

NKV-NC_003635.1 --GUGCGCG--UCACGCCACUUUUUGGCGGGAAUGUGCGAAAAUAACAGGAAAAAG 54
NKV-NC_004119.1 --CCGGUGGAAACCCACGCGUCUCUUCGU---CAAUUGACGAAAGGCCAAAAUGAAAAAC 54
ISFV-NC_027819.1 --UCACGGA--AUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAGCACUGAAG 54
ISFV-NC_008604.2 --UUUUUUU--UAAUAGUCGUUUUUUCUCGCAGAAUUGGGAAAGGACGACGGUAAAGAG 54
ISFV-NC_021069.1 --CCUUAAU--ACCUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA--GUCGGGAG 52
ISFV-NC_012671.1 --CUUUAAU--ACCUCUGGUACUAAACGUGGCGCUAUGAAGAAGAAGGA--GUCUGGAG 52
ISFV-NC_024299.2 --UUGAGAA--ACUUUUUGCAGCUUUGUGAGAAUGGAGCGAAAGAAAGGAGUAAU 54
ISFV-NC_001564.2 --GAGAGAC--CGUUGAGGUAGAACCCGGCAAGGAUGAAGAGGAAGG--AUUUGUAGG 52
ISFV-NC_005064.1 --AGUUGUU--CGGAAAAGUGACCUUGGUUCGUUAUGAAUAGAGGAA--GCUUGAAGA 52
ISFV-NC_012932.1 --AGUUGUU--UGGAAAAGAGAAUAUCUGGUUCGUUAUGAAUAGGAUGC--UCUUAGAAG 52
ISFV-NC_034204.1 --GCGACAG--AAGGAACCCAAAAAGGGAACCCAGUAGGAGCAAAACG--GAAAGUUUU 52
ISFV-NC_034017.1 --UAAAAAU--UUUUUUGGCGACGUACGGAAAGUAUGGAGCAAUCGA--GGAAGUUUU 52
ISFV-NC_027817.1 --CAAAACU--UGGUGUCCAACGUUUUGUAUUAAAAUGAGUGGAUUAAGGAGCCUUCU 54
MBFV-NC_008719.1 AAGGGUA--GAGAAAAGGAGUUUUUGU-UCUCAUGGCAACAGAGGAGCAAGCAAG 54
MBFV-NC_012735.1 AAGGAGUA--GAAGAAAGGAGAUUCAUU-CCCAAUGGCAACAAAGGGGGAUGAAUAG 54
ISFV-NC_016997.1 CAUUGAA--UAGGUUUUGAAAGAGUUGA-AUCAUAGAGAUCAUUGGUCUUCUGACGU 54
MBFV-NC_002031.1 UUGAGCGA--UUAGCAGAGAACUGACCA-GAACAUUGU-----UGGUCGUAAA 45
NKV-NC_005039.1 CGGAAGAA--CAACUGUCCGGAUUCAAA-GACGAUGA--AAAAUCGUAGACAAAA 50
MBFV-NC_002640.1 -----UGUUUGAAUAGAGAGCAGAUUCUCUG-AAAAAUG--AA--CCAACGAAAAAAG- 48
MBFV-NC_001474.2 ACAGUUUUUUAAUAGAGAGCAGAUUCUG-----AUGAAUA--CCAACGAAAAAAG- 51
MBFV-NC_001477.1 AACAGUUUUUUAAUAGAGAGCAGAUUCUG-----AUGAACAA--CCAACGAAAAAAG- 51
MBFV-NC_001475.2 GACAGUUUUUUAAUAGAGAGCAGAUUCUG-----AUGAACAA--CCAACGAAAGAG- 51
MBFV-NC_012532.1 ----AUUUGGAUUUGGAAACGAGAG-UUUCU-GGUCAUGAAAAACCCAAAGAAGAAU- 53
MBFV-NC_035889.1 ----UUUUGGAUUUGGAAACGAGAG-UUUCU-GGUCAUGAAAAA-CCCAAAAAGAAU- 52
MBFV-NC_012533.1 ----GUGUUGAGAAAGAAAAACAGU-UCUUC-GGAAUAGAAA--AGAAAAAAGG-- 48
