

ISFV-NC_034204.1 -----GCGACAGAAAGGAACCCCAAAAGGGGAACC---AGAUGGAGCAAAACG---GAAA 47
ISFV-NC_034017.1 -----UAAAAUUUUUUUGGCGACGUACGGAAA---GUAUGGAGCAAUUCG---GGAA 47
ISFV-NC_024299.2 -----UUGAAAAACUUUUUGCGACGUUGUUGAGA---AGAUGGAGCGAA--A---GAAA 45
ISFV-NC_027817.1 -----CAAACUCUGGUGUCCAACGUUUGUAUUA---AAUUG-AGUGGAUUA---GGAG 46
ISFV-NC_027819.1 -----UCACGGAAUAGCUUGAGAGAAGAAGCUUAC---UUUUGUCAAGUAAGG---AAGC 47
ISFV-NC_008604.2 -----UUGUUUUAUAAUAGUCGUUUUUUCUCGCAG---AAAUGGGAAAGGACG---ACGG 47
ISFV-NC_021069.1 -----CCUUUAUUACCUCUGGUACUAAGUGGCA---CUAUGAAGAAAGAGG---A--G 45
ISFV-NC_012671.1 -----CUUUUAUUCCUCUGGUACUAACGUGGCG---CUAUGAAGAAGAAGG---A--G 45
ISFV-NC_001564.2 -----GAGAGACCGUGAGGUUGAACCCCGGCAAG---GAAUGAAGAGGAAGG---A--- 44
ISFV-NC_005064.1 -----AGUUGUUUCGGAAAAGAGUACCUUGGUUCG---UUUAGAAUAGAGGAA---G--- 44
ISFV-NC_012932.1 -----AGUUGUUUGGAAAAGAGAUUUCUGGUUC---GUAUGAAUAGGAUGC---U--- 44
TBFV-NC_001809.1 AA---AGACAGCUUAGGAGAAACAAGAGCUGGGG---AUGGGCAGGAAGACCAUCCUG 51
TBFV-NC_027709.1 AA---AGACAGCUUAGGAGAAACAAGAGCUGGGG---AUGGGCAGGAAGUCCAUCUCCUG 51
TBFV-NC_003690.1 AA---AGACAGCUUAGGAGAAACAAGAGCUGGGA---AUGGCCGGGAAGGCCGUUCUA 51
TBFV-NC_001672.1 AA---AGACAGCUUAGGAGAAACAAGAGCUGGGG---AUGGUCAAGAAGGCCAUCCUG 51
TBFV-NC_005062.1 AA---AGACAGCUUAGGAGAAACAAGAGCUGGGG---AUGGCCGGGAAGGCCAUUCUG 51
TBFV-NC_003687.1 -----AGCUUAGGAGAAACAAGAGCUGGGGAGUGGUUAUG---AUGACCACUUCU 45
MBFV-NC_001437.1 UUA AACAGUUUUUUAGAACCGGAAGUAACC---AUGACUAAAAAA---CCAG 46
MBFV-NC_002640.1 -----UGUUUGAAUAGAGA-GCAGA-UCUCUGGAAAAAUG---AA---CCA---ACGA 42
MBFV-NC_001477.1 AAC---AGUUUUUUUAUAGAGA-GCAGA-UCUCUG---AUGAACAA---CCA---ACGG 45
MBFV-NC_001475.2 GAC---AGUUUUUUUAUAGAGA-GCAGA-UCUCUG---AUGAACAA---CCA---ACGG 45
MBFV-NC_001474.2 ACA---GUUUUUUUAAUAGAGA-GCAGA-UCUCUG---AUGAAUAA---CCA---ACGG 45
MBFV-NC_012532.1 -----AUUUGGAUUUGGAAAC-GAGAG-UUUCUGG---UCAUGAAAAACCCCA---AAGA 47
MBFV-NC_035889.1 -----UUUUGGAUUUGGAAAC-GAGAG-UUUCUGG---UCAUGAAAAA---CCCA---AAAA 46
MBFV-NC_012533.1 -----GUGUUGAGAAAGAAAC-AACAAGUUCUUCG---AAUUGAAAC---CCA---A--- 42
