

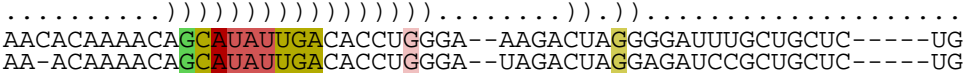
MBFV-NC_009029.2 UGAUUAACAUAACACAGUGUGAACAAAGUUUUUUGAACAG-----C---AAUGACC-- 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAAGUUUUUUGAUCAGUGG-----AU---AAUGACU-- 46
MBFV-NC_009028.2 U--U-----UAACACAGUCUGAAU--AGUUUUUU--AGCAAGGGA-UUUCC---CAUGUCU-- 46
MBFV-NC_040776.1 U--U-----UAACACAGUCUGAAU--AGUUUUUU--AGCAGAGGU-ACUAC---CAUGUCU-- 46
MBFV-NC_034151.1 -----UAACACAGUGUGAAU--AGUUUUUUUGAACAGAGGUGAAAC---CAUGUCU-- 46
MBFV-NC_001563.2 A--U-----UAACACAGUGCGAGC--UGUUUUUUGGCACGAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_009942.1 A--U-----UAACACAGUGCGAGC--UGUUUUUUGAAGCACGAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_001437.1 U--U-----AGUCAGUUUAAACA--GUUUUUUUGAACGGAAGA--UAAC---CAUGACU-- 46
MBFV-NC_007580.2 U--A-----ACUUGACUGCGAACAA--GUUUUUUAGCAGGGAUUU--ACCC---AAUGUCU-- 46
MBFV-NC_000943.1 -----UAACGCGGUUUGAAC--AGUUUUUUGGAGCUUUUGAUUUCA--AAUGUCU-- 46
MBFV-NC_006551.1 ----U-----UAACACAGUCCGGC--AGUUUCUUUGAGC--GUUGAUUUUC---AAUGUCU-- 46
MBFV-NC_012534.1 CCGU-----AAACACAGUUUGAAC--GUUUUUUUGGAGAGAG--AUUAU---UAUGUUU-- 46
MBFV-NC_015843.2 ACGU-----AAACACAGUUUGAAC--GUUUUUUUGGAUAGAG--ACAAC---UAUGUCU-- 46
MBFV-NC_009026.2 AGAA-----AAGUUUUC---GAGG--AAAGACGAG--AAGCGAAUUCUCUG--AAUGGCU-- 46
dISFV-NC_017086.1 AUAACACGUUU--GAAUAGGUUUUUAUUGUGAAUCGAAAUACCG-----AUGGCG-- 46
MBFV-NC_012532.1 AACAGGUUU-----AAUUUGGAUUUGGAAACGAGAGUUUCUGG--U---CAUG----- 43
MBFV-NC_035889.1 ACAGGUUUU-----AUUUUGGAUUUGGAAACGAGAGUUUCUGG--U---CAUG----- 43
MBFV-NC_012533.1 AACAGUUUU-----CGUGUUGGAAAGAAACAGAUUCUCUGG--A---AAUG----- 43
MBFV-NC_002640.1 -----UCUA--ACAGUUUGUUUGAAUAGAGAGCAG--AUCUCUGGAAA---AAUG----- 43
MBFV-NC_001474.2 --CGUAGUUCUA--ACAGUUU--UUUAAUUAGAGAGCAG--AUCUCUG-----AUG----- 43
ACGUAGUUCUA--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUG----- 43
MBFV-NC_001477.1 ACGUAGUGCUG--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUG----- 43
MBFV-NC_001475.2 GGAUUAAU---UGUUGUUUGAG-----AGAACAGCGAGAAGUACUACU---AUGGUUGA 48
