

ISFV-NC_001564.2 -----GAGAGACCGUGAGGUUGAACCCGGCAAGGAUGAAGAGGAAAGGAU 45
ISFV-NC_005064.1 -----AGUUGUUCGGAAAAGUGACCUUGGUUCUGUAUGAAUAGAGGAAGC 45
ISFV-NC_012932.1 -----AGUUGUUUGGAAAAGAGAUACUCUGGUUCGUAUGAAUAGGAUGCUC 45
ISFV-NC_008604.2 -----UUGUUUAUAAUAGUCGUUUUUCUCGCGAAGAAUGGGAAGGACGCAC 45
ISFV-NC_021069.1 -----CCUUAAUUAACUCUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA- 44
ISFV-NC_012671.1 -----CUUUAAUUAACUCUGGUACUAAACGUGGCGCUAUGAAGAAGAAGGA- 44
ISFV-NC_027819.1 -----UCACGCUAUGAGCUUGAGAAGAAUGUUUACUUAUGUCAAGUAAGGAA 45
ISFV-NC_027817.1 -----CAAACGUCUGGUUCCAGCGUUGUUAUAAUAGUGAGGAUUAGGA 45
ISFV-NC_034204.1 -----GCGACAGAAGGAACCCAAAAGGGGAACCAGAUG--GAGCAAACCG 43
ISFV-NC_034017.1 -----UAAAAAUUUUUUUGGCGACGUACGGAAAGUAUG--GAGCAAUCGA 43
ISFV-NC_024299.2 -----UUGAAAAACUUUUUGCGACGUUGUUGAGAAGAUG--GAGCGAA--A 41
ISFV-NC_016997.1 CAUUAGAAUAGGUUUGAAAGAGUUGAAUCA-----AUGAGAUCAAUGG-U 44
MBFV-NC_012735.1 -----AAGGAGUAGAAGAAAG-GAGAUUCAUUCCTCAUUGGCAACAAAGG-G 44
MBFV-NC_008719.1 -----AAAGGUAGAGAAAAG-GAGUUUUGCUUCUCAUGGCAACAAAGG-C 44
ISFV-NC_027999.1 -----UUUUUUGAGAGAACAGCGAGAGUAUACUACU-AUGGUUAGAGAAAG-G 44
MBFV-NC_002031.1 -----UUGAGCGAUUAGACGA-GAACUGACCGAACGAACUGUC----- 35
NKV-NC_005039.1 -----CGGAAGAAUACUGUCCGGAAUCAAGACGAUG-----AAAAA-U 39
ISFV-NC_040610.1 -----AAAGAAGUCAAAACUAGAAAAUUUA-UUU-CA-----AUGGUU---AGAC-A 41
ISFV-NC_024017.1 -----UGAU-AGAGUUGACUAGAUAAACGAUUUAU--CG-----AUGGU---ACUA-A 41
ISFV-NC_017086.1 -----UGAAUAGGUUUUAUUGUGAAUCGAAA-UAC--CG-----AUGGCG---AAUA-A 41
MBFV-NC_009028.2 -----UCUGAAUAGUUUUUU--AGCAAGGGA-UUUC--CG-----AUGUCU---AAAA-A 41
MBFV-NC_040776.1 -----UCUGAAUAGUUUUUU--AGCAGAGGU-ACUACC-----AUGUCU---AAAA-A 41
MBFV-NC_034151.1 -----UGAAUAGUUUUUU--GAACAGAGGUGAAAAC-----AUGUCU---AAAA-A 41
MBFV-NC_000943.1 -----UGAACAGUUUUUUGGAGCUUUUGAUUU--CAA-----AUGUCU---AAAA-A 41
MBFV-NC_006551.1 -----GCCGGCAGUUUCUUUGAGC-GUUGAUUU--UCA-----AUGUCU---AAGA-A 41
MBFV-NC_001563.2 -----UGCGAGCUGUUUUUGGAC-GAAGAUUU--C-G-----AUGUCU---AAGA-A 41
MBFV-NC_009942.1 -----UGCGAGCUGUUUUUAGCAC-GAAGAUUU--C-G-----AUGUCU---AAGA-A 41
