

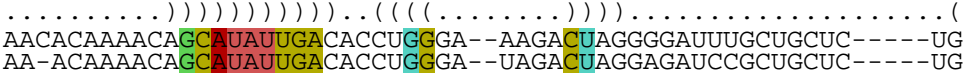
MBFV-NC_009029.2 UGAUUAACAUAACACAGUGUGAACAAAGUUUUUUGAACAG-----C---AAUGACC-- 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAAGUUUUUUGAUCAGUGG-----AU---AAUGACU-- 46
MBFV-NC_009028.2 U--U-----UAACACAGUCUGAAU--AGUUUUUU--AGCAAGGGA-UUUCC---CAUGUCU-- 46
MBFV-NC_040776.1 U--U-----UAACACAGUCUGAAU--AGUUUUUU--AGCAGAGGU-ACUAC---CAUGUCU-- 46
MBFV-NC_034151.1 -----UAACACAGUGUGAAU--AGUUUUUUUGAACAGAGUGAAAC---CAUGUCU-- 46
MBFV-NC_001563.2 A--U-----UAACACAGUGCGAGC--UGUUUUUUGGCACGAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_009942.1 A--U-----UAACACAGUGCGAGC--UGUUUUUUGAAGCAGGAAGA--UCUC---GAUGUCU-- 46
MBFV-NC_001437.1 U--U-----AGUGCAGUUUAAACA--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--GAAUAGGUUUUUAUUGGAAUACGAAAUACCG-----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-----CGUGUUGAGAAAGAAACAGAAUUCUCUGG--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 GGAUUAUU--UGUUGUUUGAG-----AGAACAGCGAGAAGUACUACU---AUGGUUGA 48
dISFV-NC_027999.1 AUUUCUACUGAAAGGGU---AG-----AGA--A--AAGGAGUUUGCUUCUCAUGGC-- 45
MBFV-NC_008719.1 -UUUCUACUG--AAAGGA---GU-----AGA--AGAAAGGAGAUUCAUCCCAAUGGC--- 45
MBFV-NC_012735.1 UAAUCGUUUCG-----UUGAG-----CGAUUAGCAGAGAACUGACCAGAACAUG-- 43
MBFV-NC_002031.1 --GAACAAUUUUUAAAGAAGACAAACUAAGAAAUUU--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-AACCAGGAAGGCGCGG--UAGAAUUGCGCGGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_032088.1 -A-----AAA-AACCAGGAAGAGCGG--CAAUCGUCGACUGUCAUAUAGUCUAAAGCGC 97
MBFV-NC_009028.2 -A-----AAA-AACCAGGAAGAGCGG--UGCAAAGCGACUGUCAUAUAGCUAAAACGC 97
MBFV-NC_040776.1 -A-----AAA-AACCAGGAGGAGCGG--CGGAAGGCGGUUGUCAUAUAGCUGAAACGC 97
MBFV-NC_034151.1 -A-----AAA-AACCAGGAGGAGCGG--AAAGAACCGGUCGUUAUAUAGCUGAAGCGC 97
MBFV-NC_001563.2 -A-----AGA-AACCAGGAGGCGCGG--UAAAAACCGGCGUCUCAUAUAGCUAAAACGC 97
MBFV-NC_009942.1 -A-----AGA-AACCAGGAGGCGCGG--CAAGAGCGCGGUCUCAUAUAGCUAAAACGC 97
MBFV-NC_001437.1 -A-----AAA-AACCAGGAGGCGCGG--UAAAAACCGGCUAUAUAUAGCUGAAGCGC 97
MBFV-NC_007580.2 -A-----AAA-AACCAGGAAGACCGG--UAGAAACCGGUUGUCAUAUAGCUAAAACGC 97
MBFV-NC_000943.1 -A-----AAA-AACCAGGAGGAGCGG--GAAGCCCAGGUCGUCAUAUAGCUAAAACGC 97