dISFV-NC_027999.1 AUUUCUACUGAAAGGGU---AG-----AGA--A--AAGGAGUUUUGCUUCUCAUGGC-- 45
MBFV-NC_008719.1 -UUUCUACUG--AAAGGA---GU-----AGA--AGAAGGAGAUUCAUUCCCAUGGC--- 45
MBFV-NC_012735.1 UAAUCGUUCG-----UUGAG-----CGAUUAGCAGAGAACUGACCAGAACAUG-- 43
dISFV-NC_02031.1 --GAACAAUUUUUAAAGAAGACAAACUAAGAAAUU--AAUUUC-----AAUGGUU-- 46
dISFV-NC_040610.1 UCGCGCAUUUUUGAUAGAGUU---GACUAGAUAAACGAUUUAUC-----GAUGGUG-- 46
dISFV-NC_024017.1 AAACA-----UAACGCAUUUGAAUAGGUUUUGAAAGAGAUUGAAUC-----AAUGAGAUC 48
dISFV-NC_016997.1 CCGUA-----AACACAGUUUGAACGUUUUUUGGAGAGAGACUACU-----AUGUCA-- 46
MBFV-NC_018705.310.....20.....30.....40.....50.....6



MBFV-NC_009029.2 -A-----AAA-AACCAGGAAGGCCGG--UAGAAUUGGGGCGGCAAAUUGCUAAAGCGC 97
MBFV-NC_032088.1 -A-----AAA-AACCAGGAAGGUCGG--CAAUCGUCGGACUGCAAAUUGCUAAAGCGC 97
MBFV-NC_009028.2 -A-----AAA-AACCAGGAAGGUCGC--UGCAAAGCGGACUGCAAAUUGCUAAAGCGC 97
MBFV-NC_040776.1 -A-----AAA-AACCAGGAGGACCGG--CGGAAGGCGGGUUGCAAAUUGCGUAAAGCGC 97
MBFV-NC_034151.1 -A-----AAA-AACCAGGAGGACCGG--AAAGAACCGGGUCGUAAUUGCGUAAAGCGC 97
MBFV-NC_001563.2 -A-----AGA-AACCAGGAGGGGCCGG--UAAAACCGGGCUGCAAAUUGCUAAAGCGC 97
MBFV-NC_009942.1 -A-----AGA-AACCAGGAGGGGCCGG--CAAGAGCGGGGCGUGCAAAUUGCUAAAGCGC 97
MBFV-NC_001437.1 -A-----AAA-AACCAGGAGGGGCCGG--UAAAACCGGGCUGCAAAUUGCGUAAAGCGC 97
MBFV-NC_007580.2 -A-----AAA-AACCAGGAAGAACCCGG--UAGAAACCGGGUUGCAAAUUGCUAAAGCGC 97
MBFV-NC_000943.1 -A-----AAA-AACCAGGAGGACCGG--GAAGCCCCGGGUGUGCAAAUUGCUAAAGCGC 97
MBFV-NC_006551.1 -A-----AGA-AACCAGGAGGGGCCGG--AAGAACCCTGGGCUCAAAUUGCGUAAAGCGC 97
MBFV-NC_012534.1 -AA--CAAAA-AACCAGGAAGAACCGG--GGCAACCCGGGUCUCAAAUUGCGUAAAGCGC 100
MBFV-NC_015843.2 -AA--CAAAA-AACCAGGAAGAACCGG--CUCAGGCCGGGUGUGCAAAUUGCGUAAAGCGC 100
MBFV-NC_009026.2 -A-----GAA-AACCAGGAAGACCGG--AGGAACCCGGGUCUGCAAAUUGCUAAAGCGC 97
dISFV-NC_017086.1 -AA--UAAACCCAA---GAAGCCGG--AAGA---CGGGCUAUGCAUUAUGAGAGACGU 94
MBFV-NC_012532.1 -AA--AAACCCCAAGAAGAAAUCCGG--AGGA-UCCGGAUUGCAAAUUGCGUAAAGCGC 97