ISFV-NC_040610.1 AAAGAAGUCAAAACUAGAAAUUUAAU-UCA-----AUGUU--AGCAACUUGGAAAG 51
ISFV-NC_024017.1 UGAU-AGAGUUGACUAGAAUACGAAUUAUCG-----AUGGUG--ACUAAAUCAAGAAAG 51
MBFV-NC_001437.1 UUAACAGUUUUUUAGAACGGAAGAUAAAC-----AUGACU--AAAAAACAGGAGGG 51
MBFV-NC_012534.1 AGUUUGAACGUUUUUUGGAGAGAGAUUAU-----AUGUUUAACAAAAAACAGGAAAA 54
MBFV-NC_015843.2 AGUUUGAACGUUUUUUGGAGAGAGACAUC-----AUGUCUAAACAAAAAACAGGAAAA 54
MBFV-NC_006551.1 -GCCGGCAGUUUCUUUGAGCGUUGAUUUUCA-----AUGUCU--AAGAAACAGGAGGG 51
MBFV-NC_009029.2 -ACACAGUGUGAACAGGUUUUUUGAACAGCA-----AUGACC--AAAAAACAGGAAGG 51
MBFV-NC_001563.2 UGCAGGCUUUUCUUUGGACGAGAAUUC-UCG-----AUGUCU--AAGAAACAGGAGGG 51
MBFV-NC_009942.1 UGCAGGCUUUUCUUAGCAGAGAAUUC-UCG-----AUGUCU--AAGAAACAGGAGGG 51
MBFV-NC_032088.1 AG-----UGUGAACAAAGUUUUUGAUCAGUG-GAUAAUGACU--AAAAAACAGGAAAG 51
MBFV-NC_009028.2 UCUGAAUAGUUUUU-AGCAAGGGA-UUUC-CCAUGUCU--AAAAAACAGGAAAG 51
MBFV-NC_040776.1 UCUGAAUAGUUUUU-AGCAGAGGU-ACUA-CCAUGUCU--AAAAAACAGGAGGA 51
MBFV-NC_034151.1 --UGAAUAGUUUUUGAACAGAGGUGAAAA-CCAUGUCU--AAAAAACAGGAGGA 51
MBFV-NC_000943.1 -UGAACAGUUUUUGGAGCUUUUGAUU-UC-----AAUUGUCU--AAGAAACAGGAGGA 51
MBFV-NC_007580.2 GCGAAGCAGUUUUU--AGCAGGGAAUUAAC-CAUUGUCU--AAAAAACAGGAAAA 51
MBFV-NC_009026.2 UCGAGGAAGAGCAGAGAACGAAUUCUCUG-----AAUGGCU--AGAAAACAGGAAGA 51
ISFV-NC_017086.1 UGAAUAGGUUUAAUUGAGAAUCGAAUUAACCG-----AUGGCGA--AUAACCCAAAGAACG 52
ISFV-NC_027999.1 UUGUUUGAGAGAAACAGCGAGAAUUAUCU-----AUGGUUGAGAAAACUAGAGUAGGA 54
TBFV-NC_003687.1 -----AGCUUAGGAGAACAAAGAGCUGGGAGUGGUUUAUG-----AUGACCACUUCUAAA 48
TBFV-NC_001672.1 AAAGACAGCUUAGGAGAAACAGAGCUGGGG-----AUGGUCAAGAAGGCCAUCCUGAAA 54
TBFV-NC_005062.1 AAAGACAGCUUAGGAGAAACAGAGCUGGGG-----AUGGCCGGGAAGGCCAUUCUGAAA 54
TBFV-NC_003690.1 AAAGACAGCUUAGGAGAAACAGAGCUGGGG-----AUGGCCGGGAAGGCCAUUCUGAAA 54
TBFV-NC_001809.1 AAAGACAGCUUAGGAGAAACAGAGCUGGGG-----AUGGCCGGGAAGGCCAUUCUGAAA 54
TBFV-NC_027709.1 AAAGACAGCUUAGGAGAAACAGAGCUGGGG-----AUGGCCGGGAAGGCCAUUCUGAAA 54
.....10.....20.....30.....40.....50.....6



