

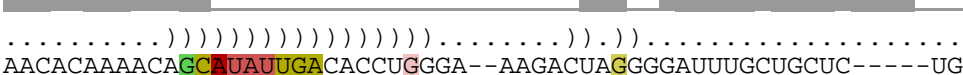
MBFV-NC_009029.2 UGAUUAACAUAACACAGUGUGAACAAAGUUUUUUGAACAG-----C---AAUGACC-- 46
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAAGUUUUUUGAUCAGUGG-----AU---AAUGACU-- 46
MBFV-NC_009028.2 U--U-----UAACACAGUCUGUAAU--AGUUUUUU--AGCAAGGGA--UUUCC---CAUGUCU-- 46
MBFV-NC_040776.1 U--U-----UAACACAGUCUGUAAU--AGUUUUUU--AGCAGAGGU--ACUAC---CAUGUCU-- 46
MBFV-NC_034151.1 -----UAACACAGUGUGAAU--AGUUUUUUUGAACAGAGGUGAAAC---CAUGUCU-- 46
MBFV-NC_001563.2 A--U-----UAACACAGUGCGGAGC--UGUUUUUUGGCACGAAGA--UCUC---CAUGUCU-- 46
MBFV-NC_009942.1 A--U-----UAACACAGUGCGGAGC--UGUUUUUUGAAGCACGAAGA--UCUC---CAUGUCU-- 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-----UAACACAGUGCCGGC--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-----CGUGUUGGAAAGAAACAGAAUUCUCUGG--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_001475.2 ACGUAGUGCUG--ACAGUUU--UUU--AUUAGAGAGCAG--AUCUCUG-----AUG----- 43
dISFV-NC_027999.1 GGAUUAAU--UGUUGUUUGAG-----AGAACAGCGAGAAGUACUACU---AUGGUUGA 48
AUUUCUACUGAAAGGGU--AG-----AGA--A--AAGGAGUUUUGCUUCUCAUGGC-- 45
MBFV-NC_008719.1 -UUUCUACUG--AAAGGA--GU-----AGA--AGAAAGGAGAUUCAUCCCAAUGGC-- 45
MBFV-NC_002031.1 UAAUCGUUUC-----UUGAG-----CGAUUAGCAGAGAACUGACCAGAACAUG-- 43
dISFV-NC_040610.1 --GAACAAUUUUUAAAGAAGACAAACUAAGAAAUU--AAUUUC-----AAUGGUU-- 46
dISFV-NC_024017.1 UCGCGCAUUUUUGAUAGAGUU--GACUAGAUAAACGAUUUAUC-----GAUGGUG-- 46
dISFV-NC_016997.1 AAACA--UAACGCAUUUGAAUAGGUUUUGAAAGAGAUUGAAUC-----AAUGAGAUC 48
MBFV-NC_018705.3 CCGUA-----AACACAGUUUGAAGGUUUUUGGAGAGAGACUACU-----AUGUCA-- 46
.....10.....20.....30.....40.....50.....6



.....((((.....((((((((((((((((.....(((
MBFV-NC_009029.2 -A-----AAA-AACCAGGAAGGCCGG--UAGAAAUCCGGGCGGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_032088.1 -A-----AAA-AACCAGGAAGGUCGG--CAAUCGUCGGACUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_009028.2 -A-----AAA-AACCAGGAAGGUCGC--UGCAAAGCGGACUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_040776.1 -A-----AAA-AACCAGGAGGACCGG--CGGAAGCGGGGUUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_034151.1 -A-----AAA-AACCAGGAGGACCGG--AAAGAACCGGGUCGUAAUAUUGCUAAGCGC 97
