGAGCGCCG--AA-CACUGUGACUGAUGGGGGGAGAAGACCACAGGA 255
MBFV-NC_006551.1 ACCACAAG-GCACAGAGCGCCG--AA-AAUUGUGGCUUGGUGGGAACUAGACCACAGGA 255
MBFV-NC_001437.1 GCUACUAG-GCACAGAGCGCCG--AA-GUAUGUAGCGUGGUGGAGGAAGAACACAGGA 255
MBFV-NC_009026.2 CCCACCCG-GCACAGAAGCGCCG--AC-A-UGGUGGAGGGGUGCUAAGACACCA--GAU 255
MBFV-NC_009029.2 GCU-CUCG-GCACAGAACGCCG--UU-GA--GGAGCCGGAGGCC--AAAACACCCGA 255
MBFV-NC_032088.1 GUU-CUAG-GCACAGAACGCCG--U-AA--UGAACUGGGGUGCC--AAAACACCCGA 255
MBFV-NC_012532.1 GCCGCGAG-GCACAGAUCCGCCG--AA-CUUCGGCGCGGUGGUGGGGAAAUCCAUGGU 255
MBFV-NC_035889.1 GCCGCGAG-GCACAGAUCCGCCG--AA-UAGCGCGCGCGUGUGGGGAAAUCCAUGGU 255
MBFV-NC_012533.1 GCCCCUAG-GCACAGAGCGCC--A-GAUCGGGGUGGUGGAGCAACCCACAGG 255
MBFV-NC_002640.1 --AUCCAG-GCACAGAGCGCCG--CA-AGAUG--GAUUGUGUGUUGAUAUCCAACAGGU 255
MBFV-NC_001477.1 --UUCAG-GCACAGAACGCCA--AA-AAAUG--GAAUGGUCUGUUGAAUACAACAGGU 255
MBFV-NC_001475.2 --UUCAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUCUGUUGAAUACAACAGGU 255
MBFV-NC_001474.2 --UUCAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUCUGUUGAAUACAACAGGU 255
dISFV-NC_024017.1 ACAGCAUUGGACAGAGCGCCUGAAGAGAGCUGUUGUUAUAAUUACAAAAAAUAACAUA 255
dISFV-NC_040610.1 ACUACAAGCAGAGCGCUUCGAGAAGUAGUUGUUCUGGAGAGAAUAGUUAUAA 255
MBFV-NC_018705.3 AACACCAC-ACAAAAACAGCA--AA--AUAGACCCUGGGAAGAGCAUAGGACCCUC 255
40.....250.....260.....270.....280.....290.....3



... 258
dISFV-NC_027999.1 UCU 258
MBFV-NC_008719.1 UCU 258
MBFV-NC_012735.1 UCU 258
MBFV-NC_002031.1 ACU 258
dISFV-NC_016997.1 UCU 258
dISFV-NC_017086.1 UCU 258
MBFV-NC_009028.1 UCU 258
MBFV-NC_040776.1 UCU 258
MBFV-NC_001563.2 UCU 258
MBFV-NC_009942.1 UCU 258
MBFV-NC_007580.2 UCU 258
MBFV-NC_034151.1 UCU 258
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MBFV-NC_009029.2 UCU 258
MBFV-NC_032088.1 UCU 258
MBFV-NC_012532.1 UCU 258
MBFV-NC_035889.1 CUU 258
MBFV-NC_012533.1 ACU 258
MBFV-NC_002640.1 UCU 258
MBFV-NC_001477.1 UCU 258
MBFV-NC_001475.2 UCU 258
MBFV-NC_001474.2 UCU 258
dISFV-NC_024017.1 UCU 258
dISFV-NC_040610.1 UGG 258
MBFV-NC_018705.3 UGU 258
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