MBFV-NC_035889.1 -AA--AAA-CCCCAAAAAGAAAUCCGG--AGGAUUCGGAUUGCAAAUUGCGUAAAGCGC 97
MBFV-NC_012533.1 -AA--AA--CCAAAAA--AGGGCCGC---AUCGCGGGCUAUGCAUUAUGCGUAAAGCGC 91
MBFV-NC_002640.1 -----AA--CCAACGAAAAAAGGUGG--UUAGACCACCUU--UCAAAUUGCGUAAAGCGC 91
MBFV-NC_001474.2 -AA--UAA--CCAACGAAAAAAGGCGA--AAAACACGCCUU--UCAAAUUGCGUAAAGCGC 94
MBFV-NC_001477.1 -AA--CAA--CCAACGAAAAAAGAGCGG--GUCGACGCUUU--UCAAAUUGCGUAAAGCGC 94
MBFV-NC_001475.2 -AA--CAA--CCAACGGAAGAAGACGG--GAAAACCGUCUA--UCAAAUUGCGUAAAGCGC 94
dISFV-NC_027999.1 GAA--AACUAGAGUAGG---AUCUCGGGUUAAAACCCGAGGUGUGCAAAUUGAU-----UA 98
AAC--AAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGCAAAUUGGU-----GA 98
MBFV-NC_008719.1 AAC--AAAGGGGAUGAAUAAGUCUCGGGUCGAUCCCGAGGCGCAAAUUGGU-----AG 98
MBFV-NC_012735.1 -----UCUGGUCGUAAAGGUCUAGGGAAAAACCCUGGGCGCAAAUUGGUACGACGA 94
MBFV-NC_002031.1 -----AGACAAUUGGAAAGACCGU--UGCAAGACGGUCUGCAAAUUGCGGAAAGCGG 97
dISFV-NC_040610.1 -----ACUAAAUCAAGAAAGCCGC--CGCAAAGCGGGCUGUAAUUGCGUAAACACGC 97
dISFV-NC_024017.1 AAUGGUCUUUCGAGCUAGGAGGCCCGU---UAAACGGGGCGGCGCAAAUUGAAUAAAGCG 103
dISFV-NC_016997.1 ----AACAAAAACAGGAAGACCCGGG--GCAACCCGGGUUGCAAAUUGCGUAAAGCGC 100
MBFV-NC_018705.3 0.....70.....80.....90.....100.....110.....1



MBFV-NC_009029.2 GGUGCGUCCCGG-----NNNNNNNN--AGAGGAGA---CCCUCCCACAAAGAAGCGCA 145
MBFV-NC_032088.1 GGAGCAUCCCGG-----NNNNNNNN--AGAGGUUAGAGGAGACCCCCACAU--ACCGCA 147
MBFV-NC_009028.2 CUGGCUAGCGGUGU-----NNNNNNNN--GUGGAGA---CCCUCAGCAC--CAAGCGC 142
MBFV-NC_040776.1 CCAGCCCGCGUUU-----NNNNNNNN--A--GGAGA---CCCUGGCACACAAAGCGC 142
MBFV-NC_034151.1 ACAUCCCGCGCUA-----NNNNNNNN--AGAGGAGA---CCCUGGCAGUGAA--CGC 138
MBFV-NC_001563.2 GGUUAGCCCCCGG-----NNNNNNNN--GAGGAGA---CCCUGGCCA--AAAACACC 142
MBFV-NC_009942.1 GGAAUCCCGCCG-----NNNNNNNN--GGAGA---CCCUGGCCACAAACACC 141
MBFV-NC_001437.1 GGCCUACCCCGG-----NNNNNNNN--UAGAGGAGA---CCCUGCAUUUG-----137
MBFV-NC_007580.2 GCGGUUCCCGCG-----NNNNNNNN--UUAGAGGAGA---CCCUGCCGUUAAACG---143
MBFV-NC_000943.1 GGCAUACCCCGG-----NNNNNNNN--GA---CCCACUCUCAAAAGCAUC 138