MBFV-NC_006551.1 -A-----AGA-AACCAGGAGGCGCGG--AAGAAACCGGCCAUCAUAUAGCUGAAGCGC 97
MBFV-NC_012534.1 -AA--CAAAA-AACCAGGAAGACCGG--GGCAACCAGGUUGUCAUAUAGCUAAAGCGC 100
MBFV-NC_015843.2 -AA--CAAAA-AACCAGGAAGACCGG--CUCAGGCAGGUUGUCAUAUAGCUAAAGCGC 100
MBFV-NC_009026.2 -A-----GAA-AACCAGGAAGACCGG--AGGAACACGGGUCGUCAUAUAGCUAAAGCGC 97
dISFV-NC_017086.1 -AA--UAAACCCAA---GAAGCGCGG--AAGA---CGGCUAUCGAUAUAGUGAGACGU 94
MBFV-NC_012532.1 -AA--AAACCCCAAGAAGAAUUCGG--AGGA-UCGGAUUGUCAUAUAGCUAAAACGC 97
MBFV-NC_035889.1 -AA--AAA-CCCCAAAAAGAAUUCGG--AGGAUUCGGAUUGUCAUAUAGCUAAAACGC 97
MBFV-NC_012533.1 -AA--AA--CCCCAAAA--AGGGCGC---AUCGCGCCUAUAUAUAGCUGAAGCGC 91
MBFV-NC_002640.1 -----AA--CCAACGAAAAAAGGUGG---UUAGACCAUU--UCAUAUAGCUGAAACGC 91
MBFV-NC_001474.2 -AA--UAA--CCAACGAAAAAAGCGA---AAAACAACGCUU--UCAUAUAGCUGAAACGC 94
MBFV-NC_001477.1 -AA--CAA--CCAACGAAAAAAGACCG--GUCGACGCUU--UCAUAUAGCUGAAACGC 94
MBFV-NC_001475.2 -AA--CAA--CCAACGGAAGAAGACGG--GAAAACCGJCUA--UCAUAUAGCUGAAACGC 94
dISFV-NC_027999.1 GAA--AACUAGAGUAGG---AUCUCGGGUUAAAACCCGAGGUGUCAUAUAGU---UA 98
MBFV-NC_008719.1 AAC--AAGAGGAGCAAGCAAGUCUGGGUCACGACCAGGGCGUCAUAUAGGU---GA 98
MBFV-NC_012735.1 AAC--AAAGGGGAUGAAUAAGUCUGGGCUCGAUCCAGGGCGUCAUAUAGGU---AG 98
MBFV-NC_002031.1 -----UCUGGUCGUAAAGCAGGGGAAAAACCCUGGCGUCAUAUAGGUACGACGA 94
dISFV-NC_040610.1 -----AGACAACUUGGAAAGACCGU--UGCAAGACCGUCUGUCAUAUAGGAAAGCGG 97
dISFV-NC_024017.1 -----ACUAAAUCAAGAAAGCGCGC--CGCAAAGCGGCGUUAUAUAGCUAACACGC 97
dISFV-NC_016997.1 AAUGGUCUUUCGAGCUAGGAGGCGCGU---UAAACGGCCGCGCAUAUAGUAAAACGU 103
MBFV-NC_018705.3 ----AACAAAAAACAGGAAGACCGG--GCAACCAGGUUGUCAUAUAGCUAAAGCGC 100
0.....70.....80.....90.....100.....110.....1



MBFV-NC_009029.2 GGUGCGUCCCGCG-----NNNNNNNN--AGAGGAGA---CCCUCCCACAAAGAAGCGCA 145
MBFV-NC_032088.1 GGAGCAUCCCGCG-----NNNNNNNN--AGAGGUUAGAGGAGACCCCCACAU--ACCGCA 147
MBFV-NC_009028.2 CUGGCUAGCGGUGU-----NNNNNNNN--GUGGAGA---CCCUCAGCAC--CAAGCGC 142
MBFV-NC_040776.1 CCAGCCAGCGGUU-----NNNNNNNN--A--GGAGA---CCCUCUGGCACACAAGCGC 142
MBFV-NC_034151.1 ACAUCCCGCGCUA-----NNNNNNNN--AGAGGAGA---AGAGGAGACCCGAGUGAA--CGC 138
MBFV-NC_001563.2 GGUAGUCCCCCGCG-----NNNNNNNN--GAGGAGA---CCCCGUGCCA--AAAACACC 142
MBFV-NC_009942.1 GGAAUCCCCCGCG-----NNNNNNNN--GGAGA---CCCCGUGCCACAAAACACC 141