MBFV-NC_001563.2 -A-----AGA-AACCAGGAGGGGCCGG--UAAAAACCGGGCUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_009942.1 -A-----AGA-AACCAGGAGGGGCCGG--CAAGAGCCGGGCGUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_001437.1 -A-----AAA-AACCAGGAGGGGCCGG--UAAAAACCGGGCUGUCAAUAUUGCUAAGCGC 97
MBFV-NC_007580.2 -A-----AAA-AACCAGGAAGAACCCGG--UAGAAACCGGGUUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_000943.1 -A-----AAA-AACCAGGAGGACCGG--GAAGCCCCGGGUGUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_006551.1 -A-----AGA-AACCAGGAGGGGCCGG--AAGAAACCGGGCUGUCAAUAUUGCUAAGCGC 97
MBFV-NC_012534.1 -AA--CAAAA-AACCAGGAAGAACCGG--GGCAACCCGGGUUGUCAAUAUUGCUAAAGCGC 100
MBFV-NC_015843.2 -AA--CAAAA-AACCAGGAAGAACCGG--CUCAGGCCGGGUGUGUCAAUAUUGCUAAAGCGC 100
MBFV-NC_009026.2 -A-----GAA-AACCAGGAAGACCGG--AGGAACCCGGGUCUGUCAAUAUUGCUAAGCGC 97
dISFV-NC_017086.1 -AA--UAAACCCAA--GAAGCCCGG--AAGA--CGGGCUAUGAUUAGUAGAGCU 94
MBFV-NC_012532.1 -AA--AAACCCCAAGAAGAAAUCCGG--AGGA-UCCGGAUUUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_035889.1 -AA--AAA-CCCCAAAAAGAAAUCCGG--AGGAUUCGGAUUUGUCAAUAUUGCUAAAGCGC 97
MBFV-NC_012533.1 -AA--AA--CCCCAAAA--AGGGCCGC--AUCGCGGGCUAUGAUUAGUAGAGCGC 91
MBFV-NC_002640.1 -----AA--CCAACGAAAAAAGGUGG--UUAGACCACCUU--UCAAUAUUGCUGAAACGC 91
MBFV-NC_001474.2 -AA--UAA--CCAACGAAAAAAGGCGA--AAAACACGCCUU--UCAAUAUUGCUGAAACGC 94
MBFV-NC_001477.1 -AA--CAA--CCAACGAAAAAAGACGG--GUCGACGCUU--UCAAUAUUGCUGAAACGC 94
MBFV-NC_001475.2 -AA--CAA--CCAACGGAAGAAGACGG--GAAAACCGUCUA--UCAAUAUUGCUGAAACGC 94
dISFV-NC_027999.1 GAA--AACUAGAGUAGG--AUCUCGGGUUAAAACCCGAGGUGUCAAUAUUGAU--UA 98
AAC--AAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUUGU--GA 98
AAC--AAAGGGGAUGAAUAAGUCUCGGGUCGAUCCCGAGGCGUCAAUAUUGU--AG 98
-----UCUGGUCGUAAAGUCUAGGGAAAAACCCUGGGCGUCAAUAUUGGUACGACGA 94
dISFV-NC_040610.1 -----AGACAACUUGGAAGACCGU--UGCAAGACGGUCUGUCAAUAUUGGAAAGCGC 97
dISFV-NC_024017.1 -----ACUAAAUCAAGAAAGCCGC--CGCAAAGCGGGCUGUAAUAUUGCUAACACGC 97
dISFV-NC_016997.1 AAUGGUCUUUCGAGCUAGGAGGCCCGU--UAAACGGGGCGGUGCAUAUUGAUAAACGC 103
MBFV-NC_018705.3 ----AACAAAAAACAGGAAGACCGGG--GCAACCCGGGUUGUCAAUAUUGCUAAAGCGC 100
0.....70.....80.....90.....100.....110.....1



((.....))))......((.....)).....
