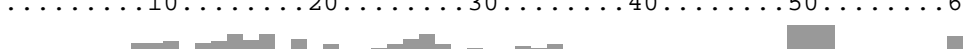


dISFV-NC_040610.1 --GAACAAUUUUUAAAGAAGUCACAAACUAGAAAU-----UUAAUU-UCAAUG----- 43
dISFV-NC_024017.1 UCGCGCAUUUUUGAUAGAGUU---GACUAGAUAA-----CGAAUUUACGAUG----- 43
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGAAACGAGAGUUUCUGG--UCAUGAAAAACCC 51
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCUGG--UCAUGAAAAA-CC 50
MBFV-NC_012533.1 AACAGUUUU-----CGUUGUGAGAAAGAAAAACAGAUUCUUCGG--AAAUGAAAA--CC 49
MBFV-NC_002640.1 -----UCUA-ACAGUUUUUUUGAAUAGAGAGCAG-AUCUCUGGAAAAAUG--AA--C 46
MBFV-NC_001477.1 ACGUAGUUUCUA-ACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACAA--C 49
MBFV-NC_001475.2 ACGUAGUGCUG-ACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUGAACAA--C 49
MBFV-NC_001474.2 -CGUAGUUUCUA-ACAGUUU-UUUAAUUAGAGAGCAG-AUCUCUG-----AUGAAUA--C 49
MBFV-NC_009028.2 U--U-----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCC-CAUG-----UC 45
MBFV-NC_040776.1 U--U-----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUAC-CAUG-----UC 45
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUUGAACAGAGGUGAAAAAC-CAUG-----UC 45
MBFV-NC_007580.2 ---U-----AACUUGACUGCGAACAGUUUUUUU-AGCAGGGAAUUACCC-AAUG-----UC 45
MBFV-NC_001437.1 ---U-----UAGUGCAGUUUAAACAGUUUUUU-AGAACCAGGAUAAC-CAUG-----AC 45
MBFV-NC_006551.1 -----U-----UAACACAGUGCGGACGAGUUUCUU-UGAGCGUAGUUUUU-AAUG-----UC 45
MBFV-NC_001563.2 A--U-----UAACACAGUGCGGAGCUGUUUCUU-GGCACG-AAGAUCUC-GAUG-----UC 45
MBFV-NC_009942.1 A--U-----UAACACAGUGCGGAGCUGUUUCUU-AGCACG-AAGAUCUC-GAUG-----UC 45
MBFV-NC_000943.1 -----U-----UAACACAGUGGUGAACGAGUUUUUUUGAGCUGUUUUUCAA-AAUG-----UC 45
MBFV-NC_012534.1 CCGU-----AAACACAGUUUGAACGUUUUUUUGGAGA---GAGAUUUU-AAUG-----UU 45
MBFV-NC_015843.2 ACGU-----AAACACAGUUUGAACGUUUUUUUGGAUA---GAGACAAC-UAUG-----UC 45
MBFV-NC_009029.2 UGAUUUACAUAACACAGUUGAACAGUUUUUUUGAAC-----AG--C-AAUG-----AC 45
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAGUUUUUUUGAUC---AGUGGAU-AAUG-----AC 45
MBFV-NC_009026.2 -----AGAAAAGUUUUCGAGGAAAGACGAGAAGCGAAUUCUCUGAAUG-----GC 45
dISFV-NC_016997.1 AAAC-----AUAACGCAUUGAAUAGGUUU---GAAAGAGUUGA--AUCAAUGAGAUAUAA 51
dISFV-NC_017086.1 -----AUAACACGUUUGAAUAGGUUUUUAUGUGAAUUCGAAAU--ACCGAUG----- 43
