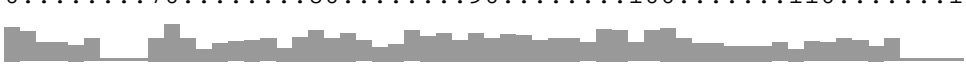


ISFV-NC_008604.2 ----UUGUUUA----UAAUAGUCGUUUUUCUCGCAGAAAUGGGAAAGGACGACGGUAA 51
ISFV-NC_021069.1 ----CCUUUAU----ACCUCUGGUACUAAAAGUGGCACUAUGAAGAAGAAGGA--GUCGG 49
ISFV-NC_012671.1 ----CUUUUAU----ACCUCUGGUACUAAACGUGGCGCUAUGAAGAAGAAGGA--GUCUG 49
ISFV-NC_024299.2 ----UUGAAAA--ACUUUUGCGACGUUGUUGAGAAAGUAGGACGAAAAGAAAGGAGUA 51
ISFV-NC_027819.1 ----UCACGGA--AUAGCUUGAGAAGAAGCUUACUUUAUGUCAAGUAAGGAAGCAGUC 51
ISFV-NC_001564.2 ----GAGAGAC--CGUGAGGUUGAACCCCGGCAAGGAAUGAAGAGGAAGG--AUUUGG 49
ISFV-NC_005064.1 ----AGUUGUU----CGGAAAAGAGACCUUGGUUUCGUUAUGAAUAGGAGGA--GCUUGA 49
ISFV-NC_012932.1 ----AGUUUGU--UGGAAAAGUAUAUCUGGUUUCGUUAUGAAUAGGAGUC--UCUUGA 49
ISFV-NC_034204.1 ----GCGACAG--AAGGAACCCAAAAAGGGAAACCAGAUGGAGCAAAACG--GAAAGU 49
ISFV-NC_034017.1 ----UAAAAAU--UUCUUUGGCCAGCUACCGGAAAGUAUGGAGCAAUUGCA--GGAGU 49
ISFV-NC_027817.1 ----CAAAACU--UGGUGUUGCAACGUUUGUAUUAUAAAUGAGUGGAUUAAGGACCUU 51
NKV-NC_003635.1 ----GUGCGCG--UCACGCCACUUUUUGUGGCGGAAUGUGCGAAAAUAACAGGAAA 51
NKV-NC_004119.1 ----CCGGUCGAAACCACGUCGCUUCUCG--CAAAUGACGAAAGGCAAAAUUGAAA 51
NKV-NC_005039.1 ----CGGAAGAACAACUGUCUGCGGAUCAAAGACGAUGAAA--AAUCG 41
MBFV-NC_002031.1 ----UUGAGCGAUUAGCAGAGAACUGACCAGCAACAUUCU-----GGUCG 41
ISFV-NC_016997.1 ----CAUUAGAAUAGGUUUGAAAGAGUUAGAAUCAGAGAUCAUAUGG--UCUUUCG 50
MBFV-NC_008719.1 ----AAAGGUUAGAGAAAAGGAGUUUUGCUUCUCAUGGCAACAAGAG--GAGCAAG 50
MBFV-NC_012735.1 ----AAGGAGUAGAAGAAAGGAGAUUCAUUTCCCAAUGGCAACAAAGG--GGAUGAA 50
MBFV-NC_012533.1 ----GUGUUGAGAAAGAAAACAGAU--UCUUCGGA----AAUGA-----A--ACCCAAA 43
MBFV-NC_002640.1 ----UGUUUGAAUAGAGAGCAGAUUCUCUGGAAA--AAUG--AA--CCA--ACGAAA 45
MBFV-NC_001477.1 AACAGUUUUUUUAUAGAGAGCAGAUUCUCUG-----AUGAACAA--CCA--ACGAAA 48
MBFV-NC_001475.2 GACAGUUUUUUUAUAGAGAGCAGAUUCUG-----AUGAACAA--CCA--ACGGAAG 48
MBFV-NC_001474.2 ACAGUUUUUUUAUAGAGAGCAGAUUCUCUG-----AUGAAUAA--CCA--ACGAAA 48
MBFV-NC_012532.1 AUUUGGAUUUGGA--AACGAGAGUUUCUGG--U----CAUGAAAAACCCCA--AAGAGA 50
MBFV-NC_035889.1 UUUUGGAUUUGGA--AACGAGAGUUUCUGG--U----CAUGAAAA--CCCA--AAAAAG 49