MBFV-NC_001437.1 GGCCUACCCCGCG-----NNNNNNNN--UAGAGGAGA---CCCGCAUUUG----- 137
MBFV-NC_007580.2 GGCUGAUCCCGCG-----NNNNNNNN--UUAGAGGAGA---CCCUCUGCCGUUAACG--- 143
MBFV-NC_000943.1 GGCAUACCCCGCG-----NNNNNNNN--GA---CCCCACUCUCAAAAGCAUC 138
MBFV-NC_006551.1 GGCAUACCCCGCG-----NNNNNNNN--A---CCCCGCA--UCAUAAGCAUC 136
MBFV-NC_012534.1 GGUAGUCCUCC-----NNNNNNNN--GGAGA---CCCUCUGCGAUUAACACC 141
MBFV-NC_015843.2 GGAACGUCUCC-----NNNNNNNN--AGGAGA---CCCUCUGCGAGUGAACACC 142
MBFV-NC_009026.2 ACGGCGGCCAACG-----NNNNNNNN--C---CCCUU--GGCAAAACAGUUAACGCAC 142
dISFV-NC_017086.1 CGGUUGCCUCGCGUCU-----NNNNNNNN--AGAGGAGACCCCCCAAAAU--- 138
MBFV-NC_012532.1 GGAGUAGCCCGUG-----NNNNNNNN--GGUAGAGGAGACCCCCCGGAAAAC--- 142
MBFV-NC_035889.1 GGAGUAGCCCGUG-----NNNNNNNN--GUUAGAGGAGACCCCCCGGAAAAC--- 141
MBFV-NC_012533.1 UUUUGAUUAGCGCGUGGCCNNNNNNNN-----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 GGCUGUACUGGA---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 AUUAGUCCCGCU-----NNNNNNNN-----ACCCUCCAGUUAUU--- 131
dISFV-NC_024017.1 AUCACCAAGCGUG-----NNNNNNNN-----CCCCACU--GCUGAG--- 130
dISFV-NC_016997.1 AAGCGUC-----NNNNNNNN-----AGGAAACCCCCCAAAAUAGU--- 138
MBFV-NC_018705.3 GGCGCAUCC-----NNNNNNNN-----CCCCCGGAAAUCA-- 131
20.....130.....140.....150.....160.....170.....1



MBFV-NC_009029.2 AACACAAAAACGCAUAUUGACACCUUGGA--AAGAUAAGGGGAUUUGCUGCUC-----UG 198
MBFV-NC_032088.1 AA--ACAAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGAUCCGUGCUC-----UG 199
MBFV-NC_009028.2 GAA--ACAAAACAGCAUAUUGACGCCUUGGA--AAGACCGGGAGAUCCUCUGCU---U-UCC 195
MBFV-NC_040776.1 AAAGACAACACAGCAUAUUGACACCUUGGA--AAGACCGGGAGACCCUCUGCU---C-UGC 196
MBFV-NC_034151.1 AAUAUAAAAACAGCAUAUUGACACCUUGGA--UAGACUGGGAGACCCUCUGCUUA-U-UUC 194
MBFV-NC_001563.2 A--AAAGAAACAGCAUAUUGACACCUUGGA--UAGACUAGGGGAUUCUUC-UGCUC-UGCAC 197
MBFV-NC_009942.1 ACAACAAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGAUUCUUC-UGCUC-UGCAC 197
MBFV-NC_001437.1 --CAUCAACACAGCAUAUUGACACCUUGGAA-UAGACUGGGAGAUUCUUCUGCUCU-AUCUC 193
MBFV-NC_007580.2 --CAACAACACAGCAUAUUGACACCUUGGAA--AGACAG--GAGAUCCCCUGCU-UUUUC 194
MBFV-NC_000943.1 ---AAACAACACAGCAUAUUGACACCUUGGAA-AAGACUAGGAGAUUCUUCUGCUCU-AUUC 193
MBFV-NC_006551.1 ---AAAAAACACAGCAUAUUGACACCUUGGAAUAGACUAGGAGAUUCUUCUGCUCU-AUUC 193