dISFV-NC_027999.1 -----GGAUUAAUUGUUGUUUGAGAGAAACAGCGAGAAUUAUUCU- AUGGUU---A 48
MBFV-NC_008719.1 -----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCAAUGGCAA---C 48
MBFV-NC_012735.1 -----UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCAAUGGCAA---C 48
MBFV-NC_002031.1 -----UAAUCGUUCGUAGAGCG--AUUAGCAGAGAGAACUGACAGAAACAGUC----- 45
MBFV-NC_018705.3 CCGUA-----AACACAGUUUGAACGUUUUUUUGAGAGAGACUACU---AUG-----UC 45
.....10.....20.....30.....40.....50.....6



.....(((.....((((.....((((.....((((..... 102
dISFV-NC_040610.1 -GUUAGACAACUUGGAAAGACCUGUGCAAGACGGUCUGUCAAUUGCGAAAGCGGAUUAAC 102
dISFV-NC_024017.1 -GUGACUAAAUCAGAAAGCCCGCGCAAAGCGGGCUGUUAUUUUGUAUAACGCAUAC 102
MBFV-NC_012532.1 CAAAGAAGAA-----AUCCGAGGA-UCCGGAUUGUCAAUUGCGUAAAACGCGGAGU 102
MBFV-NC_035889.1 CAAAAAAGAA-----AUCCGAGGAUUCGCGAUUGUCAAUUGCGUAAAACGCGGAGU 102
MBFV-NC_012533.1 CAAAAAAGG-----GCCGCAUCGCGGCCUAUCAAUUGCGUGAAGCGCUUUGG 96
MBFV-NC_002640.1 CAACGAAAAA-----AGGUG--GUUAGACCACCUUCAAUUGCGUGAAACGCGGAGAG 96
MBFV-NC_001477.1 CAACGGAATA-----AGACG--GGUCGACCGUCUCAAUUGCGUGAAACGCGCGAG 99
MBFV-NC_001475.2 CAACGGAAGA-----AGACG--GAAACCCGUCUAUCAAUUGCGUGAAACGCGGAGAG 99
MBFV-NC_001474.2 CAACGGAATA-----AGGCG--AAAAACACGCCUCAAUUGCGUGAAACGCGGAGAG 99
MBFV-NC_009028.2 U---AAAAAACACAGGAAAGUCCUGGCAAGACGCGACUGUCAAUUGCGUAAAACGCGGCG 102
MBFV-NC_040776.1 U---AAAAAACACAGGAGGACCCCGCGAAGCGGGUUGUCAAUUGCGUGAAACGCGCCAG 102
MBFV-NC_034151.1 U---AAAAAACACAGGAGGACCCCGAAGAACCGGGUCUGUCAAUUGCGUGAAAGCGGCAU 102
MBFV-NC_007580.2 U---AAAAAACACAGGAAACCCCGUAGAAACCGGGUUGUCAAUUGCGUAAAACGCGGCGU 102
MBFV-NC_001437.1 U---AAAAAACACAGGAGGGCCCGGUAACAAACCGGGCUAUCAAUUGCGUGAAACGCGGCC 102
MBFV-NC_006551.1 U---AAGAAACACAGGAGGACCCCGAAGAACCGGGCCUCAAUUGCGUGAAACGCGGCAU 102
MBFV-NC_001563.2 U---AAGAAACACAGGAGGGCCCGGUAACAAACCGGGCUGUCAAUUGCGUAAAACGCGGUA 102
MBFV-NC_009942.1 U---AAGAAACACAGGAGGGCCCGCAAGAGCCGGGUCUGUCAAUUGCGUAAAACGCGGAU 102
MBFV-NC_000943.1 U---AAAAAACACAGGAGGACCCCGAAGAACCGGGUCUGUCAAUUGCGUGAAACGCGGCAU 102
MBFV-NC_012534.1 UAACAAAAAACACAGGAAACCCCGGCAACCCGGGUUGUCAAUUGCGUAAAGCGCGGUGC 105
MBFV-NC_015843.2 UAACAAAAAACACAGGAAGACCCCGCUCAGGCGGGUUGUCAAUUGCGUAAAGCGCGGAAC 105
MBFV-NC_009029.2 C---AAAAAACACAGGAGGACCCCGAAGAACCGGGCCUGUCAAUUGCGUAAAGCGCGGUG 102