ISFV-NC_017086.1 UGA AUAGGUUUUAUUGUGAAUCGAAA-UACCC---AUGGCGAAUAAA---CCCA 46
ISFV-NC_040610.1 AAAGAAGUCAAAACUAGAAAUUUUA-UUUC---AUGGUUAGACAA---CUUG 46
ISFV-NC_024017.1 UGAU-AGAGUUGACUAGAUAACGAAUUAUCG---AUGGUGACUAAA---UCA 46
MBFV-NC_009029.2 ACAC-AGUGUGAACAGUUUUUUUGAACACAG---AUGACC-AAAAAA---C---CAG 46
MBFV-NC_032088.1 -----AGUGUGAACAAAGUUUUUUUGAACAGUGGAU---AAUGACU-AAAAAA---C---CAG 46
MBFV-NC_012534.1 -----AGUUUGAACGUUUUUUUGGAGAGAGAUUAU---UAUGUUUAACAAAA---AACCAG 49
MBFV-NC_015843.2 -----AGUUUGAACGUUUUUUUGGAGAGAGACAAC---UAUGCUAACAAAA---AACCAG 49
MBFV-NC_007580.2 GC---AAGACGUUUUUU-AGC-AGGGAUUUACCC---AAUGUCUAAAAA---CCAG 46
MBFV-NC_009028.2 UC-U-GAAUAGUUUUUAGCA-AGGGAUUUCC---AUGUCUAAAAA---CCAG 46
MBFV-NC_040776.1 UC-U-GAAUAGUUUUUAGCA-GAGGUACUAC---AUGUCUAAAAA---CCAG 46
MBFV-NC_000943.1 -U-G-AACAGUUUUUUGGAGCUUUUGAUUUCAA---AUGUCUAAAAA---CCAG 46
MBFV-NC_006551.1 GC-C-GGCAGUUUCUUGAGC-GUUGAUUUUA---AUGUCUAGAAA---CCAG 46
MBFV-NC_001563.2 UG-C-GAGCUGUUUCUUGGCACGAAGUUC-G---AUGCUAAGAAA---CCAG 46
MBFV-NC_009942.1 UG-C-GAGCUGUUUCUUGACGAGAGAUUC-G---AUGUCUAGAAA---CCAG 46
MBFV-NC_009026.2 U---CGAGGA-AAGACGAGAAGCGAAUUCUCG---AAUGGCUAGAAA---CCAG 46
MBFV-NC_034151.1 U---GAUAG-UUUUUUGACAGAGGUGAAAAC---CAUGUCUAAAAA---CCAG 46
MBFV-NC_002031.1 -----UUGAGCGAUUAGCAGAGAACUGACCAGA---ACAUGUCUGU---CGUA 43
ISFV-NC_016997.1 -----CAUUAGAAUAGGUUUUGAAAGAGUUGAAUC---AAUGA-GAUCAAUGGUCUUUCG 50
ISFV-NC_027999.1 -----UUGUUUGAGAGAGAACGCGAAGUACUAC---U-AUGGUUGAGAAAACUAGAGUA 51
MBFV-NC_008719.1 -----AAGGGUAGAGAAAAG-GAGUUUUGCUUC---UCAUGG-CAACAAGAGGAGCA 50
MBFV-NC_012735.1 -----AAGGAGUAGAAGAAAG-GAGAUUCAUUC---CAAUGG-CAACAAGGGGAUGAA 50
NKV-NC_005039.1 CGGAAGAACACUCUGCCGGAUCAAAGACG-----AUGAAAAUUCGUA---GACA 47
NKV-NC_004119.1 C---CGGUGGAAACCGACGUCGCUUCGUA---AAUGACGAAAGGCA---AAUUG 48
NKV-NC_003635.1 G---UGCGCGUCACGCCACUUUUUGUGGCGGG---AAUGUGCGAAAAUA---ACAGG 48
.....10.....20.....30.....40.....50.....6



ISFV-NC_034204.1 GUUUUUCCCGACCGGGAG-GUGGUCGG--GCCCCGCCAGAGAGGGGCGUGGGGAAGA---- 100
ISFV-NC_034017.1 GUUUUUCCCGGCCCGGGAG-GCGGUCGG--GCGGGGCCCAAAGAGGGGCGUGGCGAAGA---- 100