MBFV-NC_006551.1 GGCAUACCCCGG-----NNNNNNNN--A---CCCUGCA--UCAUAAGCAUC 136
MBFV-NC_012534.1 GGUUAGCUCC-----NNNNNNNN--GGAGA---CCCUGCGAGUUAACACC 141
MBFV-NC_015843.2 GGAACGCUCC-----NNNNNNNN--AGGAGA---CCCUGCGAGUGAACACC 142
MBFV-NC_009026.2 ACGGCGGCCAACG-----NNNNNNNN--C---CCCUU--GGCAAAACAGUUAACGCAC 142
dISFV-NC_017086.1 GCGUUGCCCGCGGUCU-----NNNNNNNN--AGAGGAGACCCCCCAAAAU---138
MBFV-NC_012532.1 GGAGUAGCCCGUG-----NNNNNNNN--GGUAGAGGAGACCCCCCGGAAAC---142
MBFV-NC_035889.1 GGAGUAGCCCGUG-----NNNNNNNN--GUUAGAGGAGACCCCCCGGAAAC---141
MBFV-NC_012533.1 UUGUAGUUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCGCGCCCAUAACCAAC---142
MBFV-NC_002640.1 GAGAGAAACCGCGUAUCAANNNNNNNNACUAGAGGUUAGAGGAGACCCCCCAACA---146
MBFV-NC_001474.2 GAGAGAAACCGCGUGU---NNNNNNNN--CUAGAGGUUAGAGGAGACCCCCC--CGAAC---146
MBFV-NC_001477.1 GCGAGAAACCGCGUGU---NNNNNNNN--UAGAGGUUAGAGGAGACCCCCCGCAAC---146
MBFV-NC_001475.2 GUGAGAAACCGUGUGU---NNNNNNNNACUAGAGGUUAGAGGAGACCCCCCG--CAAA---146
dISFV-NC_027999.1 GCGUACUGGA---A---NNNNNNNN-----UGGCGAACCCCAUUGGAAAU---138
MBFV-NC_008719.1 CGGCCAAGGCC---A---NNNNNNNN-----GAGGAGACCCUCCGAGCAC--U-138
MBFV-NC_012735.1 CGGCUAGAGUG---A---NNNNNNNN-----C--CAUUCUCCUCCGAGCACAC--138
MBFV-NC_002031.1 GGAGUUCGCUCCUUGU---NNNNNNNN-----GAG---ACCCUCCAGGGAACAA--136
dISFV-NC_040610.1 AUUACCAACCGCU---NNNNNNNN-----ACCCCUCAUUUAU---131
dISFV-NC_024017.1 AUCACCAAGCGUG---NNNNNNNN-----CCCCACU--GCUGAG---130
dISFV-NC_016997.1 AAGCGUC---NNNNNNNN-----AGGAAACCCCCCAAAAUAGU---138
MBFV-NC_018705.3 GCGCAUCC-----NNNNNNNN-----CCCCCGGAAAUCA--131
20.....130.....140.....150.....160.....170.....1



MBFV-NC_009029.2 AACACAAAAACGCAUAUUGACACCUUGGA--AAGACUAGGGGAUUUGCUGCUC-----UG 198
MBFV-NC_032088.1 AA--ACAAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGUCCCGCUGCUC-----UG 199
MBFV-NC_009028.2 GAA--ACAAACAGCAUAUUGACGCCUGGA--AAGACCGGAGAUCCUCUGCUC---U-UCC 195
MBFV-NC_040776.1 AAAGACAACACAGCAUAUUGACACCUUGGA--AAGACCGGAGACCCUCUGCUC---C-UUC 196