MBFV-NC_007580.2 -----GCGAACAGUUUUUU--AGCAGGGAAUUAACCA-----AUGUCU---AAAA-A 41
MBFV-NC_009026.2 -----UCG-----AGGAAGACGAGAGCGAAUUCUCUGA-----AUGGCU---AGAA-A 41
MBFV-NC_012534.1 -----AGUUUGAAC--GUUUUUUGGAGAGAGAUUAU-----AUGUUUAACAAA-A 44
MBFV-NC_015843.2 -----AGUUUGAAC--GUUUUUUGGAUAGAGACAACU-----AUGUCUAACAAA-A 44
MBFV-NC_009029.2 ACACAGUGUGAACAGUUUUUUUGAACAGCA-----AUG--ACCAAAA-A 41
MBFV-NC_032088.1 -----AGUGUGAACAGUUUUUUUGAUCAGUG--GAUA-----AUG--ACUAAAA-A 41
MBFV-NC_001437.1 -----UUAACAGUUUUUUUAGAACCGGAAGUAACC-----AUGACU---AAAA-A 41
MBFV-NC_012532.1 -----AUUUGGAUUU--GGAAACGAGAG-UUUCUGG--UCAUGAAAAACCCCA-A 44
MBFV-NC_035889.1 -----UUUUGGAUUU--GGAAACGAGAG-UUUCUGG--UCAUGAAAAA-CCCA-A 43
MBFV-NC_002640.1 -----UGUUUGAAUAGAGAGCAGA-UCUCUGGAAAAAUG--AA--CCA-A 39
MBFV-NC_001477.1 -----AACAGUUUUUUUAUAGAGAGCAGA-UCUCUG-----AUGAACAA--CCA-A 42
MBFV-NC_001475.2 -----GACAGUUUUUUUAUAGAGAGCAGA-UCUCUG-----AUGAACAA--CCA-A 42
MBFV-NC_001474.2 -----ACAGUUUUUUUAUAGAGAGCAGA-UCUCUG-----AUGAAUAA--CCA-A 42
MBFV-NC_012533.1 -----GUGUGUGAAAGAAAACAGAUUCUUCGGAA--AUGA-----A-A 37
TBFV-NC_001809.1 -----AAAGACAGUUAAGGAGAACAGAGCUGGGG-----AUGGGCAGGAAGACC 45
TBFV-NC_027709.1 -----AAAGACAGCUUAGGAGAACAGAGCUGGGG-----AUGGGCAGGAAGUCC 45
TBFV-NC_003690.1 -----AAAGACAGCUUAGGAGAACAGAGCUGGGG-----AUGGCCGGGAAGGCC 45
TBFV-NC_001672.1 -----AAAGACAGCUUAGGAGAACAGAGCUGGGG-----AUGGCUAAGAAAGCC 45
TBFV-NC_005062.1 -----AAAGACAGCUUAGGAGAACAGAGCUGGGG-----AUGGCCGGGAAGGCC 45
TBFV-NC_003687.1 -----AGCUUAGGAGAACAGAGCUGGGGAGUGGUUAUG-----AUGACC 39
NKV-NC_004119.1 -----CCGGUGGAAACACGUCGCUUCUGUCAAAUAGACGAAAGGCAAA 45
NKV-NC_003635.1 -----GUGCGCGUCACGCCACUUUUUGUGGCGGGAAUGUGCGAAAAUAAC 45
.....10.....20.....30.....40.....50.....6



ISFV-NC_001564.2 UUGGAGGCG--CGAGGAAAAG--CGCCAGGGAGGGAUUUUCCACCCCGUUCUGGGGAA- 100
ISFV-NC_005064.1 UUGAAGAAG--GGGCCUCCGG--GCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCCG- 100
ISFV-NC_012932.1 UUGAAGAAA--GGGCCUUCGG--GCCAAACAUUCUGGUCGGCAUCGACCAGCCGCCCG- 100