ISFV-NC_024299.2 GGAGUAAUUUAUUGGGAG-AAGGGAAGCUAUCCCGCCCCCUCCAUAUAGGAGAAAA---- 100
ISFV-NC_027817.1 GCCUU-UUUUCCUCCG-AGGGAAGAAAAAGAAAGCCCCUGUGAUACAUAUCACAGG---- 100
ISFV-NC_027819.1 ACUGAAGAAGAAGGGAAUUUCCCU-GGGAGGAGGAGGAAAACGGGGCUUUGAG--A---- 100
ISFV-NC_008604.2 UAAGAAGAAGAAGGGGCGCGGCUC--CUCCGGAUGGUUGUUAACACCG-GGGCGCG--- 100
ISFV-NC_021069.1 UCGGGGAGUAGUGGAGAGGAGGACGC-GGUACGCGCCUCCCGACCAACCGUGGAGGAA--- 100
ISFV-NC_012671.1 UCUGGAGUGGUGGAGAGAGGACGU-GGAACGCGUCCUCCUGCCAAUCGUGGAGGAG--- 100
ISFV-NC_001564.2 UUUGGAGGCGCGAGGAAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAA--- 100
ISFV-NC_005064.1 CUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGCGCAUCGACCAGACCGCCGG--- 100
ISFV-NC_012932.1 CUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGCGCAUCGACCAGACCGCCGG--- 100
TBFV-NC_001809.1 AAAGGCAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_027709.1 AAAGGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_003690.1 AAAGGAAAGGGGGGGGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA----- 100
TBFV-NC_001672.1 AAAGGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_005062.1 AAAGGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCCGCA----- 100
TBFV-NC_003687.1 AAAGGAAAGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCCGCAAAUAAGU--- 100
MBFV-NC_001437.1 GAGGGCCCGGU---AAAAACCGGCUAUCAAUUAGCUGAAACGCGGCCUACCCCGCG--- 100
MBFV-NC_002640.1 AAAAAAGGUGG---UUAAGACCCACUUCAAUUAGCUGAAACGCGGAGAGAAAC--CGC-GUA 96
MBFV-NC_001477.1 AAAAAAGACGG---GUCGACCGUCUUUCAAUUAGCUGAAACGCGCGAGAAAC--CGC-GUG 99
MBFV-NC_001475.2 AAGAAGACGG---GAAAAACCGCUAUCAAUUAGCUGAAACGCGUGAGAAAC--CGU-GUG 99
MBFV-NC_001474.2 AAAAAAGGCGG---AAAACCGGCUUCAAUUAUGCUGAAACGCGGAGAGAAAC--CGC-GU 99
MBFV-NC_012532.1 AGAAAUCCGGA---GGAUCC-GGAUUGUCAAUUAGCUGAAACGCGGAGUAGC--CGG-UG- 100
MBFV-NC_035889.1 AGAAAUCCGGA---GGAUCCCGGAUUGUCAAUUAGCUGAAACGCGGAGUAGC--CGG-UG- 100
MBFV-NC_012533.1 AAAAGGGCGCG---AUCG-CGGCCUAUCAAUUAGCUGAAAGCGCUUUGGAUCU--AGC-GCU 96