MBFV-NC_032088.1 U---AAAAAACACAGGAAAGUCCCGCAUCGUCGAGCUGUCAAUUGCGUAAAGCGCGGAGC 102
MBFV-NC_009026.2 U---AGAAACACAGGAAGACCCCGAGGAAACCGGGUCUGUCAAUUGCGUAAAGCGUACGGC 102
dISFV-NC_016997.1 GGUCUUUCGAGCUAGGAGGGCCGUUAAA---CGGGCCGUGCAUUAUAAACGUAAGCU 108
dISFV-NC_017086.1 -GCGAAUAAACCCAAGAAGCCCGAAGA---CGGGCUAUGCAUUAUGAGACGUGCGUU 99
dISFV-NC_027999.1 GAAACUAGAGUAGAUUCUCGGUUUAAAACCCGAGGUGUCAAUUGAUUAGGCGUACUG 107
MBFV-NC_008719.1 AAGAGGACAGAGCAAGAGUCUCGGGUCACGACCCGGGGCGUCAAUUGGUGACGGCCAGG 107
MBFV-NC_012735.1 AAAGGGGAUGAAUAGUCUCGGGUCUGAUCCCGAGGGCGUCAAUUGGUGAGCGGCUAGAG 107
MBFV-NC_002031.1 -----UGGUCGUAAAGCUCAGGAAACAAACCCUGGGCGUCAAUUGGUGUACGACGAGAG 99
MBFV-NC_018705.3 AAACAAAAAACACGGAAGACCCCGGCAACCCCGGGUUGUCAAUUGCUAAAGCGGCGCG 105
0.....70.....80.....90.....100.....110.....1



..... 138
dISFV-NC_040610.1 CACCCGCU-----NNNNNNN-----ACCCCUCAUUAUUUAUUA 138
dISFV-NC_024017.1 CAAGCGUG-----NNNNNNN-----CCCACUGCUGAGAG-CAAA 137
MBFV-NC_012532.1 AGCCCGUG-----NNNNNNN-----GCUUAGAGGAGACCCCCCGGAAACGCAAAA 148
MBFV-NC_035889.1 AGCCCGUG-----NNNNNNN-----GUUAGAGGAGACCCCCCGGAAACGCAAAA 147
MBFV-NC_012533.1 AUCUAGCGCUGGCCNNNNNNN-----GAGACCCCGCGCCCAUAACCAUAUAAA 148
MBFV-NC_002640.1 AAACCGCGUAUCAANNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCAACA--CAAAA 152
MBFV-NC_001477.1 AAACCGCGUGU--NNNNNNN-----UAGAGGUUAGAGGAGACCCCCGCACAACAACAAA 152
MBFV-NC_001475.2 AAACCGUGUGU--NNNNNNN--ACUAGAGGUUAGAGGAGACCCCCG--CAAAUAAAA 152
MBFV-NC_001474.2 AAACCGCGUGU--NNNNNNN--CUAGAGGUUAGAGGAGACCCCC--CGAAACAAAAA 152
MBFV-NC_009028.2 UAGCGUGU-----NNNNNNN--GUGGAGACCCUCAGCAC--CAAGCGCAA--ACAAA 150
MBFV-NC_040776.1 CAGCGUUU-----NNNNNNN--A--GGAGACCCUUGGCACACAAGCGCAA--GACAAA 151
MBFV-NC_034151.1 CCGCGCUA-----NNNNNNN-----AGACCCUUGCCAGUGAA--CGAAA-UAAAA 147
MBFV-NC_007580.2 AUCCCGCG-----NNNNNNNUUAGAGGAGACCCCUUGCCGUUA--CGC-AAACAA 150
MBFV-NC_001437.1 ACCCCGCG-----NNNNNNN--UAGAGGAGACCCCG--CAUU--UGCAUC---AAA 144
MBFV-NC_006551.1 ACCCCGCG-----NNNNNNN-----ACCCG--CAUCAUAAGCAUCA-AAAAA 143
MBFV-NC_001563.2 GCCCCGCG-----NNNNNNN--GAGGAGACCCCGUGCCA--AAAACACCA-A-AAGAAA 150
MBFV-NC_009942.1 GCCCCGCG-----NNNNNNN--GGAGACCCCGUGCCACAAAAACACCACA-ACAAA 150
MBFV-NC_000943.1 ACCCCGCG-----NNNNNNN-----GACCCA--CUCUAAAAGCAUC-AAACAA 144