ISFV-NC_008604.2 GGUAAGAAGAAGAGAGGGGCGG--GGCUCUCCGGAGUGGUUUUACCA--CCGGGGCGCG- 100
ISFV-NC_021069.1 -GUCGGGAGUAGUGGAGAGAG--GACGCGGUACGCGCCUCCGACCAACCGUGGAGGAA- 100
ISFV-NC_012671.1 -GUCUGGAGUGGUGGAGAGAG--GACGUGGAACGCGUCCUGCCAAUCGUGGAGGAG- 100
ISFV-NC_027819.1 GCACUGAAGAAGAGAGGGAUU--UCCUGGGGAGGAGGAGGAAACGGGGCUUUGAG--A- 100
ISFV-NC_027817.1 GGCCUUCUUCUCCCGUCCGAGGGGAAGAAAAGAAAG----CCCCUGUGAUACAUCACAGG- 100
ISFV-NC_034204.1 GAAAGUUUUCUCCGACCGGGAGGUGGUCG--GCCCGGCCACAGAGGGGCGUGGGGAAGA- 100
ISFV-NC_034017.1 GGAAGUUUUUCCCGCCGGGAGGCGGUCG--GCCGGCCCCAAGAGGGGCGUGGCGAAGA- 100
ISFV-NC_024299.2 GAAAGGAGUAAAUUUGGGGAGAAGGGAAGCUAUCCGCCCCUCCCAUUAAGGAGAAAA- 100
ISFV-NC_016997.1 CUUUCGAGCUAGGAGGCCCGUUAACGGGCGGUCGAUUAUAAUAAAACGUAAGCUGC---- 100
MBFV-NC_012735.1 AGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUUUGGU-----GACGGCCAA 95
MBFV-NC_008719.1 UAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUUUGAU-----UAGGCGUAC 95
MBFV-NC_002031.1 UGGUCGUAAAGCUCAGGGAUAAACCCUGGGCGUCAAUUUGGUACGACGAGGAGUUCGUC 95
NKV-NC_005039.1 CGUAGACAAAACACCCG----AAUUCGGGUGUCAAUUAUAAAAAAGCAAAAUCCAAAAA 95
ISFV-NC_040610.1 ACUUGGAAAGACC-----GUUGCAAGACGGUCUGUCAAUUUGGCAAAGC--GGAUACCAAC 96
ISFV-NC_024017.1 ACCAGGAAAGCCC-----GCCGAAAGCGGUCGUUAUUAUAGCUAACAC--GCAUACCAAG 96
ISFV-NC_017086.1 ACCCAAGAAGCCC-----GG--AAGACGGGCUAUCGAUUAUGAGAGACGUGCGUUGCCUCG 95
MBFV-NC_009028.2 ACCAGGAAAGUCC-----GCUGCAAAGCGGACUGUCAAUUUGCUAAAACGCCUGGCUAGCGU 98
MBFV-NC_040776.1 ACCAGGAGGACCC-----GCCGGAAGGCGGUGUCAAUUAGCUAAAACGCCAGCCAGCGU 98
MBFV-NC_034151.1 ACCAGGAGGACCC-----GGAAAGAACC GGUGUCAAUUAGCUAAGCGCACAUCCCGCGC 98
MBFV-NC_000943.1 ACCAGGAGGACCC-----GGGAAGCCCGGGUGUCAAUUAGCUAAAACGCCGCAUACCCCG 98
MBFV-NC_006551.1 ACCAGGAGGCGCC-----GGAAGAAACCGGGCUAUAUAGCUAAGCGCGGCAUACCCCG 98
MBFV-NC_001563.2 ACCAGGAGGCGCC-----GGUAAAAACCGGGCUGUCAAUUAGCUAAAACCGCGUAUGCCCG 98
MBFV-NC_009942.1 ACCAGGAGGCGCC-----GGCAAGAGCCGGGUGUCAAUUAGCUAAAACCGCGGAUUGCCCG 98
MBFV-NC_007580.2 ACCAGGAAACACC-----GUGAAGAAACCGGGUGUCAAUUAGCUAAAACCGCGGCAUCCCG 98