ISFV-NC_017086.1 ----UGAAUAGGUUUUAUUGUAACGAAAUACCGAUGGCG--AAUA--AACCCAA 47
ISFV-NC_024017.1 ----UGAUAGAGUUGACUAGAUAACGAAUUAUCGAUGGUG--ACUA--AAUCAAAG 47
ISFV-NC_040610.1 ----AAAGAAGUCAAAAACUAGAAAUUUAAUUUCAUGGUU--AGAC--AACUUGG 47
MBFV-NC_009026.2 ----UCGAGGAAA--GACGAGAAGCGAAUUCU--CGAAUGGCU--AGAA--AACCAGG 47
MBFV-NC_009028.2 ----UCUGAAUAGUUUUUAGCAAGCGAUUUC--C--AUGUCU--AAAA--AACCAGG 47
MBFV-NC_040776.1 ----UCUGAAUAGUUUUUAGCAGAGGUACUA--CC--AUGUCU--AAAA--AACCAGG 47
MBFV-NC_006551.1 ----GCCG--GCAGUUUCUUUGAG--CGUUGAUU--UUCAAUGUCU--AAGA--AACCAGG 47
MBFV-NC_001563.2 ----UGCGAGCUGUUUUUCUGGCA--CGAAGA--U--CUGAUGUCU--AAGA--AACCAGG 47
MBFV-NC_009942.1 ----UGCGAGCUGUUUCUAGCA--CGAAGA--U--CUGAUGUCU--AAGA--AACCAGG 47
MBFV-NC_012534.1 ----AGUUUGAACGUUUUUUGGAGAGAGAUUA--UU--AUGUUUAACAAAA--AACCAGG 50
MBFV-NC_015843.2 ----AGUUUGAACGUUUUUUGGAUAGAGACAA--CU--AUGUCUAAACAAAA--AACCAGG 50
MBFV-NC_034151.1 ----UGAAUAGUUUUUUGAACAGAGGUGAA--AACCAUGUCU--AAAA--AACCAGG 47
MBFV-NC_009029.2 ACACAGUGUGAACAGUUUUUUGAACAG--CA----AUGACC--AAAA--AACCAGG 47
MBFV-NC_032088.1 ----AGUGUGAACAGUUUUUUGAACAGUGGAUA--AUGACU--AAAA--AACCAGG 47
MBFV-NC_000943.1 ----UGAACAGUUUUUUGGAGCUUUUGAUU--UCAAAUGUCU--AAAA--AACCAGG 47
MBFV-NC_007580.2 ----GCGAACAGUUUUUU--AGCAGGGAUUACCCAAUGUCU--AAAA--AACCAGG 47
MBFV-NC_001437.1 ----UUAAACAGUUUUUUUUGAACGGAAGAUAA--AUGACU--AAAA--AACCAGG 47
ISFV-NC_027999.1 ----UUGUUUGAGAGAAACAGCGAGAAGUACUUAUGGUUGAGAAAA--CUAGAGU 50
TBFV-NC_001672.1 AAAGACAGCUUAGGAGAAC--AAGAGCUGGGG-----AUGGUCAGAAGGCCAUUCUG 51
TBFV-NC_005062.1 AAAGACAGCUUAGGAGAAC--AAGAGCUGGGG-----AUGGCCGGAAGGCCAUUCUG 51
TBFV-NC_003690.1 AAAGACAGCUUAGGAGAAC--AAGAGCUGGGG-----AUGGCCGGAAGGCCAUUCUG 51
TBFV-NC_027709.1 AAAGACAGCUUAGGAGAAC--AAGAGCUGGGG-----AUGGGCAGGAAGUCCAUUCUG 51
TBFV-NC_001809.1 AAAGACAGCUUAGGAGAAC--AAGAGCUGGGG-----AUGGGCAGGAAGUCCAUUCUG 51
TBFV-NC_003687.1 ----AGCUUAGGAGAAC--AAGAGCUGGGAGUGGUUAUG-----AUGACCACUUCU 45
.....10.....20.....30.....40.....50.....6



ISFV-NC_008604.2 AAGAAGAAGGGGCGGGGCUCCUCCGG-----AUGGUUGUUACCA--CCGGGGCG 98
