

dISFV-NC_027999.1 -----GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAAGUACUACU---AUGGUUG 47
MBFV-NC_008719.1 -----UUUCUACUGAAAGGGUAGAGA--A--AAGGAGUUUUGCUUUCUCAUGGCAA 47
MBFV-NC_012735.1 -----UUUCUACUG--AAAGGAGUAGA--AGAAAGGAGAUUCAUUCCCAAUGGCAA 47
MBFV-NC_002031.1 -----UAAUCGUUCGUUGAGCGAUUAGCAGAGAAACUGACCAGAACAUUC-- 45
dISFV-NC_016997.1 -----A-AACAUAACGCAUAGAAUAGGUUUGAAAGAGUUGAAUCA---AUGAGAU 47
dISFV-NC_017086.1 -----AUAACACGUUUGAAUAGGUUUUUAUUGUG-AAUCGAAAUACCG--AUG----- 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-GAGGUAAAAAC--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 -----UUAGUGCAGUUUAAACAGUUUUUUGAACCGGAAGAU-ACC---AUGACU- 46
MBFV-NC_009026.2 -----AGAAAAAGUUUUCGAGGAAAGACGAGAA-GCGAAUUUCUCUG--AAUGCU- 46
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAGUUUUUUGAACAG----C-----AAUGACC- 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAGUUUUUUGAUCAGUGAU-----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-----GUGUUGAGAAAGAAAACAGAUUCUUCGG--AAAUGAAA- 46
MBFV-NC_002640.1 -----UCUAACAGUUUGUUUGAAUAGAGAGAGCAG-AUCUCUGGAAAAAUG-- 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 -----GAACA AUUUUAAAGAGUCUAAAACUAGAGAAUUUAUU-UC---AAUGUU- 46
MBFV-NC_018705.3 CC----GUAAACACAGUUUGAACAGUUUUUUGGAGAGAGACUACU-----AUGCAC- 46
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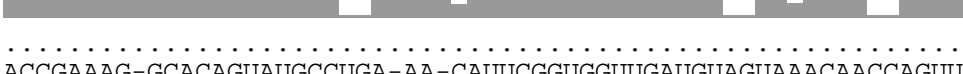
.....((((.....((((.....((((..... 105
dISFV-NC_027999.1 AGAAACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAGAU--UAGGCGUAC 105
MBFV-NC_008719.1 CAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUGGU--GACGGCCAA 105
MBFV-NC_012735.1 CAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUUGGU--AGCGGCUAG 105
MBFV-NC_002031.1 -----UGGUCGUAAGAGCUCAGGGAUAAACCCUGGGCGUCAAUUGGUUACGACGAGGAG 98
dISFV-NC_016997.1 CA AUGGUCUUUCGAGCUAGGAGGCGCGUUAACCGGGCGCUGAUUAUAUAAACGUAAGC 107
dISFV-NC_017086.1 -----GCGAAUAAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGU 98
MBFV-NC_009028.2 --A-----AAAAACAGGAAAGUCGCGUGCAAAGCCGAGCAGUCAAUUGCUAAAAACGCCUGG 101
MBFV-NC_040776.1 --A-----AAAAACAGGAGAGACCCGCGGAAGGCGGGUUGUCAAUUGCUAAAAACGCCAG 101
MBFV-NC_001563.2 --A-----AAAAACAGGAGGGGCCCGGUAUAAACCGGGCUGUCAAUUGCUAAAAACGCCGUA 101
MBFV-NC_009942.1 --A-----AGAAACAGGAGGGGCCCGCAAGAGCCGGGCUGUCAAUUGCUAAAAACGCCGAA 101
MBFV-NC_007580.2 --A-----AAAAACAGGAAACCCGUAAGAAACCGGGUUGUCAAUUGCUAAAAACGCCGGC 101
MBFV-NC_034151.1 --A-----AAAAACAGGAGAGACCCCGGAAGAACCCGGGUGCUCAAUUGCUAAAAACGCCACA 101
MBFV-NC_012534.1 --AACA AAAAACAGGAAACCCCGGGCAACCCGGGUUGUCAAUUGCUAAAAGCGCGGUG 104
MBFV-NC_015843.2 --AACAAAAAACAGGAAGACCCCGCUCAGGCCGGGUUGUCAAUUGCUAAAAGCGCGGAA 104
MBFV-NC_000943.1 --A-----AAAAACAGGAAGACCCCGGAGAACCCCGGUGCUCAAUUGCUAAAAGCGCGCA 101