ISFV-NC_017086.1 AGAAGCCCGG---AAGACGGGCUAUCAUUAUAGUGAGACGUGCGUUGCCUCGCGUCU- 100
ISFV-NC_040610.1 GAAAGACCGGU---GCAAGACGGGUCUGUCAAUUAGCGAAAGC--GGAUUACACCCG-CU- 100
ISFV-NC_024017.1 GAAAGCCCGG---GCAAGCGGGCUGUUAUUAUAGCUGAAACGCGGAGAGAAAC--CGC-UG- 100
MBFV-NC_009029.2 GAAGGCCCGGU---AGAAAUCCGGCCGUCAAUUAUGCUAAAGCGCGGUGCGUC--CCG-CG- 100
MBFV-NC_032088.1 GAAAGUCCCGG---AAUCGUCGGAGCUGUCAAUUAGCUGAAAGCGCGGAGCAUC--CCG-CG- 100
MBFV-NC_012534.1 GAAAACCCCGG---GCAACCGGGCUGUUAUUAUAGCUGAAACGCGGAGAGAAAC--CGC-CG- 100
MBFV-NC_015843.2 GAAGACCCGGC---UCAGGCCGGGUUGUCAAUUAGCUGAAAGCGCGGAACGUC--CC----- 100
MBFV-NC_007580.2 GAAAACCCCGU---AGAAACCGGGUUGUCAAUUAGCUGAAACGCGGGCGUAUC--CCG-CG- 100
MBFV-NC_009028.2 GAAAGUCCGCU---GCAAGCGGAGCUGUCAAUUAGCUGAAACGCGGAGAGAAAC--CGC-GU- 100
MBFV-NC_040776.1 GAGGACCCGCC---GGAAGCGGGUUGUCAAUUAGCUGAAACGCGCCAGCCAG--CGU-UU- 100
MBFV-NC_000943.1 GAGGACCCGGG---AAGCCCGGGGUCGUCAAUUAGCUGAAACGCGGGCAUACC--CCG-CG- 100
MBFV-NC_006551.1 GAGGGCCCGG---AGAAACCGGGCCAUCAAUUAGCUGAAACGCGGAGAGAAAC--CCG-CG- 100
MBFV-NC_001563.2 GAGGGCCCGGU---AAAACCGGGCUGUCAAUUAGCUGAAACGCGGUAUGCC--CCG-CG- 100
MBFV-NC_009942.1 GAGGGCCCGG---AAGAGCCGGGUCGUCAAUUAGCUGAAACGCGGAAUGCC--CCG-CG- 100
MBFV-NC_009026.2 GAAGACCCGGA---GGAACCCGGGUCGUCAAUUAGCUGAAAGCGCUACGGCGG--CAA-CG- 100
MBFV-NC_034151.1 GAGGACCCGGA---AAGAACCGGGUCGUUAAUUAUGCUGAAGCGCACAUCCCG--CGC-UA- 100
MBFV-NC_002031.1 AAGCUCAGGGA---AAAACCCUGGGCUGUCAAUUAGGUACGACGAGGAGUUCGCUCCUUGU- 100
ISFV-NC_016997.1 AGCUAGGAGGCCCGGUUAAACCGGGCCGUCGAUUAUAAACGUAAGCUGC----- 100
ISFV-NC_027999.1 GGAUCUCGGGUU---AAAACCCGAGGUGUCAAUUAGAUUAGGCGUACUGGAA----- 100
MBFV-NC_008719.1 CAAGUCUCGGGUCACGACCCCGGGGCGUCAAUUAGGUAGCGGCGCAAGGCA----- 100
MBFV-NC_012735.1 UAAGUCUCGGGUCAGUCCCGGAGGCGUCAAUUAGGUAGCGGCUAGAGUGA----- 100
NKV-NC_005039.1 AAACACCCG---AAUUUCG---GGUGUCAUUAUAAAAAGCAAAAUCCAAAAACAAA 99
NKV-NC_004119.1 AAAAACCGCAUAAAGAAAGGUAUCAAAGAAAGAAUUGCUGAAGGAUUG----- 100
NKV-NC_003635.1 AAAAAGAACCUGGAAAGGGUCAAAGAAGAAAGAAUUGCUGAAGGAUUG----- 100
0.....70.....80.....90.....100.....110.....1