NKV-NC_003635.1 AACCUCUGGAAAGGGUUC--AAAGAAGAAAAAGAAUGCUGGAAGGAAU-----GNN 102
NKV-NC_004119.1 GCGAAUAAAAAGAAAGC--AAAAAACUCUAAGACCAAUCCUGGUCGA-----ANN 102
ISFV-NC_027819.1 AAGAAGGGAAUUUCCUGGGAGGAGGAGGAAACGGGGCUUUGAG-----ANN 102
ISFV-NC_008604.2 AAGAAGGGGCGGGCUCUCCUGGGAUGGUUGUUACCA--CCGGGCGC-----GNN 102
ISFV-NC_021069.1 UAGUGGAGAGAGGACGCGGUACGCGCCUCCGACCACCCGUGGAGGA-----ANN 102
ISFV-NC_012671.1 UGUUGGAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGA-----GNN 102
ISFV-NC_024299.2 UAUGGGAGAAAGGGGAAGC--UAUCCCGCCCCUCCAUUAGGAGAAA-----ANN 102
ISFV-NC_001564.2 CGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGA-----ANN 102
ISFV-NC_005064.1 AGGGGCUCCCGGGCCCAACAUCCAACCGUCCGUGCGGUUGGACCGCC-----GNN 102
ISFV-NC_012932.1 AAGGGCCUUCGGGCCAAAACAUUCGUGUCGGCCAUCCAGCAGACGCCG-----GNN 102
ISFV-NC_034204.1 CCCGACCGGGAGGUGGUCGGGCCCAGAGAGGGGCGUGGGGAAG-----ANN 102
ISFV-NC_034017.1 CCCGCGCCGGGAGCGGUCGGGCGGGCCCCAAAGAGGGGCGUGGCGAAG-----ANN 102
ISFV-NC_027817.1 CCCUCCAGGGGGAAGAA--AAAGAAAGCCCCUGUGAUACAAUCAGACAG-----GNN 102
MBFV-NC_008719.1 UCUCGGGUCACGACCCGGGGCGUCAUAUUGGUGACGGCCAAGG-----C--C-ANN 102
MBFV-NC_012735.1 UCUCGGGUCGUAUCCCGAGGCGUCAUAUUGGUAGCGGCUAGAG-----U--G-ANN 102
ISFV-NC_016997.1 AGGAGGCCCGUUAACCGGGCUGCAUAUAUAAUAAACGUAAGC-----U--G-CNN 102
MBFV-NC_002031.1 GCUCAGGGAAAAACCCUGGGCGUCAUAUUGGUACGACGAGGAGUUCGCUCCUU--G-UNN 102
NKV-NC_005039.1 CACCCG-----AAUUUCGGGUGUCAUAUAAAAAAGCAGAAAUCAAAAAACAAA-ANN 102
MBFV-NC_002640.1 -GU---G--GUUAGACCUUUAUUAUUGCUGAAACCGCGAGAGAAACCGGUAUCANN 102
MBFV-NC_001474.2 -GC---G--AAAAACAGCCUUUCAUAUAGCUGAAACGCGAGAGAAACCGCGUGU---NN 102
MBFV-NC_001477.1 -AC---G--GGUCGACCGUCUUUCAUAUAGCUGAAACGCGCGAGAAACCGCGUGU---NN 102
MBFV-NC_001475.2 -AC---G--GGAAAGCGGCUUAUCAUAUAGCUGAAACCGCGUGAGAAACCGCGUGU---NN 102
MBFV-NC_012532.1 -CC---GGAGGA-UCCGGAUUGUCAUAUAGCUAAAACGCGGAGUAGCCCGUG-----NN 102
MBFV-NC_035889.1 -CC---GGAGGAUUCGGAUUGUCAUAUAGCUAAAACGCGGAGUAGCCCGUG-----NN 102
MBFV-NC_012533.1 -----CGCGCAUCGCGGGCUAUAUAGCUGAAGCGCUUUGGAUUCAGCGUCCNN 102