MBFV-NC_009029.2 GGUGCGUCCCGCG-----NNNNNNNN--AGAGGAGA--CCCUCCACAAAGAAGCGCA 145
MBFV-NC_032088.1 GGAGCAUCCCGCG-----NNNNNNNN--AGAGGUUAGAGGAGACCCCCACAU--ACCGCA 147
MBFV-NC_009028.2 CUGGCUAGCGUGU-----NNNNNNNN--GUGGAGA--CCCUCAGCAC--CAAGCGC 142
MBFV-NC_040776.1 CCAGCCCGCGUUU-----NNNNNNNN--A--GGAGA--CCCUCUGGCACACAAGCGC 142
MBFV-NC_034151.1 ACAUCCCGCGCUA-----NNNNNNNN--AGAGGAGA--CCCUCUGGCACAGUAA--CGC 138
MBFV-NC_001563.2 GGUUAGCCCCCGCG-----NNNNNNNN--GAGGAGA--CCCUGUGCCA--AAAACACC 142
MBFV-NC_009942.1 GGAAUCCCGCGCG-----NNNNNNNN--GGAGA--CCCUGUGCCACAAACACC 141
MBFV-NC_001437.1 GGCCUACCCCGCG-----NNNNNNNN--UAGAGGAGA--CCCUGCAUUUG-----137
MBFV-NC_007580.2 GGCUGAUCCCGCG-----NNNNNNNN--UUAGAGGAGA--CCCUCUGCCGUUAACG---143
MBFV-NC_000943.1 GGCAUACCCCGCG-----NNNNNNNN--GA--CCCACUCUCAAAAGCAUC 138
MBFV-NC_006551.1 GGCAUACCCCGCG-----NNNNNNNN--A--CCCUGCA--UCAUAAGCAUC 136
MBFV-NC_012534.1 GGUUAGCUCC-----NNNNNNNN--GGAGA--CCCUCUGCGAGUUAACACC 141
MBFV-NC_015843.2 GGAACGCUCC-----NNNNNNNN--AGGAGA--CCCUCUGCGAGUGAACACC 142
MBFV-NC_009026.2 ACGGCGGCCAACG-----NNNNNNNN--C-----CCCUU--GGCAAAACAGUUAACGCAC 142
dISFV-NC_017086.1 GCGUUGCCUCGCGUCU-----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--CUAGAGGUUAGAGGAGACCCCC--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-----ACCCUCCAGUUAUU---131
dISFV-NC_024017.1 AUCACCAAGCGUG-----NNNNNNNN-----CCCCACU--GCUGAG---130
dISFV-NC_016997.1 AAGCUGC-----NNNNNNNN-----AGGAAACCCCCCAAAAUAGU---138
MBFV-NC_018705.3 GCGCAUCC-----NNNNNNNN-----CCCCCGGGAUAUCA--131
20.....130.....140.....150.....160.....170.....1



.....))))))))......)).....
MBFV-NC_009029.2 AACACAAAACGCAUAUUGACACCUUGGA--AAGACUAGGGGAUUUUGCUGCUC-----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 AAAGACAACACGCAUAUUGACACCUUGGA--AAGACCGGAGACCCUCUGCUC--C--UGC 196
MBFV-NC_034151.1 AAUAUAAAACAGCAUAUUGACACCUUGGA--UAGACUGGAGACCCUCUGCUCU--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--UAGACUGGAGAGAUUCUUGCUCU--AUCUC 193
MBFV-NC_007580.2 --CAACAACACAGCAUAUUGACACCUUGGAA--AGACAG--GAGAUCCCCUGCUCU--UUUUC 194
MBFV-NC_000943.1 ---AAACAACACAGCAUAUUGACACCUUGGAA--AAGACUAGGAGAUUCUUGCUCU--AUUCC 193