MBFV-NC_009026.2 ACCAGGAAGACC-----GGAGGAAACCGGGUGUCAAUUAGCUAAAAGCGUACGGCGGCCAA 98
MBFV-NC_012534.1 ACCAGGAAACACC-----GGGCAACCCGGGUUGUCAAUUAGCUAAAAGCGCGGUGCAUCCC- 100
MBFV-NC_015843.2 ACCAGGAAAGACC-----GGUCAGGCGGGGUUGUCAAUUAGCUAAAACCGCGGAGUCCCC- 100
MBFV-NC_009029.2 ACCAGGAAGGCC-----GGUAGAAAUCCGGGCGUCAAUUAGCUAAAAGCGCGGUGCGUCCG 98
MBFV-NC_032088.1 ACCAGGAAAGUCC-----GGCAUUCGUCGGACUGUCAAUUAGCUAAAAGCGCGGAGCAUCCCG 98
MBFV-NC_001437.1 ACCAGGAGGCGCC-----GGUAAAACCGGGCUAUAUAGCUAAGCGCGGCAUACCCCG 98
MBFV-NC_012532.1 AGAAGAAAUCC-----C-----GGAGGA-UCCGGAUUGUCAAUUAGCUAAAACCGCGAGUAGCCG 98
MBFV-NC_035889.1 AAAAGAAAUCC-----C-----GGAGGAUUCGGAUUGUCAAUUAGCUAAAACCGCGGAGUAGCCCG 98
MBFV-NC_002640.1 CGAAGAAAGG--U-----GUUAGACC--ACCUUUAUUAUAGCUAAGCGCGGCAUACCCCG 92
MBFV-NC_001477.1 CGGAAAAAGA--C-----GGGUCGACC--GUCUUUCAAUUAGCUAAGCGCGGAGAAACCG 95
MBFV-NC_001475.2 CGGAAGAAGA--C-----GGGAAACCC--GUCUAUCAAUUAGCUAAGCGCGGAGAAACCG 95
MBFV-NC_001474.2 CGGAAAAAGG--C-----GAAAAACAC--GCCUUUCAAUUAGCUAAGCGCGGAGAAACCG 95
MBFV-NC_012533.1 CCCAAAAAAG--G-----GCCGCAUCGCGGCCUAUCAAUUAGCUAAGCGCUUUGGAUCUAG 92
TBFV-NC_001809.1 AUCCUGAAAGGCAAGGGGGGGCGUCCCCCUCGACGAGUGUCGAAAGAGACCCGGA----- 100
TBFV-NC_027709.1 AUCCUGAAAGGUAAGGGGGGGCGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_003690.1 GUUCUAAAAGGAAAGGGGGGGGGUCCCCCUCGACGAGCGUUCGAAAGUGGCCCCAA----- 100
TBFV-NC_001672.1 AUCCUGAAAGGUAAGGGGGGGCGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_005062.1 AUUCUGAAAGGUAAGGGGGGGCGUCCCCCUCGACGAGUGUCGAAAGAGACCCGUA----- 100
TBFV-NC_003687.1 ACUUCUAAAGGAAAGGGGGGGCGUCCCCCUAGGCGCAAGCUUAAAGUGACCCGCAAAUAG 99
NKV-NC_004119.1 AUGAAAAACGCGAAUAAAAAGAAAGCAAAAAACUCUAAAGACCAUCCUGGUCGAA----- 100
NKV-NC_003635.1 AGGAAAAAGAACCUCUGGAAAGGGUUCAAAGAAAGAAAGAAUGUGGAAGGA----- 100
0.....70.....80.....90.....100.....110.....1



ISFV-NC_001564.2 -----NNNNNNNAG--CGAACCUCUCCGUUAAUGUGAAAAAGUAUGGGG--AAAGAAC 150
ISFV-NC_005064.1 -----NNNNNNNN--CGUACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAAC 148
ISFV-NC_012932.1 -----NNNNNNNNUUGCGAACCCCCUCCG-AAUUGUGAAAAAUUUGGGGAAAAGUAC 151