ISFV-NC_040610.1 ACC---GUUGCAAGACGGUCUGUCAUAUAGCGAAAGCGGAUUACCACCCGCU-----NN 102
ISFV-NC_024017.1 CCC---GCCGCAAAAGCGGGCUGUUAUAUAGCUAACACGCAUCCACCAAGCGUG-----NN 102
MBFV-NC_001437.1 CCC---GGUAAAAACCGGGCUGUCAUAUAGCUGAAGCGGUAACCGCCUACCCGCG-----NN 102
MBFV-NC_012534.1 CCC---GGGGCAACCCCGGGUUGUCAUAUAGCUAAAAGCGCGGUGCAUCCC-----NN 102
MBFV-NC_015843.2 CCC---GGCUCAGGCCGGGUUGUCAUAUAGCUAAAAGCGCGGAACGUCCC-----NN 102
MBFV-NC_006551.1 CCC---GGUAAAAACCGGGCUGUCAUAUAGCUGAAGCGGUAACCGCCGCG-----NN 102
MBFV-NC_009029.2 CCC---GGUAGAAAUCGGGGCGUCAUAUAGCUAAAAGCGCGGUGCGUCCCGCG-----NN 102
MBFV-NC_001563.2 CCC---GGUAAAAACCGGGCUGUCAUAUAGCUAAAACCGCGGUAUGCCCGCG-----NN 102
MBFV-NC_009942.1 CCC---GGCAAAAGCGGGCUGUCAUAUAGCUGAAGCGGUAUAGCCCGCG-----NN 102
MBFV-NC_032088.1 UCC---GGCAUUCGUCGGACUGUCAUAUAGCUAAAAGCGCGGAGCAUCCCGCG-----NN 102
MBFV-NC_009028.2 UCC---GCUGCAAAGCGGACUGUCAUAUAGCUAAAACGCGGUGGCUAGCGUGU-----NN 102
MBFV-NC_040776.1 CCC---GCCGGAAGCGGGCUGUCAUAUAGCUGAAGCGGUAACCGCGGUAUCCGUGU-----NN 102
MBFV-NC_034151.1 CCC---GGAAAGAACCGGGUCUGUUAUAUAGCUGAAGCGCACAUCCCGCGCUA-----NN 102
MBFV-NC_000943.1 CCC---GGGAAGCCCCGGGUGUCAUAUAGCUAAAACGCGGCAUACCCCGCG-----NN 102
MBFV-NC_007580.2 CCC---GGUAGAAACCGGGCUGUCAUAUAGCUGAAGCGGCUAUCGCCGCG-----NN 102
MBFV-NC_009026.2 CCC---GGAGGAAACCGGGUCUGUCAUAUAGCUAAAAGCGUACGGCGGCCAACG-----NN 102
ISFV-NC_017086.1 CCG---G---AAGACGGGCUAUAUAGUAGAGACGUGGUGCCUCGCGUCU---NN 102
ISFV-NC_027999.1 UCUCGGGUUAAAAACCCGAGGUGUCAUAUAGUAAUAGCGUACUGGAA-----NN 102
TBFV-NC_003687.1 GGAAAGGGGGGCGGUGCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAAGU-----NN 102
TBFV-NC_001672.1 GGAUAGGGGGGCGGUGCCCCUCGACGAGUGUCGAAAGAGACCGCA-----NN 102
TBFV-NC_005062.1 GGAAAGGGGGGCGGUGCCCCUCGACGAGUGUCGAAAGAGACCGCGA-----NN 102
TBFV-NC_003690.1 GGAAAGGGGGGCGGUGCCCCUCGACGAGUGUCGAAAGAGACCGCGA-----NN 102
TBFV-NC_001809.1 GGAAAGGGGGGCGGUGCCCCUCGACGAGUGUCGAAAGAGACCGCAA-----NN 102
TBFV-NC_027709.1 GGAAAGGGGGGCGGUGCCCCUCGACGAGUGUCGAAAGAGACCGCAA-----NN 102
0.....70.....80.....90.....100.....110.....1