MBFV-NC_006551.1 ---AAAAAACAGCAUAUUGACACCUUGGAAUAGACUAGGAGAUUCUUGCUCU--AUUCC 193
MBFV-NC_012534.1 AAAACAAAACAGCAUAUUGACACCUUGGA--GAGACUAGGAGACCCUCUGCUC--CUAAC 196
MBFV-NC_015843.2 ACAA--GAAACAGCAUAUUGACACCUUGGA--UAGACUAGGAGACCCUCUGCUC--CUAAC 196
MBFV-NC_009026.2 CAAAAGAAACAGCAUAUUGACACCUUGGA--UAGACC--GGAGAAUUGCGUCCU--CG--C 196
dISFV-NC_017086.1 ---GAAGGGCAUAUAUGCAGCUCUUGGA--AAGACCAAGGUAUCUCGUGAUU--CACCG 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 ---UAAAACAGCAUAUUGACG--CUGGGA--GAGACCAAGAUCCUG--CUGU--CUCCU 196
dISFV-NC_027999.1 ---UAGC--U--UCAUAUUGACACCAAGGAA--AAGACCGAGAAACUUCUUGCUUACUCGC 191
MBFV-NC_008719.1 --AUAGC--GGACCAUAUUGACGCGUUGG--A--AAGACCGAGACACUCCUUGAUU--CUCAC 192
MBFV-NC_012735.1 --AUAGC--GGACCAUAUUGACGCGAGGG--A--AAGACCGAGACACUCCUUGAUU--CUCAC 192
MBFV-NC_002031.1 --AUAGUGGGA--CAUAUUGACGCCAGGG--A--AAGACCGAGUGGUUCUUCGCUUUCUC 192
dISFV-NC_040610.1 --UAUAUAACAGCAUAUUGACACCAAGG--A--UAGACCGAGACAAUUCUUGGUGG 185
dISFV-NC_024017.1 --AGCAAAUAGCAUAUUGACACUUGGGA--GAGACCAAGAUU--UGAUGUUGUUUGUCA 186
dISFV-NC_016997.1 --AUAGGCUUAUAUUGACACCAUGGA--AAGACCGAGAUACCUUUGGUUUCGAG 193
MBFV-NC_018705.3 --AAGCAAGGCAUAUUGACCAACUGGUAUAACAGGGGAGAUACUAGGUUUGGAG 189
80.....190.....200.....210.....220.....230.....2



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MBFV-NC_009029.2 GACUUCGCGCUCUC--GGCACAGAACCGG--UUGA--GGACCGGAGGCCCAAA--- 247
MBFV-NC_032088.1 GAACCCGAGUCUUA--GGCACAGAACCGG--UUA--UGAACUGGGUGGCCAAA--- 247
MBFV-NC_009028.2 AUCACCAAGCACUA--GGCACAGAUCCCG--CAAGUAGUGGUGGUGGUGAAACA 249
MBFV-NC_040776.1 AACACCAAGCACUA--GGCACAGAUCCCG--AAAUUCUGGUGGUGGUGGAGAAACA 250
MBFV-NC_034151.1 AACACCAAGCACUC--GGCACAGAACCGG--UAAGU--GUGCGUGGUGGUGAA--AAUA 246
MBFV-NC_001563.2 A--ACCAAGCACAC--GGCACAGUCCCGG--ACAUAAGUGGUGGUG--GUGCUAGA 247
MBFV-NC_009942.1 A--ACCAAGCACAC--GGCACAGUCCCGG--ACAUAAGUGGUGGUG--GUGCGAGA 247
MBFV-NC_001437.1 AACAUACGCUACUA--GGCACAGAGCCCG--AAGUAUGUAGUGGUGGUGGAGGA 247
MBFV-NC_007580.2 AACACCAAGCACAA--GGCACAGAGCCCG--UAAAGUGUGGUGGUGGUGAAAA--A 247
MBFV-NC_000943.1 AACAUACGUCACAA--GGCACAGAGCCCG--AACACUGUAGACUAGUGGGGAGAGA 247