ISFV-NC_034204.1 -----NNNNNNNNACG---ACCACCCUCUGAAUUGUAAUAAUAAACUGUGCGG-AAAAAUCCA 152
ISFV-NC_034017.1 -----NNNNNNNNGCG---ACCACCCUCUGAAUUGUAAUAAUAAUUGUGCGG-AAAAAUCCA 152
ISFV-NC_024299.2 -----NNNNNNNNACG---A-CACCCCCCGAAUUGUGUAUUAUAGCCCAUGUU-GAGUCUAUA 151
ISFV-NC_027817.1 -----NNNNNNNAGU---AGCGUACCCCCCGGUUGUGAAACAGAUUGCGA-CUAGAACUA 152
ISFV-NC_027819.1 -----NNNNNNNNUUA-GCCACACCCCCGAAACGUGAUUUUAU-AUGAGAC-AAGA-AUCA 152
ISFV-NC_008604.2 -----NNNNNNNNCUG-ACGCCACCUCCCGAAUUGUGUAUAGU-AACAGAGC-AUGC-CUGC 152
ISFV-NC_021069.1 -----NNNNNNNNGUG-ACGCCACCCCCGAAUUGUGUAUUGU-GACAG-GC-UUGC-UCGC 151
ISFV-NC_012671.1 -----NNNNNNNNGUG-ACGCCACCCCCGAAUUGUGUAUUGU-GACAG-GC-UUGC-UCGC 151
ISFV-NC_001564.2 -----NNNNNNNN-AGCGAACCUCCUCCGUAAUUGUGAAAAAGUAUGGGG-A-AAGACUCA 153
ISFV-NC_005064.1 -----NNNNNNNNCGU---ACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGA-AAGAACCCA 151
ISFV-NC_012932.1 -----NNNNNNNUUGCGAACACCCUCCG-AAUUGUGAAAAAUUAGGGGAA-AAGUACCCA 154
TBFV-NC_001809.1 -----NNNNNNNNG-UUUUUCUCCUCCUAUACCAAUUC---CCCUCGACAGAGGGGGG 152
TBFV-NC_027709.1 -----NNNNNNNNGAUUUUUCUCCUCCUAUACCAAU-UCC---CCCUCGACAGAGGGGG 152
TBFV-NC_003690.1 -----NNNNNNNNGAUUUUUCUCCUCCUAUACCAAU-UCC---CCCUCGACAGAGGGGGG 152
TBFV-NC_001672.1 -----NNNNNNNNUAUACAAAUAUCCCCUCCGU-AGAGGGG---GGGCGGUUCUUGUUCUC 152
TBFV-NC_005062.1 -----NNNNNNNNUAUACAAAUAUCCCCUCCAUUC-UGAGGGG---GGGCGGUUCUUGUUCUC 152
TBFV-NC_003687.1 -----NNNNNNNN-AUUUUUUCGCGUCCUAUACUAAUUC---CCAGGAAACUGGGGGG 152
MBFV-NC_001437.1 -----NNNNNNNNCAUUAUGACACCCUGGGAAUAGACUGGGAGAUUCUCG-----CUCUAUC 151
MBFV-NC_002640.1 UCAANNNNNNNNCCCCCAACA--CAAAAACAGCAUAUUGACG-CUGGGA--AAGACCA-G 150
MBFV-NC_001477.1 U---NNNNNNNNCCCGCAAC--CAAAAACAGCAUAUUGACG-CUGGGA--GAGACCA-G 150
MBFV-NC_001475.2 U---NNNNNNNNCCCCCGCAAA--UAAAAACAGCAUAUUGACG-CUGGGA--GAGACCA-G 150
MBFV-NC_001474.2 U---NNNNNNNNCCCCCGAAAC--AAAAACAGCAUAUUGACG-CUGGGA--AAGACCA-G 150
MBFV-NC_012532.1 -----NNNNNNNNCCCGGAAAC--CAAAAACAGCAUAUUGAC-G-UUGGGA--AAGACCA-G 150
MBFV-NC_035889.1 -----NNNNNNNN-CCCGGAAACGCAAAACAGCAUAUUGAC-G-CUGGGA--AAGACCA-G 150