ISFV-NC_008604.2 -----NNNNNNNNCUG-ACGCCACUCCCGAAUUGUGUAUGUAACAGAGC-AUGCCU 150
ISFV-NC_021069.1 -----NNNNNNNNGUG-ACGCCACCCCCCGAAUUGUGUAUUGUGACAG-GC-UUGCUC 149
ISFV-NC_012671.1 -----NNNNNNNNUUA-GCCACACCCCCGAAACGUGAUUUUAU-AUGAGACAGAAU 150
ISFV-NC_027819.1 -----NNNNNNNNUAGCGUCACCCCCCG-UUGUGAAACGAUUGCGACUAGAAU 151
ISFV-NC_034204.1 -----NNNNNNNNACGA--CCACCCUCUGAAUUGUUAUAAUAAAACUGUGCGGAAAAAU 149
ISFV-NC_034017.1 -----NNNNNNNNGCGA--CCACCCUCUGAAUGUUUAUAAUAGUUGUGCGGAAAAAU 149
ISFV-NC_024299.2 -----NNNNNNNNACGA--CACCCCCGAAUUGUGUAUUAUAGGCCAUUGAGUCU 148
ISFV-NC_016997.1 -----NNNNNNNNCAA--AAUUGUAUAGGCUAUUAUUAUCGACA-CCUAGGAAAGACUG 149
MBFV-NC_012735.1 AGUGA--NNNNNNNNCC--GAGCACAUAGCGGACCAUUAUUGAGC-CCAGGG-AAAGACC 149
MBFV-NC_008719.1 GGCCA--NNNNNNNNCC--GAGCAC-UAUAGCGACCAUUAUUGAGC-CCUGGG-AAAGACC 148
ISFV-NC_027999.1 UGGAA--NNNNNNNNCCAUGGGAUUUAGCU--UCAUAUUGACA-CCAGGGAAAAGACC 149
MBFV-NC_002031.1 CUUGU--NNNNNNNNAGGGAACAAAU-AGUGGGACCAUUAUUGAGC-CCAGGG-AAAGACC 149
NKV-NC_005039.1 CAAA--NNNNNNNNUAGCGUAAGAUUUUGC--AAUAUUGUGACA-UCAGGGAAAGACCG 149
ISFV-NC_040610.1 CG-CU--NNNNNNNNCUCAUUAUUU-AUAUAACAGCAUUAUUGACA-CCAGGG-AUAGACC 149
ISFV-NC_024017.1 CG-UG--NNNNNNNNNCU-GCUGAGA-GCAAAUAGCAUUAUUGACA-CUUGGGAAGAGACC 149
ISFV-NC_017086.1 CGUCU--NNNNNNNNCCCAAAUUG-A-AGGGCAUUAUAGGACACUUGGG--AAAGACC 148
MBFV-NC_009028.2 GU-----NNNNNNNNCAAGCGCGAA--ACAAACAGCAUUAUUGAGC-CCUGGGA--AAGAC 147
MBFV-NC_040776.1 UU-----NNNNNNNN--AAGCGCAA-GACAAACAGCAUUAUUGACA-CCUGGGA--AAGAC 147
MBFV-NC_034151.1 UA-----NNNNNNNNGAA-CGCAA-AAAGAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_000943.1 CG-----NNNNNNNNAAAAGCAUC--AAACAACAGCAUUAUUGACA-CCUGGGAA-AAGAC 148
MBFV-NC_006551.1 CG-----NNNNNNNN--UAAGCAUCA-AAAAACAGCAUUAUUGACA-CCUGGGAUUAGAC 149
MBFV-NC_001563.2 CG-----NNNNNNNNAAACACC-AA-AGAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_009942.1 CG-----NNNNNNNNAAACACCA-CA-ACAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_007580.2 CG-----NNNNNNNNCCGUAAACG-AAACAACAGCAUUAUUGACA-CCUGGGA--AGAC 147