MBFV-NC_034151.1 AAUAUAAAAACAGCAUAUUGACACCUUGGA--UAGACUGGAGACCCUCUGCUUA--U-UGC 194
MBFV-NC_001563.2 A--AAAGAAACAGCAUAUUGACACCUUGGA--UAGACUAGGGGAUUCUUC--UGCUC--UGCAC 197
MBFV-NC_009942.1 ACAACAAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGAUUCUUC--UGCUC--UGCAC 197
MBFV-NC_001437.1 --CAUCAAACAGCAUAUUGACACCUUGGAA--UAGACUGGAGAGAUUCUUGCUCU--AUCUC 193
MBFV-NC_007580.2 --CAACAACAGCAUAUUGACACCUUGGAA--AGACAG---GAGAUCCCCUGCU--UUUUC 194
MBFV-NC_000943.1 ---AAACAACAGCAUAUUGACACCUUGGAA--AAGACUAGGAGAUUCUUGCUCU--AUUCC 193
MBFV-NC_006551.1 ---AAAAAACAGCAUAUUGACACCUUGGAAUAGACUAGGAGAUUCUUGCUCU--AUUCC 193
MBFV-NC_012534.1 AAAACAAAACAGCAUAUUGACACCUUGGA--GAGACUAGGAGACCCUCUGUC---CUAAC 196
MBFV-NC_015843.2 ACAA--GAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGACCCUCUGUC---CUAAC 196
MBFV-NC_009026.2 CAAAGAAACAGCAUAUUGACACCUUGGA--UAGACC--GGAGAAUUGCGUCCU--CG--C 196
dISFV-NC_017086.1 ---GAAGGGCAUAUAUGCAGACUUGGGA--AAGACCAAGGACUUCGUGAUU--CACC 192
MBFV-NC_012532.1 ---GCAAAACAGCAUAUUGACG--UGGGA--AAGACCAAGACUCCAUGAGUUU--CCACC 194
MBFV-NC_035889.1 ---GCAAAACAGCAUAUUGACG--UGGGA--AAGACCAAGACUCCAUGAGUUU--CCACC 194
MBFV-NC_012533.1 ---AUAAAACAGCAUAUUGACACCUUGGA--AAGACCGGAGACUCCUUGGCUU--UCCCC 196
MBFV-NC_002640.1 ---CAAAACAGCAUAUUGACG--CUGGGA--AAGACCAAGAUCCUG---CUGU--CUCUG 196
MBFV-NC_001474.2 ---AAAAACAGCAUAUUGACG--CUGGGA--AAGACCAAGAUCCUG---CUGU--CUCUG 196
MBFV-NC_001477.1 ---AACAAACAGCAUAUUGACG--CUGGGA--GAGACCAAGAUCCUG---CUGU--CUCUA 196
MBFV-NC_001475.2 ---UAAACAGCAUAUUGACG--CUGGGA--GAGACCAAGAUCCUG---CUGU--CUCCU 196
dISFV-NC_027999.1 ---UAGC--U--UCAUAUUGACACCAAGGAA--AAGACCGAGAAACUUCUUGCUUACUCGC 191
MBFV-NC_008719.1 --AUAGC--GGACCAUAUUGACGCUUGG--A--AAGACCGAGACACUCCUUGAUU--CUCAC 192
MBFV-NC_012735.1 --AUAGC--GGACCAUAUUGACGCGAGGG--A--AAGACCGAGACACUCCUUGAUU--CUCAC 192
MBFV-NC_002031.1 --AUAGUGGGA--CAUAUUGACGCCAGGG--A--AAGACCGAGUGGUUCUUGCUUUUCCUC 192
dISFV-NC_040610.1 --UAUAUAACAGCAUAUUGACACCAAGG--A--UAGACCGAGACUCCUUGAGUU--CUCAC 185