MBFV-NC_012533.1 GGCCNNNNNNNNUAUACCAACA--UAAAACAGCAUAUUGACACUUGGGA--AAGACCG-G 150
ISFV-NC_017086.1 -----NNNNNNNNCCAAAUAUG-A-AGGGCACAUAUUGCAGACUUGGG--AAAGACCA-G 150
ISFV-NC_040610.1 -----NNNNNNNNCUCAUUAUUU-AUAUAACAGCAUAUUGACA-CCAGGG-AUAGACCG--G 151
ISFV-NC_024017.1 -----NNNNNNNNCU-GCUGAGA-GCAAAAAGCAUAUUGACA-CUUGGGAAGAGACCA--G 151
MBFV-NC_009029.2 -----NNNNNNNN-AGCGCAAG-A-AAAAACAGCAUAUUGACA-CCUGGGA--AAGACCA-G 150
MBFV-NC_032088.1 -----NNNNNNNN-UACCGCAAA-ACAAAACAGCAUAUUGACA-CCUGGGA--UAGACUA-G 150
MBFV-NC_012534.1 -----NNNNNNNN-AACACCAA-ACAAAACAGCAUAUUGACA-CCUGGGA--GAGACUA-G 150
MBFV-NC_015843.2 -----NNNNNNNNGAACACCAACA-A-GAAACAGCAUAUUGACA-CCUGGGA--UAGACUA-G 150
MBFV-NC_007580.2 -----NNNNNNNNCCGUUAACGC-AAACAACAGCAUAUUGACA-CCU-GGA--AAGACAG-G 150
MBFV-NC_009028.2 -----NNNNNNNNAACGCGCAA-ACAAAACAGCAUAUUGACG-CUGGGA--AAGACCG-G 150
MBFV-NC_040776.1 -----NNNNNNNN-AAGCGCAA-GACAACAGCAUAUUGACA-CCUGGGA--AAGACCG-G 150
MBFV-NC_000943.1 -----NNNNNNNNAAAAAGCAUCA--AACAACAGCAUAUUGACA-CCUGGGA--AAGACUA-G 151
MBFV-NC_006551.1 -----NNNNNNNNUAAGCAUCAA--AAAAACAGCAUAUUGACA-CCUGGGAUUAAGACUA-G 150
MBFV-NC_001563.2 -----NNNNNNNNUAAGCAUCAA--AAGAACAGCAUAUUGACA-CCUGGGA--UAGACUA-G 152
MBFV-NC_009942.1 -----NNNNNNNNAACACCAACA--CAAAAACAGCAUAUUGACA-CCUGGGA--UAGACUA-G 150
MBFV-NC_009026.2 -----NNNNNNNN-AGCACCAA-AAGAAACAGCAUAUUGACA-CCUGGGA--AAGACCG-G 150
MBFV-NC_034151.1 -----NNNNNNNN-GAACGCAAA-UAAAAACAGCAUAUUGACA-CCUGGGA--UAGACUGGG 151
MBFV-NC_002031.1 -----NNNNNNNNAGGGAACAAA-UAGUGGGACCAUAUUGACG-CCAGGGA--AAGACCGGA 152
ISFV-NC_016997.1 -----NNNNNNNNC---AAAAUGUAUAGGCUAUUAUUGCACA-CCUAGGA--AAGACUG-G 150
ISFV-NC_027999.1 -----NNNNNNNNCAUUGGAAAUUAGCU--UCAUAUUGACA-CCAGGGA--AAGACC-G 150
MBFV-NC_008719.1 -----NNNNNNNNCC-GAGCAC-UAUAGCGGACCAUAUUGACG-CCUGGGA--AAGACCG-G 150
MBFV-NC_012735.1 -----NNNNNNNNCC-GAGCACUAUAGCGGACCAUAUUGACG-CCAGGGA--AAGACCG-G 151
NKV-NC_005039.1 A---NNNNNNNNUAGCAUGGAUUUGCAUAUUGUUGACUC---AGGGA--AAGACCG-G 150
NKV-NC_004119.1 -----NNNNNNNNUUGCCUCAUCUUGAAUGGGGGGCGCG---UGGAU-AUAUACUCCAG 150