MBFV-NC_009026.2 CG-----NNNNNNNN-ACGACCAA-AAAGAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_012534.1 -----NNNNNNNN-AACACCAA-ACAAACAGCAUUAUUGACA-CCUGGGA--GAGAC 147
MBFV-NC_015843.2 -----NNNNNNNNNGAACACCA-A-GAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_009029.2 CG-----NNNNNNNN-AGCGCAAC-ACAAACAGCAUUAUUGACA-CCUGGGA--AAGAC 147
MBFV-NC_032088.1 CG-----NNNNNNNN-UACCGCAA-ACAAACAGCAUUAUUGACA-CCUGGGA--UAGAC 147
MBFV-NC_001437.1 CG-----NNNNNNNN--CAUUAUGACA-CCUGGGAA-UAGAC 130
MBFV-NC_012532.1 UG-----NNNNNNNNCCCGGAAAACGCAAAACAGCAUUAUUGAC--G-UGGGA--AAGAC 147
MBFV-NC_035889.1 UG-----NNNNNNNN-CCCGGAAAACGCAAAACAGCAUUAUUGAC--GCUGGGA--AAGAC 147
MBFV-NC_002640.1 CGUAUCAANNNNNNNNCCCCCAACAA-A-AAAACAGCAUUAUUGAC--GCUGGGA--AAGAC 147
MBFV-NC_001477.1 CGUGU--NNNNNNNNCCCCCGACAACAA-CAAACAGCAUUAUUGAC--GCUGGGA--GAGAC 147
MBFV-NC_001475.2 UGUGU--NNNNNNNNCCCCCGCAAAUA-AAAACAGCAUUAUUGAC--GCUGGGA--GAGAC 147
MBFV-NC_001474.2 CGUGU--NNNNNNNNCCCCCGCAAAACAA-AAAACAGCAUUAUUGAC--GCUGGGA--AAGAC 147
MBFV-NC_012533.1 CGCUGGCCNNNNNNNNNAUACCAA--CAUAAAACAGCAUUAUUGACA-CCUGGGA--AAGAC 147
TBFV-NC_001809.1 -----NNNNNNNNG-UUUUUCUCCUCCUAUACCAAAUUU-CCCCUCGACAGAGGGG 150
TBFV-NC_027709.1 -----NNNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UU-CCCCUCGACAGAGGGG 150
TBFV-NC_003690.1 -----NNNNNNNNGAUUAUUUCCUCCUCCUAUACCAAA-UU-CCCCUCGACAGAGGGG 150
TBFV-NC_001672.1 -----NNNNNNNN--UAUACAAA-UU-CCCCUCGUAAGAGGGG 135
TBFV-NC_005062.1 -----NNNNNNNN--UAUACAAA-UU-CCCCUCGUAAGAGGGG 135
TBFV-NC_003687.1 U-----NNNNNNNN-AUUUUUCCGCCUCCUAUACUAAUUU-CCCCCAGAACUGGGG 150
NKV-NC_004119.1 -----NNNNNNNNUUGCCUCAUCUUGAAUGGGGGGCG-----GCGUGGAUUAUUAAC 145
NKV-NC_003635.1 -----NNNNNNNNUUUCCAAAUUCC-CGUUACCGGG-----GGUGGGGUGAUUAG 144
20.....130.....140.....150.....160.....170.....1



ISFV-NC_001564.2 UCAUCUUAACCUCCAC----- 167
ISFV-NC_005064.1 CCAUCUUAACCUCCAC-----G 167
ISFV-NC_012932.1 CCAUCUUA--AGCCUCC-----A 167
ISFV-NC_008604.2 GCAGCAGCAGGUCUCCA----- 167
ISFV-NC_021069.1 GCAGCAGCAAAUCUCCAC----- 167
ISFV-NC_012671.1 GCAGCAGCAAAUCUCCAC----- 167