dISFV-NC_024017.1 --AGCAAAUAGCAUAUUGACACUUGGGA--GAGACCAAGAUU--UGAUGUUGUUUGUCA 186
dISFV-NC_016997.1 --AUAGGCUUUAUAUUGACACCAUGGA--AAGACCGAGAUACCUUUGGUUUCAGGAG 193
MBFV-NC_018705.3 --AAGCAAGGCAUAUUGACCAACUGUAUAACACGGG--AGAAGACUAGGUUUCUUGGC 189
80.....190.....200.....210.....220.....230.....2



MBFV-NC_009029.2 GACUUCGCGGUCUC---GGACAGAACCGG---UUGA--GGACCGGAGGCCCAAA----- 247
MBFV-NC_032088.1 GAACCCGAGUUCUA---GGCAGAGAACCGG---UAA--UGAACUGGGGUGGCCAAA----- 247
MBFV-NC_009028.2 AUCACCGAGCUCUA---GGCAGAGAACCGG---CAAGUAGUGGUGGUGGUGAAAAACA 249
MBFV-NC_040776.1 AACACCGAGCUCUA---GGCAGAGAACCGG---AAAUUCUGGUGGUGGUGGAGAAAAACA 250
MBFV-NC_034151.1 AACACCGAGCUCUC---GGCAGAGAACCGG---UAAGU--GUGCGUGGUGGUGAA--AAUA 246
MBFV-NC_001563.2 A--ACCAGCAGAC---GGCAGAGUCCCGG---ACAUAAGUGGUGGUG--GUGCUAGA 247
MBFV-NC_009942.1 A--ACCAGCAGAC---GGCAGAGUCCCGG---ACAUAAGUGGUGGUG--GUGCGAGA 247
MBFV-NC_001437.1 AACAUAGCUCUA---GGCAGAGAACCGG---AAGUAUGUAGUGGUGGUGGAGGAAGA 247
MBFV-NC_007580.2 AACACCGAGCACA---GGCAGAGAGCGCG---UAAAGUGUGGUGGUGGUGAAAAA--A 247
MBFV-NC_000943.1 AACAUAGCUCACA---GGCAGAGAGCGCG---AACACUGUAGCUGAUGGGGGGAGAGA 247
MBFV-NC_006551.1 AACAUAGCUCACA---GGCAGAGAGCGCG---AAAAUUGGUGGUGGUGGAGUAGA-- 247
MBFV-NC_012534.1 AACACCGAGCAUCC---GGCAGAGAGCGCG---UAAG--UGUGGUGGUGGUGGUGA-- 247
MBFV-NC_015843.2 AACACCGAGCAUU---GGCAGAGAGCGCG---AAAG--UGUGGUGGUGGUGGUGA-- 247
MBFV-NC_009026.2 AACACCUCCACCC---GGCAGAGAACCGG---ACA--UGUGGAGGGGUGGUGAAGACA 249
dISFV-NC_017086.1 CCACCAAGATACAC---GGCAGAGCGCGCG---GAAAAGGUAUUGGAGGUUGUAAAA 246
MBFV-NC_012532.1 ACGCUGGCGGCCA---GGCAGAGUCCCGG---AACUUCGCGCGCGGUGUGGGGAAAU 247
MBFV-NC_035889.1 ACACCGGCGGCCA---GGCAGAGUCCCGG---AAUAGCGCGCGCGGUGGUGGGGAAAU 247
MBFV-NC_012533.1 ACACCAACCCCU---GGCAGAGAGCGCG---AGAUCGCGCGGUGGUGGAGGAGAAC 247
MBFV-NC_002640.1 CAACAUAUAUCCA---GGCAGAGAGCGCG---CAAGAUGGAUUGGUGUUGUUG--AUC 247
MBFV-NC_001474.2 CAGCAUAUAUCCA---GGCAGAGAACCGA---GAAAAGGUAUUGGUGCUGUUG--AAU 247
MBFV-NC_001477.1 CAGCAUAUAUCCA---GGCAGAGAACCGA---AAAAUAGGAUUGGUGGUGUUG--AAU 247