NKV-NC_003635.1 -----NNNNNNNNUUUCCAAAUUCC-CGUUACCGGGGGU---GGGGUGAUUAGCCAUGG 150
20.....130.....140.....150.....160.....170.....1



ISFV-NC_034204.1 AAGACUGUCC-CCACC- 167
ISFV-NC_034017.1 AAGAAUGUCC-CCACC- 167
ISFV-NC_024299.2 CAACAGGCUU-CCACCG 167
ISFV-NC_027817.1 AAGUCGAGAG-UCUCC- 167
ISFV-NC_027819.1 GAUGAAGAU-CCCCA- 167
ISFV-NC_008604.2 AGCAGCAGGU-CUCCA- 167
ISFV-NC_021069.1 AGCAGCAAAU-CUCCAC 167
ISFV-NC_012671.1 AGCAGCAAAU-CUCCAC 167
ISFV-NC_001564.2 UCUUA--ACC-UCCAC 167
ISFV-NC_005064.1 GCUAAAUCUC-CCACCG 167
ISFV-NC_012932.1 UCUA--AGC-CUCCCA 167
TBFV-NC_001809.1 GCGGUUCUU-GUUCUC- 167
TBFV-NC_027709.1 GCGGUUCUU-GUUCUC- 167
TBFV-NC_003690.1 GCGGUUCUU-GUUCUC- 167
TBFV-NC_001672.1 CCUGAGCCA-CCAUCA- 167
TBFV-NC_005062.1 CCUGAGCCA-CCAUCA- 167
TBFV-NC_003687.1 GCGGUUCUU-GUUCUC- 167
MBFV-NC_001437.1 UCAACAUCU-GCUACUA 167
MBFV-NC_002640.1 AGAUCCUGCUGUCUCUA 167
MBFV-NC_001477.1 AGAUCCUGCUGUCUCUA 167
MBFV-NC_001475.2 AGAUCCUGCUGUCUCU 167
MBFV-NC_001474.2 AGAUCCUGCUGUCUCU 167
MBFV-NC_012532.1 AGACUCCAUGAGUUUCC 167
MBFV-NC_035889.1 AGACUCCAUGAGUUUCC 167
MBFV-NC_012533.1 AGACUCUGUUGGUUUU 167
ISFV-NC_017086.1 AGGUACUCGUGAUUUA 167
ISFV-NC_040610.1 AGAAUAACACAAUUC- 167
ISFV-NC_024017.1 AGAAUUG-AUGUUGUU 167
MBFV-NC_009029.2 GGGAUUUGCUGUCUGG 167
MBFV-NC_032088.1 GAGAUCGCGUCUCUGG 167
MBFV-NC_012534.1 GAGACCCUCUGUCUAA 167
MBFV-NC_015843.2 GAGACCCUCUGUCUAA 167
MBFV-NC_007580.2 AGAUCCCUUGUUUUU 167
MBFV-NC_009028.2 GAGAUCUUGCUGUUUC 167
MBFV-NC_040776.1 GAGACCCUCUGCUGUC 167
MBFV-NC_000943.1 GAGAUCUUC-UGCUCUA 167
MBFV-NC_006551.1 GAGAUCUUC-UGCUCUA 167
MBFV-NC_001563.2 GGAUCUUCUGCUGUC 167
MBFV-NC_009942.1 GAGAUCUUCUGCUCUGC 167
MBFV-NC_009026.2 AGAAUUUGCUGCCUGC 167
MBFV-NC_034151.1 AGACCCUCUG-CUUAU 167
MBFV-NC_002031.1 GUG-GUUCUC-UGCUCU 167
ISFV-NC_016997.1 AGAUACCUUUGGUUUU 167
ISFV-NC_027999.1 GAGAAACUUCUGCUUA 167
MBFV-NC_008719.1 AGACACUCCUUGAUUCU 167
MBFV-NC_012735.1 AGACA-CUCCUGAUUC 167
NKV-NC_005039.1 GUGUUUGACGCUUCGG 167
NKV-NC_004119.1 CCGAAGAACACUACAGU 167
NKV-NC_003635.1 UCGACAGAUCACAGUC 167
80.....190.....