MBFV-NC_006551.1 --A-----AGAAACAGGAGGGGCCCGGAAGAAACCGGGCCAUCAAUUGCUGAAACCGCGCA 101
MBFV-NC_001437.1 --A-----AAAAACAGGAGGGGCCCGUAUAAACCGGGCUAUCAAUUGCUGAAACCGCGCC 101
MBFV-NC_009026.2 --A-----GAAACACAGGAAGACCCCGGAGAAACCCGGGUGCUCAAUUGCUAAAAGCGUACGG 101
MBFV-NC_009029.2 --A-----AAAAACAGGAAGGGCCCGUAGAAAUCCGGGCCGUCAAUUGCUAAAAGCGCGGUG 101
MBFV-NC_032088.1 --A-----AAAAACAGGAAAGUCGCGCAAUCGUCGGACUGUCAAUUGCUAAAAGCGCGGAG 101
MBFV-NC_012532.1 --A-----ACCCCAAAGAGAAGAAUCCCGAGGA-UCCGGAUUGUCAAUUGCUAAAACGC---G 98
MBFV-NC_035889.1 --A--A-CCCAAAAAGAAAUCCGAGGAUUCGGAUUGUCAAUUGCUAAAACGC---G 98
MBFV-NC_012533.1 --A-----CCCAAAAAGG-----GCCCAUCGCGGCCUAUCAAUUGCUGAAAGCGCUUUG 95
MBFV-NC_002640.1 --A--A-CCAACGAAAAAAGGUG--GUUAGACCACCUUUCAAUUGCUGAAACCGCGGA 95
MBFV-NC_001477.1 --A--A-CCAACGGA AAAAGACG--GGUCGACCGUCUUUCAAUUGCUGAAACCGCGGA 98
MBFV-NC_001475.2 --A--A-CCAACGGAAGAAGACG--GGAAACCGGUCUAUCAAUUGCUGAAACCGCGGA 98
MBFV-NC_001474.2 --A--A-CCAACGGA AAAAGGCG--AAAAACACGCCUUUCAAUUGCUGAAACCGCGGA 98
dISFV-NC_024017.1 -----ACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUAAUAUGCUAACACGCAUCA 101
dISFV-NC_040610.1 -----AGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUUGCUGAAAGCGGAUUA 101
MBFV-NC_018705.3 --AACAAAAAACAGGAAGACCCCGGGCAACCCGGGUUGUCAAUUGCUAAAAGCGCGCGC 104
0.....70.....80.....90.....100.....110.....1



.....NNNNNNNN-----UGGCGAACC--CCAUGGAAAAUUUAGC- 142
dISFV-NC_027999.1 UGG-----AA--NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UUAGC- 143
MBFV-NC_008719.1 GGC-----CA--NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UUAGC- 143
MBFV-NC_012735.1 AGU-----GA--NNNNNNNN-----C-CCAUUCUCCUCCCGAGCACACAUAGC- 143
MBFV-NC_002031.1 UUCGCUCCUUGU-----NNNNNNNN-----G--AGACCUCCAGGGAACAAUAGUG 142
dISFV-NC_016997.1 UGC-----NNNNNNNN-----A-GGAACCCCCCAAAAUGUAUAGG- 143
dISFV-NC_017086.1 UGCCUCGCGUCU--NNNNNNNN-----A-GAGGAGACCCCCCAAAAUGAAGG- 143
MBFV-NC_009028.2 CUAGCGUGU-----NNNNNNNN-----GUGGAGACCCUCCAGCAC-CAAGCGCGAA-ACAA 149
MBFV-NC_040776.1 CCAGCGUUU-----NNNNNNNN-----A-GGAGACCCUUGGCACACAAGCGCAA-GACAA 150
MBFV-NC_001563.2 UGCCCCGCG-----NNNNNNNN-----GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAA 149
MBFV-NC_009942.1 UGCCCCGCG-----NNNNNNNN-----GGAGACCCCGUGCCACAAACACCACA-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-----AGGAGACCCUUGCGAGUGAACACCACA-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-----UAGAGGAGACCCCGCAUUGC-AU---CAA 143
MBFV-NC_009026.2 CGGCCAACG-----NNNNNNNN-----CCCUUGGC AAAACAGUUAACCGACCAA-AAGAA 150
MBFV-NC_009029.2 CGUCCCGCG-----NNNNNNNNAGAGG--AGACCCUCCACAAAGAAGCGCAAACACAAA 153
MBFV-NC_032088.1 CAUCCCGCG-----NNNNNNNNAGAGGUUAGAGGAGACCCCAUUAACCGCAA-ACAAA 154
MBFV-NC_012532.1 GAG-UAGCCC--GUGNNNNNNNN-----GGUAGAGGAGACCCCGGAAAACGCAAA 147
MBFV-NC_035889.1 GAG-UAGCCC--GUGNNNNNNNN-----GUUAGAGGAGACCCCGGAAAACGCAAA 146