MBFV-NC_012534.1 AUCC-----NNNNNNN--GGAGACCCCUUGCGAGUUACACCAAA-ACAAA 150
MBFV-NC_015843.2 GUCC-----NNNNNNN--AGGAGACCCUUGCGAGUGAACACCACA-A-GAAA 150
MBFV-NC_009029.2 GUCCCGCG-----NNNNNNNAGAG--AGACCCUCCCAAGAAAGCGCAACACAAAA 154
MBFV-NC_032088.1 AUCCCGCG-----NNNNNNNAGAGGUUAGAGGAGACCCCCCAUACCGCAA-ACAAA 155
MBFV-NC_009026.2 GGCCAACG-----NNNNNNNCCCUUGG--CAAAACAGUUAACGACCAAAAGAAA 151
dISFV-NC_016997.1 GC-----NNNNNNN--AGAAACCCCCCAAAAUAGUUAAGGC----- 144
dISFV-NC_017086.1 GCCUCGCGUCU--NNNNNNNAGAGGAGACCCCCCAAAAUAG--AAGGG----- 144
dISFV-NC_027999.1 -----GAA--NNNNNNN-----UGGCGAACCCCAUUGGAAAUUA--G 141
MBFV-NC_008719.1 -----CCA--NNNNNNN-----GAGGAGACCCUCCGAGCAC-UAUAGC-G 144
MBFV-NC_012735.1 -----UGA--NNNNNNN-----C-CAUUCUCCCGGAGCACAUAGC-G 144
MBFV-NC_002031.1 UCGCUCCUUGU--NNNNNNN-----G--AGACCCUCCAGGGAACAAUAGUG 143
MBFV-NC_018705.3 AUCC-----NNNNNNN-----CCCCGCGAAUCAAAGCAAGGCCAACCC 146
20.....130.....140.....150.....160.....170.....1



.))))))..)))))..)).....))..))..... 195
dISFV-NC_040610.1 CAGCAUA-UUGACACCAGGG-A-UAGACGGAGAAUACACAAUUCUGGGCACUAAACA 195
dISFV-NC_024017.1 UAGCAUA-UUGACACUUGGAA-GAGACCAGAGAUUAUGUUGUUUGUCAUAGCUGACA 195
MBFV-NC_012532.1 CAGCAUA-UUGACG--UGGGA--AAGACAG--AGACUCCAUGAGUUUCCACCAC--GCU 199
MBFV-NC_035889.1 CAGCAUA-UUGACG-CUGGGA--AAGACAG--AGACUCCAUGAGUUUCCACCAC--GCU 199
MBFV-NC_012533.1 CAGCAUA-UUGACACCUGGGA--AAGACCG--AGACUCUGUUGCUUUUCCCCAC--ACC 201
MBFV-NC_002640.1 CAGCAUA-UUGACG-CUGGGA--AAGACAG--AGAUCUCUGCU-GUCUCUGCAAC--AUC 203
MBFV-NC_001477.1 CAGCAUA-UUGACG-CUGGGA--GAGACAG--AGAUCUCUGCU-GUCUCUACAGC--AUC 203
MBFV-NC_001475.2 CAGCAUA-UUGACG-CUGGGA--GAGACAG--AGAUCUCUGCU-GUCUCCUAGC--AUC 203
MBFV-NC_001474.2 CAGCAUA-UUGACG-CUGGGA--AAGACAG--AGAUCUCUGCU-GUCUCCUAGC--AUC 203
MBFV-NC_009028.2 CAGCAUA-UUGACGCGUGGGA--AAGACCG--GAGAUCUCUCUGCU--UUCCAUC--ACC 201
MBFV-NC_040776.1 CAGCAUA-UUGACACCUGGGA--AAGACCG--GAGACCCUCUGCU--CUGCAAC--ACC 202
MBFV-NC_034151.1 CAGCAUA-UUGACACCUGGGA--UAGACUGG--GAGACCCUCUGCU-UAUUUCAAC--ACC 200
MBFV-NC_007580.2 CAGCAUA-UUGACACCUGGAA--AGACAG--GAGAUCUCCUGCUUU--UUCAAC--ACC 200
MBFV-NC_001437.1 CAGCAUA-UUGACACCUGGGA--UAGACUGG--GAGAUCUUCUGCUUAUCUACAAC--AUC 199
MBFV-NC_006551.1 CAGCAUA-UUGACACCUGGGA--UAGACUAG--GAGAUCUUCUGCUUAUCCAAC--AUC 199