ISFV-NC_021069.1 GAGUAGUGGAGAGAGGACGCGGUACG-----CGCCCUCCGACCAACCGUGGAGG 98
ISFV-NC_012671.1 GAGUUGUGGAGAGAGGACGUGGAACG-----CGUCCUCCUGCCAUCUGUGGAGG 98
ISFV-NC_024299.2 AAUUAUGGGAGAAGGGAGG--UAUC-----CCGCCCCUCCCAUUAAGGAGAA 98
ISFV-NC_027819.1 AAGAAGAAGGGAAUUUCCUGGGAGG-----AGGAGGAAAACGGGGCUUUGAG-- 99
ISFV-NC_001564.2 AGGCGCGAGGAAAAGCGCCAGGGAGG-----GAUUUUUCCACCCCGUUCUGGGG 98
ISFV-NC_005064.1 AGAAGGGGCGUUAAGCGGGAAGC--UAUC-----AACCGUCCGUGGUUGGACCGCC 98
ISFV-NC_012932.1 AGAAAGGGGCCUUCGGGCCAAACAUCU-----GGUCGGCCAUCGACCAGACCGCC 98
ISFV-NC_034204.1 UUUCCCCGACCGGAGGUGGUCGGGCG-----CGGCCAGAGAGGGGCUUGGGAA 98
ISFV-NC_034017.1 UUUCCCCGCGGAGGCGGUCGGGCG-----GGCCCCAAGAGGGGCGUGCGAA 98
ISFV-NC_027817.1 CUUCCCCUCCGAGGGAAGAA--AAAG-----AAAGCCCCUGUGAUACAUAACA 98
NKV-NC_003635.1 AAGAACCUCGAAAAGGGUUC--AAAG-----AAGAAAAAGAAUGCUGGAAGGAA 98
NKV-NC_004119.1 AACGCGAAUAAAGAAAGC--AAAA-----AACUCUAAAGACAAUCCUGGUCG 98
NKV-NC_005039.1 UAGACAAAACACCCGAAUUUCGG--GUGUCAAUUAAAAAAGCAAAAU--CCAAAAACAA 98
MBFV-NC_002031.1 UAAAGCUCAGGGAAAAAACCCUGGGCGUCAAUUGGUACGACGAGGAGU--UC--GCUCCU 98
ISFV-NC_016997.1 AGCUAGGAGGCCGUUAAAGAGCUGCAUUAUAAUAAAACG--AAAG--CU-- 98
MBFV-NC_008719.1 CAAGUCUCGGGUC--ACGACCCGGGGCGUCAAUUGGUGACGGCCAAG--GC----- 98
MBFV-NC_012735.1 UAAGUCUCGGGCUCAUCCGAGGCGUCAAUUGGUAG--CGGCUUGA--GU----- 98
MBFV-NC_012533.1 AAAG-----GGCCGAAUCGCGGUGUCAAUUGCUGAAGCGCUUUGAUAGCGCUGG 98
MBFV-NC_002640.1 AAGG-----UGGUUAGACC--ACCUUUCAAUUGCUGAAACGCGAGAGAAACCGCGUAUC 98
MBFV-NC_001477.1 AAGA-----CGGGUCGACC--GUCUUUCAUUAUGCUGAAACGCGCGAGAAACCGCGUGU- 100
MBFV-NC_001475.2 AAGA-----CGGGAAGAAC--GUCUAUCAUUAUGCUGAAACGCGGAGAGAAACCGUGU- 100
MBFV-NC_001474.2 AAGG-----CGAAAAACAC--GCCUUUCAUUAUGCUGAAACGCGAGAGAAACCGCGUGU- 100
MBFV-NC_012532.1 AAUC-----CGGAGGA--UCCGAAUUGUCAAUUGCUAAAAACGCGGAGUAGCCCGUG- 100
MBFV-NC_035889.1 AAUC-----CGGGAAGUUCGGAAUUGUCAAUUGCUAAAAACGCGGAGUAGCCCGUG- 100
ISFV-NC_017086.1 GAAGC-----CCGGAAGACGGGCUAUCGAUUAUGUGAGACGUGCGUUGCCUCGCGUCU- 100
ISFV-NC_024017.1 AAAGCC-----CGCCGCAAAGCGGGC--UGUUAAUUAUGCUAACACGCAUCACCAAGCGU--G- 100