ISFV-NC_027819.1 CAGAUGGAAGAUCUCCCA----- 167
ISFV-NC_027817.1 AAAGUCGAGAGUC-----C 167
ISFV-NC_034204.1 CCAAAGACUGUCCCAAC----- 167
ISFV-NC_034017.1 CCAAAGAAUGUCCCAAC----- 167
ISFV-NC_024299.2 AUACAACAGGCUUCCAC-----G 167
ISFV-NC_016997.1 GAGAUACCUUUGGUUCU----- 167
MBFV-NC_012735.1 GGAGACACUCCUUGAUUC----- 167
MBFV-NC_008719.1 GGAGACACUCCUUGAUUC----- 167
ISFV-NC_027999.1 GGAGAAACUUCUGCUU-A----- 167
MBFV-NC_002031.1 GGAGUGGUUCUUGCUU-U----- 167
NKV-NC_005039.1 GGUGUUUGUCGGU-UCCGG----- 167
ISFV-NC_040610.1 GGAGAAUACACAAUUC----- 167
ISFV-NC_024017.1 AGAGAAUUG-ACAUUGUU----- 167
ISFV-NC_017086.1 AGAGGUACUCGUGAUUCA----- 167
MBFV-NC_009028.2 CGGGAGAU-CCUCUGCUUUC----- 167
MBFV-NC_040776.1 CGGGAGAC-CCUCUGCUUG----- 167
MBFV-NC_034151.1 UGGGAGAC-CCUCUGCUUAAU----- 167
MBFV-NC_000943.1 UAGGAGAU-CUUCUGCUUA----- 167
MBFV-NC_006551.1 UAGGAGAU-CUUCUGCUU----- 167
MBFV-NC_001563.2 UAGGGGAU-CUUCUGCUUG----- 167
MBFV-NC_009942.1 UAGGAGAU-CUUCUGCUUG----- 167
MBFV-NC_007580.2 AGGAGAU-CCUCGUUUUU----- 167
MBFV-NC_009026.2 CGGAGAAU-UUGCGUCCUG----- 167
MBFV-NC_012534.1 UAGGAGAC-CCUCUGCUUAA----- 167
MBFV-NC_015843.2 UAGGAGAC-CCUCUGCUUAA----- 167
MBFV-NC_009029.2 UAGGGGAU-UUGCGUCUUG----- 167
MBFV-NC_032088.1 UAGGAGAU-CCGCGUCUUG----- 167
MBFV-NC_001437.1 UGGGAGAU-CUUCUGCUU----- 167
MBFV-NC_012532.1 CAGAGACUCCAUAGAGUUUC----- 167
MBFV-NC_035889.1 CAGAGACUCCAUAGAGUUUC----- 167
MBFV-NC_002640.1 CAGAGAUCUCCUGGUCUUG----- 167
MBFV-NC_001477.1 CAGAGAUCUCCUGGUCUUA----- 167
MBFV-NC_001475.2 CAGAGAUCUCCUGGUCUUC----- 167
MBFV-NC_001474.2 CAGAGAUCUCCUGGUCUUC----- 167
MBFV-NC_012533.1 CGGAGACUCUGUUGGCUUUC----- 167
TBFV-NC_001809.1 GGGCGGUU-CUUGUUCU----- 167
TBFV-NC_027709.1 GGGCGGUU-CUUGUUCU----- 167
TBFV-NC_003690.1 GGGCGGUU-CUUGUUCU----- 167
TBFV-NC_001672.1 GGGCGGUU-CUUGUUCU-----A 167
TBFV-NC_005062.1 GGGCGGUU-CUUGUUCU-----A 167
TBFV-NC_003687.1 GGGCGGUU-CUUGUUCU----- 167
NKV-NC_004119.1 CCAUGCCAGAAAGACUCAGAU----- 167
NKV-NC_003635.1 CCAUGGCGCAACAGAUCAAGU-----C 167
80.....190.....200.....210.....