MBFV-NC_001563.2 CAGCAUA-UUGACACCUGGGA--UAGACUAG--GGAUUCUUCUGCUUAUCCAAC--AUC 199
MBFV-NC_009942.1 CAGCAUA-UUGACACCUGGGA--UAGACUAG--GAGAUCUUCUGCUUAUCCAAC--AUC 199
MBFV-NC_012534.1 CAGCAUA-UUGACACCUGGGA--GAGACUAG--GAGACCCUCUGUCCUA--ACAAC--ACC 202
MBFV-NC_015843.2 CAGCAUA-UUGACACCUGGGA--UAGACUAG--GAGACCCUCUGUCCUA--ACAAC--ACC 202
MBFV-NC_009029.2 CAGCAUA-UUGACACCUGGGA--AAGACUAG--GGAUUUGCUGCUCUG--GAC--UUC 204
MBFV-NC_032088.1 CAGCAUA-UUGACACCUGGGA--UAGACUAG--GAGAUCGUGCUGCUCUG--GAA--CCC 205
MBFV-NC_009026.2 CAGCAUA-UUGACACCUGGGA--UAGACCG--AGAAUUGCUGGCCUCG--CAAC--ACC 202
dISFV-NC_016997.1 UAUAUA-UUGACACCUAGGA--AAGACUGGAGAUAC--CUUUGGUUUUUGCAGAC--UAA 198
dISFV-NC_017086.1 CACUAUA-UUGACACUUGGGA--AAGACAGAGGUACUCGUGAUUACCGCCAC--CAG 199
dISFV-NC_027999.1 CUAUAUA-UUGACACCAGGGA--AAGACGGAGAAACUUCUGCUUAUCCGACG--AAC 197
MBFV-NC_008719.1 GAGCAUA-UUGACGCGGGA--AAGACGGAGACACUCCUUGAUU--CUCACCUU--UCU 198
MBFV-NC_012735.1 GAGCAUA-UUGACGCGGGA--AAGACGGAGACACUCCUUGAUU--CUCACCUU--UCU 198
MBFV-NC_002031.1 GAGCAUA-UUGACGCGGGA--AAGACGGAGGUUUCUGCUGCUUUUCCGAG--AG 197
MBFV-NC_018705.3 UGCAUA-UUGCUAAACAGGGGAAGACACUAGAGGUUAGGACACCCUUGCAGUGAGA 206
80.....190.....200.....210.....220.....230.....2



ACUACAA--GCACAGAGCGCUUCGAGAAAGUAGUUGUUUCUUGACGAGAGAAUUGUAUAU 253
dISFV-NC_024017.1 ACAGCCAUGGCACAGAGCGCCUGAAGA--GAGCUGUUGUUUAUAAUUUACAAAAAUACA 253
MBFV-NC_012532.1 GGCCGCCAGGCACAGAUCCGCCAACUUCGGCGGCCGGUGUGGGGAA-AUCCAUGGUU--- 255
MBFV-NC_035889.1 GGCCGCCAGGCACAGAUCCGCCAACUAGCGCGGCCGGUGUGGGGAA-AUCCAUGGUU--- 255
MBFV-NC_012533.1 AGCCCGUGGCACAGAGCGCC--AGAUCCGGGUUGUGUGGAGACA-ACCAACAGG--- 255
MBFV-NC_002640.1 A--AUCCAGGCACAGAGCGCCGCAAGAUG--GAUUGGUGUUGUUGA-UCCAACAGGU--- 255
MBFV-NC_001477.1 A--UUCAGGCACAGAACGCCAAAAAUG--GAAUGGUGUGUUGA-AUCAACAGGU--- 255
MBFV-NC_001475.2 A--UUCAGGCACAGAACGCCAAAAAUG--GAAUGGUGUGUUGA-AUCAACAGGU--- 255
MBFV-NC_001474.2 A--UUCAGGCACAGAACGCCAAAAAUG--GAAUGGUGUGUUGA-AUCAACAGGU--- 255
MBFV-NC_009028.2 AGCCACUAGGCACAGAUCCGCCAAGUAGUGGUGGUGGUGGAAAA-ACACA-UGGA--- 255
MBFV-NC_040776.1 AGCCACUAGGCACAGAUCCGCCAAGUAGUGGUGGUGGUGGAAAA-ACACA-UGGA--- 255
MBFV-NC_034151.1 AGCCACUCCGCACAGAACGCCGUAAAGU-GUGGUGGUGGUGGAAAAUAAACA-CAGAG--- 255
MBFV-NC_007580.2 AGCCACUAGGCACAGAGCGCCGUAAAGUUGGUGGUGGUGGAAAA-AAAAUCA-CAGGA--- 255