NKV-NC_003635.1 NNNNNUUU-----UCCAAAUUCCGUUACCGGGGUGGGGUGAUUAGCCAUGGUCGCAGA 157
NKV-NC_004119.1 NNNNNUUG-----CCUCAUCUUGAAUGGGGCGCGGUGGAUAUAUACUCCAGCCAGAAA 157
ISFV-NC_027819.1 NNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUUAUGAUGACAAGAAUCAGAUGGAAG 160
ISFV-NC_008604.2 NNNNN-CUGACGCCACUCCCG-AAAUGUGUAUAGUAACAGAGAUCCUGCAGCAGCAG 160
ISFV-NC_021069.1 NNNNN-GUGACGCCACCCCCCG-AAUUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAA 159
ISFV-NC_012671.1 NNNNN-GUGACGCCACCCCCCG-AAUUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGCAA 159
ISFV-NC_024299.2 NNNNN---ACGACACCCCCCG-AAUUGUGUAUUAUGCCAAUGUGAGUCUAACAGAA 157
ISFV-NC_001564.2 NNNNN--AGCGAACUCCUCCGUUAUUGUAAUUAUGGUG-AAAAGACUCAUCUUAA 159
ISFV-NC_005064.1 NNNNN---CGUACCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCCAGCUAAA 157
ISFV-NC_012932.1 NNNNN-UUGCGAACCCUCCUG-AAAUGUGAAAAUUAUGGGGAAAAGUACCCAUCUAAG 160
ISFV-NC_034204.1 NNNNN---ACGCAACCCUCCUG-AAUUGUUAUAAACUGUGCGGAAAAAUCCAAAGACU 158
ISFV-NC_034017.1 NNNNN--GCGACCACCCUCCUG-UAAUGUUUAUUAAGUUGUGCGGAAAAAUCCAAGAAU 158
ISFV-NC_027817.1 NNNNNUAGUGACGUCACCCUCCGUG--UGUGAAAAAGCAUUGCGACUAGAACUAAAGCAG 160
MBFV-NC_008719.1 NNNNNCC-GAGCA-UUAUG-CGGACCAUUAUGACGUCUGGGAA--AGACCAGGAGACAC 156
MBFV-NC_012735.1 NNNNNCC-GAGCACACAUAG-CGGACCAUUAUGACGCCAGGGAA--AGACCAGGAGACAC 157
ISFV-NC_016997.1 NNNNN-C-AAAAUGUAUAG-GCUUAUUAUUGACACCUUAGGAA--AGACUGGAGAUAC 156
MBFV-NC_002031.1 NNNNNA--GGGAACAAAAGUGGGGACCAUUAUGACGCGCAGGGAA--AGACCAGGAGUGU 157
NKV-NC_005039.1 NNNNN-U-ACGCAUGGAU-UUUGCAAUAUGUUGACAUCAGGGAA--AGACCGGGUGUUU 156
MBFV-NC_002640.1 NNNNNCCCCCAACACA-A-AAACAGCAUAUUGACG--CUGGGAA--AGACCAGAGAUC 156
MBFV-NC_001474.2 NNNNNCCCCCGAAACAA-A-AAACAGCAUAUUGACG--CUGGGAA--AGACCAGAGAUC 156
MBFV-NC_001477.1 NNNNNCCCCGACAAACA-A-AAACAGCAUAUUGACG--CUGGGAG--AGACCAGAGAUC 156
MBFV-NC_001475.2 NNNNNCCCCCGGAAACGA-A-AAACAGCAUAUUGACG--CUGGGAG--AGACCAGAGAUC 156
MBFV-NC_012532.1 NNNNNCCCCCGGAAACGCA-AAACAGCAUAUUGACG--UUGGAA--AGACCAGAGACUC 156
MBFV-NC_035889.1 NNNNN-CCCGGAAAACGCA-AAACAGCAUAUUGACG--CUGGGAA--AGACCAGAGACUC 156