ISFV-NC_040610.1 AAAGAC--CGUUGCAAGACGGGUCUGUCAAUUAGUCG--AAAGCGAAUUAACCCUGGUCG- 100
MBFV-NC_009026.2 AAGACC--CGGAGGAAACCGGGUCGUCAAUUAGCUAAAGCGUACGGCGGCCAACG-- 100
MBFV-NC_009028.2 AAAGUC--CGCUGCAAAGCGGACUGUCAAUUAGCUAAAAACGCGUGGCUAGCGUGU-- 100
MBFV-NC_040776.1 AGGACC--CGCCGAAGAGCGGGUUGUCAAUUAGCUAAUAGCGCAUACCCCGUGUU- 100
MBFV-NC_006551.1 AGGGCC--CGGAAGAAACCGGGCCAUAUUAUGCUGAAACGCGGCAUACCCCGCG- 100
MBFV-NC_001563.2 AGGGCC--CGGUAAAAACCGGGCUGUCAAUUAGCUAAAAACGCGGUAUGCCCGCG- 100
MBFV-NC_009942.1 AGGGCC--CGGCAAGAGCCGGGUCGUCAAUUAGCUAAAAACGCGGAUAGCCCGCG- 100
MBFV-NC_012534.1 AAAACC--CGGGGCAACCCGGGUUGUCAAUUAGCUAAAGCGCGGUGCAUCCC- 100
MBFV-NC_015843.2 AAGACC--CGGCUCAGGCGGGGUUGUCAAUUAGCUAAAGCGCGGAACGUCCC- 100
MBFV-NC_034151.1 AGGACC--CGGAAGAAACCGGGUCGUCAAUUAGCUAAUAGCGCAUACCCCGCUA- 100
MBFV-NC_009029.2 AAGGCC--CGGUAGAAAUCCGGGCGUCAAUUAGCUAAAGCGCGGUGCGUCCCGCG- 100
MBFV-NC_032088.1 AAAGUC--CGGCAUUCGUCGGACUGUCAAUUAGCUAAAGCGCGGAGCAUCCCGCG- 100
MBFV-NC_000943.1 AGGACC--CGGAAGCCCGGGUCGUCAAUUAGCUAAUAGCGCAUACCCCGCG- 100
MBFV-NC_007580.2 AAAACC--CGGUAGAAACCGGGUUGUCAAUUAGCUAAAGCGCGGUGCAUCCC- 100
MBFV-NC_001437.1 AGGGCC--CGGUAAAAACCGGGCUAUAUUAUGCUGAAACGCGGCGUACCCCGCG- 100
ISFV-NC_027999.1 AGGAUCUCGGGUAAAAACCGGAGGUGUCAAUUAGUUAUAGGCGUACUGGAA- 100
TBFV-NC_001672.1 AAAGGUAAGGGGGCGGUCUCCCCUCGACGAGUGUCGAAAGAGACCCGCA- 100
TBFV-NC_005062.1 AAAGGAAAGGGGGCGGUCUCCCCUCGACGAGUGUCGAAAGAGAGACCCGCA- 100
TBFV-NC_003690.1 AAAGGAAAGGGGGCGGUCUCCCCUCGACGAGUGUCGAAAGAGAGACCCGCA- 100
TBFV-NC_027709.1 AAAGGUAAGGGGGCGGUCUCCCCUCGACGAGUGUCGAAAGAGACCCGCA- 100
TBFV-NC_001809.1 AAAGGCAAGGGGGCGGUCUCCCCUCGACGAGUGUCGAAAGAGAGACCCGCA- 100
TBFV-NC_003687.1 AAAGGAAAGGGGGCGGUCUCCCCUAGGCGCUAAAGAGUAGCCGCAAAUAGU- 100
0.....70.....80.....90.....100.....110.....1



ISFV-NC_008604.2 CGNNNNNNNN--CUGACGCCACCUCUCCG--AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCA 156
ISFV-NC_021069.1 AANNNNNNNN--GUGACGCCACCCCCCG--AAUUGUGUAUUGUGACAG--GCUUGCUCGCAGCA 155
ISFV-NC_012671.1 AGNNNNNNNN--GUGACGCCACCCCCCG--AAUUGUGUAUUGUGACAG--GCUUGCUCGCAGCA 155