MBFV-NC_012533.1 GAUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCGCGCCCAUAAACCAUAAA 147
MBFV-NC_002640.1 GAAACCGCUAUC AANNNNNNNN--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 GAAACCGGAGU--NNNNNNNN--CUAGAGGUUAGAGGAGACCCCGCA-AACAAAA 151
dISFV-NC_024017.1 CCAAGCGUG-----NNNNNNNN-----CCCAUCUGUGAGUG-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



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dISFV-NC_027999.1 U--UCAAUAUUGACACCAGGGAAAGACCGGAGAAACUUCUUGCUUACUCGCAGC--AACC 198
MBFV-NC_008719.1 GGACCAUAUUGACGCCAGGG-AAAGACCGGAGACACUCCUUGAUU-CUCACCUU-UCUC 199
MBFV-NC_012735.1 GGACCAUAUUGACGCCAGGG-AAAGACCGGAGUGGUUCUCUGCUU-UUCCUCC--AGAG 197
MBFV-NC_002031.1 CUAUUAUAUCGACACCUAGG-AAAGACUGGAGAUAC-CUUUGUUUCUUGCAGAC--UAAC 199
dISFV-NC_016997.1 GCACUAUAUCGACACUUGGG-AAAGACAGAGGUACUCGCUGAUUCACCGCCAC--CAG- 199
dISFV-NC_017086.1 ACAGCAUAUUGACGCCUGGGA--AAGACCGGGAGAUCUUCUGCU--UUCCAUC--ACCA 202
MBFV-NC_009028.2 ACAGCAUAUUGACACCUUGGGA--AAGACCGGGAGACCCUCUGCU--CUGCAC--ACCA 202
MBFV-NC_040776.1 ACAGCAUAUUGACACCUUGGGA--AAGACAGGAGAUCCUUCUGCU--CUGCAC--ACCA 202
MBFV-NC_001563.2 ACAGCAUAUUGACACCUUGGGA--AAGACAGGAGAUCCUUCUGCU--CUGCAC--ACCA 202
MBFV-NC_009942.1 ACAGCAUAUUGACACCUUGGGA--AAGACAGGAGAUCCUUCUGCU--CUGCAC--ACCA 202
MBFV-NC_007580.2 ACAGCAUAUUGACACCUUGGGA--AAGACAGGAGAUCCUUCUGCU--UUUCAAC--ACCA 201
MBFV-NC_034151.1 ACAGCAUAUUGACACCUUGGGA--AAGACUGGGAGACCCUCUGCUU-AUUUCAAC--ACCA 201
MBFV-NC_012534.1 ACAGCAUAUUGACACCUUGGGA--GAGACUAGGAGACCCUCUGUCC--UAACAAC--ACCA 203
MBFV-NC_015843.2 ACAGCAUAUUGACACCUUGGGA--AAGACUAGGAGACCCUCUGCU--UAACAAC--ACCA 203
MBFV-NC_000943.1 ACAGCAUAUUGACACCUUGGGA--AAGACUAGGAGAUUCUUCGCUUAUCCAAC--AUCA 200
MBFV-NC_006551.1 ACAGCAUAUUGACACCUUGGGAUAGACUAGGAGAUUCUUCGCUUAUCCAAC--AUCA 200
MBFV-NC_001437.1 ACAGCAUAUUGACACCUUGGGA--AAGACUGGGAGAUUCUUCGCUUAUCCAAC--AUCA 200
MBFV-NC_009026.2 ACAGCAUAUUGACACCUUGGGA--AAGACCGGAGAUUUGUGUCC--UCCGCAAC--ACCU 203
MBFV-NC_009029.2 ACAGCAUAUUGACACCUUGGGA--AAGACUAGGGGAUUGUGUGCU--CUGGACU--UCCG 206
MBFV-NC_032088.1 ACAGCAUAUUGACACCUUGGGA--AAGACUAGGAGAUCCGUGCU--CUGGAAC--CCCA 207
MBFV-NC_012532.1 ACAGCAUAUUGACG--UGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCAC--GCUG 200
MBFV-NC_035889.1 ACAGCAUAUUGACG-CUGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCAC--GCUG 200
MBFV-NC_012533.1 ACAGCAUAUUGACACCUUGGGA--AAGAC-CGGAGACUCUGUUGGCUUUCCCAC--ACCA 202
MBFV-NC_002640.1 ACAGCAUAUUGACG-CUGGGA--AAGAC-CAGAGAUCCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001477.1 ACAGCAUAUUGACG-CUGGGA--GAGAC-CAGAGAUCCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001475.2 ACAGCAUAUUGACG-CUGGGA--GAGAC-CAGAGAUCCUGCU-GUCUCUACAGC--AUCA 204
MBFV-NC_001474.2 ACAGCAUAUUGACG-CUGGGA--AAGAC-CAGAGAUCCUGCU-GUCUCUACAGC--AUCA 204