MBFV-NC_012533.1 NNNNNA--UAACCAACAA-A-AAACAGCAUAUUGACG--CUGGGAA--AGACCAGGAGACUC 156
ISFV-NC_040610.1 NNNNNC--UCAUUAUUUAUUAUAAACAGCAUAUUGACAC---CAGGUAUAGCCCG-AGAA 156
ISFV-NC_024017.1 NNNNNC--UGCUGAGAG--CAAAAUAGCAUAUUGACACUUGGGAAGAGACCAG-AGAU 156
MBFV-NC_001437.1 NNNNN-----CAUAUUGACAC-CUGGGAA-UAGACUUGGAGAGU 139
MBFV-NC_012534.1 NNNNNAA--CACCAAAACA-AAACAGCAUAUUGACAC-CUGGGAA-GAGACUAGGAGACC 156
MBFV-NC_015843.2 NNNNNGA--ACACCACAAG-AAACAGCAUAUUGACAC-CUGGGAA-UAGACUAGGAGACC 156
MBFV-NC_006551.1 NNNNNUA--AGCAUACAAA-AAACAGCAUAUUGACAC-CUGGGAAUUAUAGACUAGGAGAU 158
MBFV-NC_009029.2 NNNNNAG--CGCAAAACA-AAACAGCAUAUUGACAC-CUGGGAA--AGACUAGGGGAU 156
MBFV-NC_001563.2 NNNNNAA--ACACCAAAAG-AAACAGCAUAUUGACAC-CUGGGAA-UAGACUAGGGGAU 156
MBFV-NC_009942.1 NNNNNAA--CACCAACA-AAACAGCAUAUUGACAC-CUGGGAA-UAGACUAGGAGAU 156
MBFV-NC_032088.1 NNNNNUA--CCGCAAAACA-AAACAGCAUAUUGACAC-CUGGGAA-UAGACUAGGAGAU 156
MBFV-NC_009028.2 NNNNNCA--AGCGCGAA-ACAAACAGCAUAUUGACG-CUGGGAA--AAGACCGGGAGAU 156
MBFV-NC_040776.1 NNNNN-A--AGCGCAAGACAAACAGCAUAUUGACAC-CUGGGAA--AAGACCGGGAGAC 156
MBFV-NC_034151.1 NNNNNGA--A-CGCAAAUAAAAACAGCAUAUUGACAC-CUGGGAA-UAGACUUGGAGACC 156
MBFV-NC_000943.1 NNNNNAA--AAGCAUCAAAACAGCAUAUUGACAC-CUGGGAA-AAGACUAGGAGAU 157
MBFV-NC_007580.2 NNNNNCCGUUAACGCAAAAC--AACAGCAUAUUGACAC-CUGGGAA-AAGAC-AGGAGAU 155
MBFV-NC_009026.2 NNNNNA--CGCACAAAAGAAACAGCAUAUUGACAC-CUGGGAA-UAGACCGGAGAAU 156
ISFV-NC_017086.1 NNNNNC---CCAAAAUAGAAGGGCACUAUUAUCGACACUUGGGAA--AGACCAGAGGUAC 156
ISFV-NC_027999.1 NNNNNCCAUUGGAAAAUUU--AGCUUCAUUAUGACCCAGGGAAA--AGACCAGGAGAAC 157
TBFV-NC_003687.1 NNNNN-AUUUUUCCGCCUCCUUAUACAAUUU--CCCCGAGAAACUGGGGGGCGGUUC 159
TBFV-NC_001672.1 NNNNN-----UAUACAAA-UU-CCCCCUCGUAAGAGGGGGGGCGGUUC 144
TBFV-NC_005062.1 NNNNN-----UAUACAAA-UU-CCCCCUCGUAAGAGGGGGGGCGGUUC 144
TBFV-NC_003690.1 NNNNNGAUUUUUCCUCCUCCUAUACCAAAU--CCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_001809.1 NNNNNG-UUUUUUCCUCCUCCUAUACCAAAU--CCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_027709.1 NNNNNGAUUUUUCCUCCUCCUAUACCAAA-UU--CCCCCUCGACAGAGGGGGGGCGGUUC 159