ISFV-NC_024299.2 AANNNNNNNN----ACGACACCCCCCG--AAUUGUGUAUUAUGCCAUUGUAGUCUAUACA 153
ISFV-NC_027819.1 -ANNNNNNNN--UUAGCCACACCCCCG--AAACGUGAUUUUAUGAUGACAGAAUACAGUAG 156
ISFV-NC_001564.2 AANNNNNNNN--AGCGAACCUCUCCGUUAAUGUGAAAAAGUAUGGGG--AAAGAUCUCAUC 155
ISFV-NC_005064.1 CGNNNNNNNN--CGUACCCCCUCCG--UAAUGUGAAAAAGUAUGGGGAAAAAGAACCCAGC 153
ISFV-NC_012932.1 GGNNNNNNNN--UUGCGAACCCCCUCCG--AAUUGUGAAAAAUUAGGGAAGAAUACCCAU 156
ISFV-NC_034204.1 GANNNNNNNN--ACGACCACCCUCUG--AAUUGUUAUUAUAAACUGUGCGGAAAAAUCCAAA 154
ISFV-NC_034017.1 GANNNNNNNN--GCGACACCCUCUG--UAAUGUUUAUUAAGUUGUGCGGAAAAAUCCAAA 154
ISFV-NC_027817.1 GGNNNNNNNNAGUAGCGUACCCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAAGU 156
NKV-NC_003635.1 UGNNNNNNNNUU--UCCAAAUUCCCGUUAACCGGGGGUGGGGUGAUUAGCCAUGGUCG 153
NKV-NC_004119.1 AANNNNNNNNUUG--CCUACUUGUAAUGGGGGGCGGCGUGGAUAUAUACUCCAGCA 153
NKV-NC_005039.1 AANNNNNNNN--UACGCAUGGAG--U--UUGCAUUAUGUAGCAUACGGGAA--AGACC--GG 150
MBFV-NC_002031.1 GUNNNNNNNN--AGGGAACAAAUAG--UGGGACCAUUAUGACGCCAGGGAA--AGACCCGAGU 154
ISFV-NC_016997.1 GCNNNNNNNN--CAAAAAGUUAUAG--GCU--AUUAUUAUCGACACCUAGGAA--AGACUGGAGA 153
MBFV-NC_008719.1 AANNNNNNNN--UAGCCACACCCGAC--UAUAG--CGG--ACCAUUAUGACGCCUGGGAA--AGACCCGAGA 153
MBFV-NC_012735.1 GANNNNNNNN--UAGCCACACCAUAG--CGG--ACCAUUAUGACGCCAGGGAA--AGACCCGAGA 154
MBFV-NC_012533.1 CCNNNNNNNN--AACCAA--CAUAAAACAGCAUUAUGACACCUUGGAA--AGACCCGAGA 153
MBFV-NC_002640.1 AANNNNNNNN--UAGCAACAAAA--AAAACAGCAUUAUGACG--CUGGGAA--AGACCAGAGA 153
MBFV-NC_001477.1 --NNNNNNNNCC--CGCACAAACA--CAAACAGCAUUAUGACG--CUGGGAG--AGACCAGAGA 153
MBFV-NC_001475.2 --NNNNNNNNCC--CCCGAAAAA--AAAACAGCAUUAUGACG--CUGGGAG--AGACCAGAGA 153
MBFV-NC_001474.2 --NNNNNNNNCC--CCGGAACAA--AAAACAGCAUUAUGACG--CUGGGAA--AGACCAGAGA 153
MBFV-NC_012532.1 --NNNNNNNNCC--CCGGAACAA--AAAACAGCAUUAUGACG--UGGGAA--AGACCAGAGA 153
MBFV-NC_035889.1 --NNNNNNNN--C--CGGAAAAACGCAAAACAGCAUUAUGACG--CUGGGAA--AGACCAGAGA 153
ISFV-NC_017086.1 --NNNNNNNN--CCAAAUAUGA--AGGCAUUAUUGACACUUGGGAA--AGACCAG--AG 152