MBFV-NC_006551.1 AACAUACGUCACAA--GGCACAGAGCCCG--AAAAUUGGUGGUGGUGGAGACUA 247
MBFV-NC_012534.1 AACACCAAGCAUCC--GGCACAGAGCCCG--UAAG--UGUGGUGGUGGUGUAGA-- 247
MBFV-NC_015843.2 AACACCAAGCACUU--GGCACAGAGCCCG--AAAG--UGUGGUGGUGGUGUAGA-- 247
MBFV-NC_009026.2 AACACCUCCACCC--GGCACAGAACCGG--ACA--UGUGGAGGGGUGGUAAGACA 249
dISFV-NC_017086.1 CCACCAAGACUACAC--GGCACAGCGCCCG--GAAAAGGUAGUUGGAGGUUGUAAAA 246
MBFV-NC_012532.1 ACGCUGGCGGCCA--GGCACAGAUCCCG--AACUUCGCGCGCGUGUGGGGAAAU 247
MBFV-NC_035889.1 ACACCAAGCCGCA--GGCACAGAUCCCG--AAUAGCGCGCGCGUGUGGGGAAAU 247
MBFV-NC_012533.1 ACACCAAGCCCU--GGCACAGAGCCCG--AGAUCGCGCGUGGUGGUGGAGCAAC 247
MBFV-NC_002640.1 CAACAUAUAUCCA--GGCACAGAGCCCG--CAAGAUGGAUUGGUGUUGUUG--AUC 247
MBFV-NC_001474.2 CAGCAUAUAUCCA--GGCACAGAACCGA--GAAAAGGAUUGGUGUGUUG--AAU 247
MBFV-NC_001477.1 CAGCAUAUAUCCA--GGCACAGAACCGA--AAAAUAGGAUUGGUGUGUUG--AAU 247
MBFV-NC_001475.2 CAGCAUAUAUCCA--GGCACAGAACCGA--GAAAAGGAUUGGUGUGUUG--AAU 247
dISFV-NC_027999.1 AGCAACCAAGCAAA--GGCACAGUAGCCU--GAAACAUUGGUGGUGUAGUAGUAAA 246
MBFV-NC_008719.1 CUUUCUACCCUUA--AGCACAGAUUGCU--G--AAUGCAGGUGGGGAAGUUGGGAAC 246
MBFV-NC_012735.1 CUUUCUUGCCCUAA--AGCACAGAUUGCU--G--AAUGCAGGCGGGGAAGUUGAGAAC 246
MBFV-NC_002031.1 C--AGAGCUUCUGU--AGCACAGUUGCU--CAAGAUAAGACAGACUUCUUG--GAUGACAAA 246
dISFV-NC_040610.1 GCACUAUAUAUACA--AGCACAGUUGCU--UAGAGAAGUAGUAGUUGACGAGAGAA 244
dISFV-NC_024017.1 AUGCGACAACAGCCAU--GGCACAGAGCCCGUAGAGAGAGCGUUGUUAUAAUUUACAAAA 246
dISFV-NC_016997.1 ACUAACUCCUAU--GGCACAGAGCCCGAUAAGAACCGAGUUGAUGGAAAAAGACU 249
MBFV-NC_018705.3 ACCCCUUGCGAG--UGAACACCACAA--AAAACAGCAUAUUGACUUGGGAAGAAACU 244
40.....250.....260.....270.....280.....290.....3



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MBFV-NC_009029.2 -ACACCAGAUUC--- 258
MBFV-NC_032088.1 -ACACCAGAUUC--- 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 UAG--GUGUU--UCU--- 258
MBFV-NC_035889.1 CCAUGGGU--CUU--- 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 CACAAAACACUCU--- 258
dISFV-NC_040610.1 UAU--GUAAUAUCUUG 258
dISFV-NC_024017.1 AAUAACAUUAUCU--- 258
dISFV-NC_016997.1 UA---AGUAUCU--- 258
MBFV-NC_018705.3 AGGAGAGCCUCU--GU 258
00.....310...