MBFV-NC_012534.1 AAAACAAAACAGCAUAUUGACACCUUGGA--GAGACUAGGAGACCCUCUGUC---CUAAC 196
MBFV-NC_015843.2 ACAA--GAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGACCCUCUGUC---CUAAC 196
MBFV-NC_009026.2 CAAAAGAAACAGCAUAUUGACACCUUGGA--UAGACC--GGAGAUUUUGCGUCCU-CG--C 196
dISFV-NC_017086.1 ---GAAGGGCAUAUAUCGACACUUGGA--AAGACAGAGGAUUCUGCUGAUU-CACCG 192
MBFV-NC_012532.1 ---GCAAAACAGCAUAUUGACG--UGGGA--AAGACAGAGACUCCAUGAGUUU-CCACC 194
MBFV-NC_035889.1 ---GCAAAACAGCAUAUUGACG--UGGGA--AAGACAGAGACUCCAUGAGUUU-CCACC 194
MBFV-NC_012533.1 ---AUAAAACAGCAUAUUGACACCUUGGA--AAGACCGGAGACUCCUUGGCUU-UCCCC 196
MBFV-NC_002640.1 ---CAAAAACAGCAUAUUGACG--CUGGGA--AAGACAGAGAUCCUG---CUGU-CUCUG 196
MBFV-NC_001474.2 ---AAAAACAGCAUAUUGACG--CUGGGA--AAGACAGAGAUCCUG---CUGU-CUCCU 196
MBFV-NC_001477.1 ---AACAAACAGCAUAUUGACG--CUGGGA--GAGACAGAGAUCCUG---CUGU-CUCUA 196
MBFV-NC_001475.2 ---UAAAACAGCAUAUUGACG--CUGGGA--GAGACAGAGAUCCUG---CUGU-CUCCU 196
dISFV-NC_027999.1 ---UAGC-U--UCAUAUUGACACCAAGGAA-AAGACCGGAGAAACUUCUUGCUUACUCGC 191
MBFV-NC_008719.1 --AUAGC--GGACCAUAUUGACGCCUUGG-A-AAGACCGGAGACACUCCUUGAUU-CUCAC 192
MBFV-NC_012735.1 --AUAGC--GGACCAUAUUGACGCCAGGG-A-AAGACCGGAGACACUCCUUGAUU-CUCAC 192
MBFV-NC_002031.1 --AUAGUGGGA--CAUAUUGACGCCAGGG-A-AAGACCGGAGUGGUUCUCUGCUUUUCCUC 192
dISFV-NC_040610.1 --UAUAUAACAGCAUAUUGACACCAAGG-A-UAGACCGGAGAGAAU--ACACAUUUUCUGG 185
dISFV-NC_024017.1 --AGCAAAAUAGCAUAUUGACACUUGGGAA-GAGACAGAGAUUA-UGAUGUUGUUUGUCA 186
dISFV-NC_016997.1 --AUAGGCUUAUAUUGACACCAUAGGA--AAGACUGGAGAUACCUUUGGUUACUGGAG 193
MBFV-NC_018705.3 ---AAGCAAGGCAUAUUGACCAAGGUGUAUACCAAGUGGAGAUACUAGGUUUCAGGC 189
80.....190.....200.....210.....220.....230.....2



MBFV-NC_009029.2 (((247
MBFV-NC_032088.1 GACUUCGCGUUCUC---GGCACAGAACCGG---UUGA--GGACCGGAGGCCCAAA--- 247
MBFV-NC_009028.2 GAUCCCGAGUUCUA---GGCACAGAACCGCG---UAA--UGAACUGGGGGGCCAAA--- 247
MBFV-NC_040776.1 AUCACCGCGACUA---GGCACAGAUCGCG---CAAGUAGUGGUGUGUGGUGAAACA 249
MBFV-NC_034151.1 AACACCGCGACUA---GGCACAGAUCGCG---AAAUUCUGGUGUGUGUGGUGGACAAA 250
MBFV-NC_001563.2 AACACCGCGACUC---GGCACAGAACCGCG---UAAGU--GUGCGUGGUGUGAA--AAUA 246
MBFV-NC_009942.1 A--ACGACCGACAC---GGCACAGUGCGCG---ACAUAAGUGGUGUGUG--GUGCUAGA 247