MBFV-NC_001475.2 CAGCAUAUAUCCA---GGCAGAGAACCGA---GAAAAGGUAUUGGUGCUGUUG--AAU 247
dISFV-NC_027999.1 AGCAACACACGAAA---GGCAGAGUAGCGU---GAAACAUUGGUGGUGGUGAUGAUA 246
MBFV-NC_008719.1 CUUUCUACCCUUA---AGCAGAGAUUGCU---G--AAUGCAGGUGGGGAAGUUGGGAAC 246
MBFV-NC_012735.1 CUUUCUUGCCCUAA---AGCAGAGAUUGCU---G--AAUGCAGGCGGGGAAGUUGAGAAC 246
MBFV-NC_002031.1 C--AGAGGUGUGU---AGCAGAGUUGCU---CAAGAUAAGACGAGCUCUUUG--GAUGACAAA 246
dISFV-NC_040610.1 GCACUAAGCUCUACA---AGCAGAGUUGCU---GAGAGAAGUAGUAGUUCUUGACGAGAGA 244
dISFV-NC_024017.1 AUGCGACAACAGCCAU---GGCAGAGAGCGCGUGAAGAGAGCGUUGUUAUAAUUUACAAAA 246
dISFV-NC_016997.1 ACUAACUCCUAU---GGCAGAGAGCGCGAUAAGAACCGGAGUUAUGAUGGAAAAAGAC 249
MBFV-NC_018705.3 ACCCCUUGCGAG---UGAACACCACAA--AAAACAGCAUAUUGACACUUGGGAAAGACU 244
40.....250.....260.....270.....280.....290.....3



MBFV-NC_009029.2 -ACACCAGAUUC--- 258
MBFV-NC_032088.1 -ACACCGAAUUC--- 258
MBFV-NC_009028.2 CAUGGA---UCU--- 258
MBFV-NC_040776.1 CA--GGA---UCU--- 258
MBFV-NC_034151.1 AACACAGAGUCU--- 258
MBFV-NC_001563.2 -ACACAGGAUCU--- 258
MBFV-NC_009942.1 -ACACAGGAUCU--- 258
MBFV-NC_001437.1 -ACACAGGAUCU--- 258
MBFV-NC_007580.2 -UCACAGGAUCU--- 258
MBFV-NC_000943.1 C--CACAGGAUCU--- 258
MBFV-NC_006551.1 C--CACAGGAUCU--- 258
MBFV-NC_012534.1 -ACACAGUGUCU--- 258
MBFV-NC_015843.2 -ACACAGGAUCU--- 258
MBFV-NC_009026.2 C---CAGAUUCU--- 258
dISFV-NC_017086.1 CAACAAGUAUCU--- 258
MBFV-NC_012532.1 CCAUGGUU--UCU--- 258
MBFV-NC_035889.1 CCAUGGGU--UCU--- 258
MBFV-NC_012533.1 CCAACAGG--ACU--- 258
MBFV-NC_002640.1 CAACAGGU--UCU--- 258
MBFV-NC_001474.2 CAACAGGU--UCU--- 258
MBFV-NC_001477.1 CAACAGGU--UCU--- 258
MBFV-NC_001475.2 CAACAGGU--UCU--- 258
dISFV-NC_027999.1 CAACCAGUUUCU--- 258
MBFV-NC_008719.1 CAACUAGUGUCU--- 258
MBFV-NC_012735.1 CAACUAGUGUCU--- 258
MBFV-NC_002031.1 CACAAAAGACUCU--- 258
dISFV-NC_040610.1 UAU--GUAAUUAUUGG 258
dISFV-NC_024017.1 AAUAACAUUAUCU--- 258
dISFV-NC_016997.1 UA---AGUAUCU--- 258
MBFV-NC_018705.3 AGGAGAGCCUCU--GU 258
00.....310...