dISFV-NC_024017.1 AUAGCAUAUUGACACUUGGGAAGAGACCAGAGAUUAUGAUUGUUUGUCAUUGCU-GACA 195
dISFV-NC_040610.1 ACAGCAUAUUGACACCAGGG-AUAGACGGAGAAU-ACACAAUUCUUGGCGACUAAACA 195
MBFV-NC_018705.3 CUGCAUAUUGAGUAAACAGGGGAAGGACGAGAGGUUACAGAAUCCUUGGUGCU--AGUG 203
80.....190.....200.....210.....220.....230.....2



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dISFV-NC_027999.1 ACCCUUAA-GCACAGAUUGCUUG--AA-UGCAGGGUGGGGAAGUUGGGAACCAACUAGUG 255
MBFV-NC_008719.1 GCCCUAAA-GCACAGAUUGCUUG--AA-UGCAGGGCGGGGAAGUUGAGAACCAACUAGUG 255
MBFV-NC_012735.1 GUCUGUA-GCACAGAUUGCUAAGAA-UAAGCAGACCUUUGGAUGACAAACCAAAAGC 255
MBFV-NC_002031.1 UCCUAAUG-GCACAGAGCGCCAU--AA-GAACCGAGUUAGUAUGGAAAAACAUAAGU 255
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MBFV-NC_009028.2 GCCACUAG-GCACAGAUCCGCC--AA-AUCUGUGGCUUGGUGUGC-AAAACACA-GGA- 255
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MBFV-NC_007580.2 GCCACUCG-GCACAGAACGCCG--UA-AGU-GUGGCUUGGUGUGGUAUAAACACAGAG 255
MBFV-NC_034151.1 GCCAUUCG-GCACAGAGCGCCG--AA-AGU-GUGGCUUGGUGUGGUGUAGAACACAGUG-- 255
MBFV-NC_012534.1 GCCACUUG-GCACAGAUCCGCC--AA-AGU-GUGGCUUGGUGUGGUAGAACACAGGA-- 255
MBFV-NC_015843.2 GUCACAAG-GCACCAGAGCGCCG--AA-CACUGUGACUGAUGGGGGAGAAGACCACAGGA 255
MBFV-NC_000943.1 ACCACAAG-GCACAGAGCGCCG--AA-AAUUGUGGCUUGGUGGGGAACUAGACCACAGGA 255
MBFV-NC_006551.1 GCUACUAG-GCACAGAGCGCCG--AA-GUAUGUAGCGUGGUGGAGGAAGAACACAGGA 255
MBFV-NC_001437.1 CCCACCCG-GCACAGAAGCGCCG--AC-A-UGGUGGAGGGGUGCUUAAGACACCA--GAU 255
MBFV-NC_009026.2 GCU-CUCG-GCACAGAACGCCG--UU-GA--GGAGCCGGAGGCC--AAAACACCCGA 255
MBFV-NC_009029.2 GUU-CUAG-GCACAGAACGCCG--U-AA--UGAACUGGGUGGCC--AAAACACCCGA 255
MBFV-NC_032088.1 GCCGCGAG-GCACAGAUCCGCC--AA-CUUCGGCGGCCGUGUGGGGAAAUCCAUGGU 255
MBFV-NC_012532.1 GCCGCGAG-GCACAGAUCCGCC--AA-UAGCGCGGCCGUGUGGGGAAAUCCAUGGU 255
MBFV-NC_035889.1 GCCCCUAG-GCACAGAGCGCC--A-GAUCGGGGUGUGGUGGAGCAACCCACAGG 255
MBFV-NC_012533.1 --AUCCAG-GCACAGAGCGCCG--CA-AGAUG--GAUUGUGUGUGUAGAUAACACAGGU 255
MBFV-NC_002640.1 --UUCGAG-GCACAGAACGCCA--AA-AAAUG--GAAUGGUGCUGUUGAAUACAAGGU 255
MBFV-NC_001477.1 --UUCGAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUGCUGUUGAAUACAAGGU 255
MBFV-NC_001475.2 --UUCGAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUGCUGUUGAAUACAAGGU 255
MBFV-NC_001474.2 ACAGCAUUGGCAAGAGCGCCUGAAGAGAGCUGUUGUUAUAAUUACA AAAAUACAUA 255
dISFV-NC_024017.1 ACUACAAGCAGAGAGCGCUUCGAGAAGUAGUUGUUUCUGGAGAGAAUAGUUAUAUC 255
dISFV-NC_040610.1 AACACCAC-ACAAAAACAGCAU--AAUAGACCCUGGGAAGAGCAUAGGACCCUC 255
MBFV-NC_018705.3 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
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dISFV-NC_024017.1 UCU 258
dISFV-NC_040610.1 UGG 258
MBFV-NC_018705.3 UGU 258
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