MBFV-NC_001437.1 AGCUACUAGGCACAGAGCGCCGAAAGUAGUGGUGGUGGUGGAGGAAGAACA-CAGGA--- 255
MBFV-NC_006551.1 AACCACAAGGCACAGAGCGCCGAAAUUGUGGUGGUGGUGGGAACUAGACCA-CAGGA--- 255
MBFV-NC_001563.2 AGCCACACGGCACAGUCGCGCCACAUAGGUGGUGGUGGUGG--CUAGAACA-CAGGA--- 255
MBFV-NC_009942.1 AGCCACACGGCACAGUGCGCCACAAGUGGUGGUGGUGGUG--CGAGAACA-CAGGA--- 255
MBFV-NC_000943.1 AGUCACAAGGCACCGAGCGCCGACUAGCUGAGCUGAUGGGGAGAGAACCA-CAGGA--- 255
MBFV-NC_012534.1 AGCCAUCCGGCACAGAGCGCCGUAAAG-UGUGGUGGUGGUGG--GUAGAACA-CAGUG--- 255
MBFV-NC_015843.2 AGCCACUUGGCACAGAUCCGCAAG-UGUGGUGGUGGUGG--GUAGAACA-CAGGA--- 255
MBFV-NC_009029.2 CGGCUCUCCGCACAGAACGCCGUUGAGGA---GCCGAGGCCAAAAACA-CCAGA--- 255
MBFV-NC_032088.1 CAGUUCUAGGCACAGAACGCCG--UAAUG---ACUGGGUGCCAAAAACA-CCGAA--- 255
MBFV-NC_009026.2 UCCCAACCGGCACAGAACGCCGACA---UGGUGGAGGGGUGCGUAAAGACACAGAU--- 255
dISFV-NC_016997.1 CUCCUAAUGGCACAGAGCGCCAUAGAAACCGAGUUAUGUAGUAAACCAUAAGUA--- 255
dISFV-NC_017086.1 -ACUACACGGCACAGCGCGCCGAAAGGUAGUUGGAGUUGUAAACCAACAGUA--- 255
dISFV-NC_027999.1 CACCGAAAGGCACAGAUUGCCUGA--AACAUCCGUGUGUAGUUGUAAACCAACAG--- 253
MBFV-NC_008719.1 CACCCUUAAGGCACAGAUUGCUUG--AAUGAGGGUGGGGAAUGUUGGAACCAACUAG--- 253
MBFV-NC_012735.1 CGCCCUAAGGCACAGAUUGCUUG--AAUGAGGGCGGGGAAUGUAGAACCAACUAG--- 253
MBFV-NC_002031.1 -GUCUGUGAGCACAGUUGCUUACAAGAAUAGGACAGACCUUUGGAUACCAACCAAA--- 253
MBFV-NC_018705.3 ACCACAAAAAACAGCAUUAUUGACUAGG--AAAGACUAGGAGACCC 253
40.....250.....260.....270.....280.....290.....3



..... 258
dISFV-NC_040610.1 CUUGG 258
dISFV-NC_024017.1 UAUG 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
MBFV-NC_009028.2 --UCU 258
MBFV-NC_040776.1 --UCU 258
MBFV-NC_034151.1 --UCU 258
MBFV-NC_007580.2 --UCU 258
MBFV-NC_001437.1 --UCU 258
MBFV-NC_006551.1 --UCU 258
MBFV-NC_001563.2 --UCU 258
MBFV-NC_009942.1 --UCU 258
MBFV-NC_000943.1 --UCU 258
MBFV-NC_012534.1 --UCU 258
MBFV-NC_015843.2 --UCU 258
MBFV-NC_009029.2 --UCU 258
MBFV-NC_032088.1 --UCU 258
MBFV-NC_009026.2 --UCU 258
dISFV-NC_016997.1 --UCU 258
dISFV-NC_017086.1 --UCU 258
dISFV-NC_027999.1 UUUU 258
MBFV-NC_008719.1 UGUU 258
MBFV-NC_012735.1 UGUU 258
MBFV-NC_002031.1 CCAC 258
MBFV-NC_018705.3 UCUG 258
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