MBFV-NC_001437.1 A--ACGACCGACAC---GGCACAGUGCGCG---ACAUAAGUGGUGUGUG--GUGCGAGA 247
MBFV-NC_007580.2 AACAUACGCUACUA---GGCACAGAGCGCG---AAGUAUGUAGUGUGUGGUGGAGAAGA 247
MBFV-NC_000943.1 AACACCGCGACAA---GGCACAGAGCGCG---UAAAGUGUGGUGUGUGGUGA---A 247
MBFV-NC_006551.1 AACAUACGCUACAA---GGCACAGAGCGCG---AACAUUGUAGUGUGUGGUGGAGAAGA 247
MBFV-NC_012534.1 AACACCGCGAUCC---GGCACAGAGCGCG---UAAG--UGUGGUGUGUGGUGUAGA--- 247
MBFV-NC_015843.2 AACACCGCGACUU---GGCACAGAGCGCG---AAAG--UGUGGUGUGUGGUGUAGA--- 247
MBFV-NC_009026.2 AACACUCCCAACU---GGCACAGAACCGCG---ACA--UGUGGAGGAGGUGGUAAGACA 249
dISFV-NC_017086.1 CCACCGAGCUACAC---GGCACAGCGCGCG---GAAAAGGUAGUUUGGAGGUUGUAAAA 246
MBFV-NC_012532.1 ACGCUGCGCGCCA---GGCACAGAUCGCG---AACUUCGCGCGCGUGUGUGGGGAAAU 247
MBFV-NC_035889.1 ACGCUGCGCGCCA---GGCACAGAUCGCG---AAUAGCGCGCGCGUGUGUGGGGAAAU 247
MBFV-NC_012533.1 ACACCGCGCGCCU---GGCACAGAGCGCG---AGAUCGCGCGUGUGUGGAGCAAC 247
MBFV-NC_002640.1 CAACAUCAAUCCA---GGCACAGAGCGCG---CAAGAUGGAUUGGUGUUGUUG--AUC 247
MBFV-NC_001474.2 CAGCAUCAUCCA---GGCACAGAACCGCA---GAAAUGGAUUGGUGUGUUG--AAU 247
MBFV-NC_001477.1 CAGCAUCAUCCA---GGCACAGAACCGCA---AAAAUGGAUUGGUGUGUUG--AAU 247
MBFV-NC_001475.2 CAGCAUCAUCCA---GGCACAGAACCGCA---GAAAUGGAUUGGUGUGUUG--AAU 247
dISFV-NC_027999.1 AGCAACACCGCAAA---GGCACAGUAGCGU---GAAAUAUUGGUGUGUAGUAGUAAA 246
MBFV-NC_008719.1 CUUUCUACCCUUA---AGCACAGAUUGCU---G-AAUGCAGGUGGGGAAGUUGGGAAC 246
MBFV-NC_012735.1 CUUUCUUGCCCUAA---AGCACAGAUUGCU---G-AAUGCAGGCGGGGAAGUUGAGAAC 246
MBFV-NC_002031.1 C--AGAGGUGUGU---AGCACAGUUGCU---CAAGAAUAGCAGACCUUUG--GAUGACAAA 246
dISFV-NC_040610.1 GCACAAACAGUACA---AGCACAGAGCGCU---GAGAAGUAGUAGUUGUUGAGAGAGAA 244
dISFV-NC_024017.1 AUGCAACAAACAGCCAU---GGCACAGAGCGCG---UGAAGAGAGCGUUGUUAUAUUUACAAA 246
dISFV-NC_016997.1 ACUAACUCCUAU---GGCACAGAGCGCG---AAAGAACAACGAGUUAAGUUGGAAAAAGAC 249
MBFV-NC_018705.3 ACCCCUUGCGAG---UGAACACCACAA---AAAACAGCAUAUAGCCUUGGGAAAGACU 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 CAUUGGUU--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 CACAAAACUCU--- 258
dISFV-NC_040610.1 UAU--GUAAUAUCUUGG 258
dISFV-NC_024017.1 AAUAACAUUAUCU--- 258
dISFV-NC_016997.1 UA---AGUAUCU--- 258
MBFV-NC_018705.3 AGGAGAGCCUCU--GU 258
00.....310...