ISFV-NC_024017.1 --NNNNNNNN--UGCUGAGAGC--AAAAUAGCAUUAUGACACUUGGGAA--GAGACCAG--AG 153
ISFV-NC_040610.1 --NNNNNNNN--UCAUUAUUUAUUAACAGCAUUAUGACACCCAGGG--A--UAGCCGG--AG 153
MBFV-NC_009026.2 --NNNNNNNN--ACGCAAAAAAGA--AACAGCAUUAUGACACCUUGGGA--UAGACCAGAGA 153
MBFV-NC_009028.2 --NNNNNNNN--CAAGCGCGAA--ACA--AACAGCAUUAUGACGCCUGGGA--AAGACCCGGAG 153
MBFV-NC_040776.1 --NNNNNNNN--AAGCGCAAGACA--AACAGCAUUAUGACACCUUGGGA--AAGACCCGGAG 153
MBFV-NC_006551.1 --NNNNNNNN--UAGCAUAAAAA--A--AACAGCAUUAUGACACCUUGGGAUUAAGACUAGGAG 155
MBFV-NC_001563.2 --NNNNNNNN--AAACACCAAAAG--A--AACAGCAUUAUGACACCUUGGGA--UAGACUAGGGG 153
MBFV-NC_009942.1 --NNNNNNNN--AACACCAAAAG--A--AACAGCAUUAUGACACCUUGGGA--UAGACUAGGGAG 153
MBFV-NC_012534.1 --NNNNNNNN--AAC--ACCAAAACA--AACAGCAUUAUGACACCUUGGGA--GAGACUAGGAG 153
MBFV-NC_015843.2 --NNNNNNNN--GAA--CACCACAAGA--AACAGCAUUAUGACACCUUGGGA--UAGACUAGGAG 153
MBFV-NC_034151.1 --NNNNNNNGA--CGCAAAUAAAA--CAGCAUUAUGACACCU--GGGA--UAGACUAGGAG 153
MBFV-NC_009029.2 --NNNNNNNN--AGCGAAACACAA--AACAGCAUUAUGACACCUUGGGA--AAGACUAGGGG 153
MBFV-NC_032088.1 --NNNNNNNN--UACCGCAAAACA--AACAGCAUUAUGACACCUUGGGA--UAGACUAGGAG 153
MBFV-NC_000943.1 --NNNNNNNN--AAGCAUCAAAACAGCAUUAUGACACCUUGGGA--AAGACUAGGAG 154
MBFV-NC_007580.2 --NNNNNNNN--CGGUUAAC--GCAAA--CAACAGCAUUAUGACACCUUG--A--AAGAC--AGGAG 152
MBFV-NC_001437.1 --NNNNNNNN--CAUUAUGA--CACCUGGGGA--UAGACUAGGGAG 136
ISFV-NC_027999.1 --NNNNNNNNCCAUUGGAAAU--UAGCUUCAUUAUGACACCCAGGGAA--AAGACCCGAGA 154
TBFV-NC_001672.1 --NNNNNNNN--U--AUACAAA--UUCCCCCUCGAGAGGGGGGGCGG 141
TBFV-NC_005062.1 --NNNNNNNN--U--AUACAAA--UUCCCCCUCAUCUGAGGGGGGGCGG 141
TBFV-NC_003690.1 --NNNNNNNNGAUUAUUCUCCUCCU--AUACAAA--UUCCCCCUCGACAGGGGGGGCGG 156
TBFV-NC_027709.1 --NNNNNNNN--UUAUUCUCCUCCU--AUACAAA--UUCCCCCUCGACAGGGGGGGCGG 156
TBFV-NC_001809.1 --NNNNNNNN--UUUUUCCUCCUCCU--AUACAAA--UUCCCCCUCGACAGGGGGGGCGG 156
TBFV-NC_003687.1 --NNNNNNNN--AUUUUCCGCCUCCU--AUACAAA--UUCCCCCAGGAAACUGGGGGGGCGG 156
20.....130.....140.....150.....160.....170.....1



ISFV-NC_008604.2 GCAGGUCUCCA----- 167