20.....130.....140.....150.....160.....170.....1



NKV-NC_003635.1 GAUCAAGC-----UC 167
NKV-NC_004119.1 AGACUCAG-----AU 167
ISFV-NC_027819.1 AUCCCA 167
ISFV-NC_008604.2 GUCUCCA 167
ISFV-NC_021069.1 AUCUCCAC----- 167
ISFV-NC_012671.1 AUCUCCAC 167
ISFV-NC_024299.2 GCUUCCAC-----CG 167
ISFV-NC_001564.2 CCUCCAC----- 167
ISFV-NC_005064.1 UCUCCCAC-----CG 167
ISFV-NC_012932.1 CCUCCCA 167
ISFV-NC_034204.1 GUCCCCAC-----C- 167
ISFV-NC_034017.1 GUCCCCAC-----C- 167
ISFV-NC_027817.1 AGUCUC-----C 167
MBFV-NC_008719.1 UCCUUGAUUCU----- 167
MBFV-NC_012735.1 UCCUUGAUUC----- 167
ISFV-NC_016997.1 CUUUGGUUCU----- 167
MBFV-NC_002031.1 UCUCUGCUU-U----- 167
NKV-NC_005039.1 GUCGGUUCUG----- 167
MBFV-NC_002640.1 UGCUGUCUCG----- 167
MBFV-NC_001474.2 UGCUGUCUCCU----- 167
MBFV-NC_001477.1 UGCUGUCUCU----- 167
MBFV-NC_001475.2 UGCUGUCUCCU----- 167
MBFV-NC_012532.1 CAUGAGUUUC----- 167
MBFV-NC_035889.1 CAUGAGUUUC----- 167
MBFV-NC_012533.1 UGUUGGUUUC----- 167
ISFV-NC_040610.1 UACACAAUUU----- 167
ISFV-NC_024017.1 UGAUGUUGUU----- 167
MBFV-NC_001437.1 UUGUGUCUAUCUCAACAUCUACUA 167
MBFV-NC_012534.1 CUCUGUCCUA-A----- 167
MBFV-NC_015843.2 CUCUGUCCUA-A----- 167
MBFV-NC_006551.1 UUCUGUCU----- 167
MBFV-NC_009029.2 UGCUGUCUCG-G----- 167
MBFV-NC_001563.2 UUCUGUCUCG-C----- 167
MBFV-NC_009942.1 UUCUGUCUCG-C----- 167
MBFV-NC_032088.1 CGCUGUCUCG-G----- 167
MBFV-NC_009028.2 CUCUGCUUC-C----- 167
MBFV-NC_040776.1 CUCUGUCUCG-C----- 167
MBFV-NC_034151.1 CUCUGCUUAU----- 167
MBFV-NC_000943.1 UUCUGUCUUA----- 167
MBFV-NC_007580.2 CCCUGCUUUU----- 167
MBFV-NC_009026.2 UGCUGC-CUCG----- 167
ISFV-NC_017086.1 UGCUGAUUC-----A 167
ISFV-NC_027999.1 UCUUGGUU----- 167
TBFV-NC_003687.1 UGUUUC----- 167
TBFV-NC_001672.1 UUGUUCUCCUG-----AGCCACCAUCA 167
TBFV-NC_005062.1 UUGUUCUCCUG-----AGCCACCAUCA 167
TBFV-NC_003690.1 UUGUUC----- 167
TBFV-NC_001809.1 UUGUUC----- 167
TBFV-NC_027709.1 UUGUUC----- 167
80.....190.....200.....