ISFV-NC_021069.1 GCAAAUCUCCAC----- 167
ISFV-NC_012671.1 GCAAAUCUCCAC----- 167
ISFV-NC_024299.2 ACAGGCUUCCACC-----G 167
ISFV-NC_027819.1 GUAAGAUCCCA----- 167
ISFV-NC_001564.2 UAAACCUCCAC----- 167
ISFV-NC_005064.1 UAAAUUCCUCCACC-----G 167
ISFV-NC_012932.1 UAAGCCUCCCA----- 167
ISFV-NC_034204.1 GACUGUCCCCACC----- 167
ISFV-NC_034017.1 GAAUGUCCCCACC----- 167
ISFV-NC_027817.1 CGAGAGUCUC-----C 167
NKV-NC_003635.1 CACAGAUCAAGCU-----C 167
NKV-NC_004119.1 GAAAAGACUCAGA-----U 167
NKV-NC_005039.1 GUGUUUGUCGGUUC-----GG 167
MBFV-NC_002031.1 GUUCUCUCUGUUU----- 167
ISFV-NC_016997.1 UACCUUUGGUUCU----- 167
MBFV-NC_008719.1 CACUCCUUGAUUC----- 167
MBFV-NC_012735.1 CACUCCUUGAUUC----- 167
MBFV-NC_012533.1 CUCUGUUGGUUUC----- 167
MBFV-NC_002640.1 UCCUGCUGUCUCUG----- 167
MBFV-NC_001477.1 UCCUGCUGUCUCUA----- 167
MBFV-NC_001475.2 UCCUGCUGUCUCU----- 167
MBFV-NC_001474.2 UCCUGCUGUCUCCU----- 167
MBFV-NC_012532.1 CUCCAUGAGUUUC----- 167
MBFV-NC_035889.1 CUCCAUGAGUUUC----- 167
ISFV-NC_017086.1 GUACUGCGUGAUUCA----- 167
ISFV-NC_024017.1 AUAUGAUGUUGUU----- 167
ISFV-NC_040610.1 AAAUACAC--AAUUUC----- 167
MBFV-NC_009026.2 AUUUGCUGCCUCG--C----- 167
MBFV-NC_009028.2 AUCCUCUGCUUUC--C----- 167
MBFV-NC_040776.1 ACCCUCUGCUCUG--C----- 167
MBFV-NC_006551.1 AUCUUCUGCUCU----- 167
MBFV-NC_001563.2 AUCUUCUGCUCUG--C----- 167
MBFV-NC_009942.1 AUUUUCUGCUCUG--C----- 167
MBFV-NC_012534.1 ACCCUCUGUCCUA--A----- 167
MBFV-NC_015843.2 ACCCUCUGUCCUA--A----- 167
MBFV-NC_034151.1 ACCCUCUGUCCUA--U----- 167
MBFV-NC_009029.2 AUUUGCUGCUCUG--G----- 167
MBFV-NC_032088.1 AUCCUGCUGCUCUG--G----- 167
MBFV-NC_000943.1 AUCCUCUGCUCUA----- 167
MBFV-NC_007580.2 AUCCCCUGCUUUUC----- 167
MBFV-NC_001437.1 AUCUUCUGCUCUAUCUCAACAUCAGCUACUA 167
ISFV-NC_027999.1 AACUUCUUGCUUA----- 167
TBFV-NC_001672.1 UUCUUGUUCUCCUG-----AGCCACCAUCA 167
TBFV-NC_005062.1 UUCUUGUUCUCCUG-----AGCCACCAUCA 167
TBFV-NC_003690.1 UUCUUGUUCUC----- 167
TBFV-NC_027709.1 UUCUUGUUCUC----- 167
TBFV-NC_001809.1 UUCUUGUUCUC----- 167
TBFV-NC_003687.1 UUCUUGUUCUC----- 167
80.....190.....200.....

