

ISFV-NC_008604.2 U--UGUUUAUAAUAGUCGUUUUUUCUCGCAGAA-----AUGGGAAAGGACGACGGUAGG 52
ISFV-NC_021069.1 C--CUUAUAUACCCUCUGGUACUAAAAGUGGCACU-----AUGAAGAAGAAGGA--GUCGGG 50
ISFV-NC_012671.1 C--UUUAUUUACCCUCUGGUACUAAACGUGGCGCU-----AUGAAGAAGAAGGA--GUCUGG 50
ISFV-NC_024299.2 U--UGAAAAACUUUUUGCGACGUUGUUGAGAAG-----AUGGAGCGAAAGAAAGGAGUAA 52
ISFV-NC_027819.1 U--CACGGAAUAGCUUUGAGAAGAAGCUUACUU-----AUGUCAAGUAAGGAAGCAGUAA 52
ISFV-NC_001564.2 G--AGAGACCGUGAGGUUGAACCCTGGCAAGGA-----AUGAAGAGGAAGG--AUUUGGA 50
ISFV-NC_005064.1 A--GUUGUUUCGGAAAAAGAGACUUGGUUCGU-----AUGAAUAGAGGAA--GCUUGAA 50
ISFV-NC_012932.1 A--GUUGUUUCGGAAAAAGAGAUUUCUGGUUCGU-----AUGAAUAGAGG--GCUUGAA 50
ISFV-NC_034204.1 G--CGACAGAAGGAACCCAAAAAGGGAACCG-----AUGGAGCAAAACG--GAAAGUU 50
ISFV-NC_034017.1 U--AAAAAUUUUUUUUGGCGACGUACGGAAAGU-----AUGGAGCAAUCGA--GGAGAUU 50
ISFV-NC_027817.1 C--AAACUCUGGUGUCCACGCUUUUGUAUUA-----AUGAGUGGAUUAGGAGCCUUC 52
TBFV-NC_003687.1 -----AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG-----AUGACCACUUCUA 46
TBFV-NC_003690.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCGGGAAGGCCGUUCUAA 52
TBFV-NC_027709.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCAGGAAGUCCAUCUGA 52
TBFV-NC_001809.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCAGGAAGACCAUCCUGA 52
TBFV-NC_001672.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCAGGAAGGCCAUCCUGA 52
TBFV-NC_005062.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCGGGAAGGCCAUUCUGA 52
MBFV-NC_001563.2 -----UGCGAGCUGUUUCUUGGCAC--GAAGA--UCUCGAUGUCU--AAGAAACCAGGAG 49
MBFV-NC_009942.1 -----UGCGAGCUGUUUCUUGAGC--GAAGA--UCUCGAUGUCU--AAGAAACCAGGAG 49
MBFV-NC_006551.1 -----GCCG--GCAGUUUCUUUGAGC--GUUGAUUUUCAAUGUCU--AAGAAACCAGGAG 49
MBFV-NC_000943.1 -----GCCC--GAACAGUUUUUUGGAGCUUUUGAUUUUCAAUGUCU--AAAAACCAGGAG 49
MBFV-NC_009029.2 ACACAGUGUGAAACAAGUUUUUGAACAG-----CAAUGACC-----AAAAACCAGGAA 49
MBFV-NC_032088.1 -----AGUGUGAAACAAGUUUUUGAUCAGUGGA-----UAAUGACU--AAAAACCAGGAA 49
MBFV-NC_007580.2 -----GCGAACAGUUUUUUUAGCAGGGAAU--UACCCAAUGUCU--AAAAACCAGGAA 49
MBFV-NC_012534.1 -----AGUUUGAACGUUUUUUGGAGAGAGAUAAU-----AUGUUUAACAAAAACCAGGAA 52
MBFV-NC_015843.2 -----AGUUUGAACGUUUUUUGGAUAGAGAACU-----AUGUCUAACAAAAACCAGGAA 52
MBFV-NC_009028.2 -----UCUGAAUAGUUUUUU--AGCAAGGGA--UUUCCCAUGUCU--AAAAACCAGGAA 49
MBFV-NC_040776.1 -----UCUGAAUAGUUUUUU--AGCAGAGGU--ACUACCAUGUCU--AAAAACCAGGAG 49
MBFV-NC_034151.1 -----UGAAUAGUUUUUUUGAACAGAGGUGAAACCAUGUCU--AAAAACCAGGAG 49
MBFV-NC_009026.2 UCGAGGAAAGA-----CGAGAAGCGAAUUCUCUCAAUGGCU-----AGAAACCAGGAA 49
ISFV-NC_040610.1 A-----AAGAAAGUCAAAACUAGAAAUUUAAUUUCAUUGGU-----AGACAACUUGGAA 49
ISFV-NC_017086.1 -----UGAAUAGGUUUUAUUGUGAAUCGAAAUACCGAUGCG-----AAUAAACCAAGA 49
MBFV-NC_001474.2 ACAGUUUUUUUA-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_001477.1 AACAGUUUUUUUA-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_001475.2 GACAGUUUUUUUA-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_002640.1 -----UGUUUGA-----AUAGAGAGCAGAUUCUCUGGAAAAU-----A--A--CCAACGAAA 45
MBFV-NC_012533.1 -----GUGUUGAG-----AAAGAAAACAGAU--UCUUCGGAAAUGAAAA--C--CCAA--AAA 45
MBFV-NC_012532.1 -----AUUUGGA-----UUUGGAAACGAGAGUUUCUGGUCUAGAAAA--ACCCCAAAGAAGA 50
MBFV-NC_035889.1 -----UUUUGGA-----UUUGGAAACGAGAGUUUCUGGUCUAGAAAA--A--CCCAAAAAAGA 49
ISFV-NC_024017.1 -----UGAUAGAGUUGACUAGAUAAACGAAUUAUCGAUGGUG-----ACUAAAUCAGAA 49
MBFV-NC_002031.1 -----UUGAGCGAU--UAGCAGAGAACUGACCAGAACAUUGUC-----UGGUCGUA 43
ISFV-NC_027999.1 -----UUGUUUGAGAGAACACAGCGAGAGUACUACU--AUGGUUGAGAAAAACUAGAGUAG 52
MBFV-NC_012735.1 -----AAGGAGUAGAAAGAAAG--GAGAUUCAUUCCAAUGGCAACAAAGGGGAGUAAUA 52
MBFV-NC_008719.1 -----AAAGGUAAGAGAAAAG--GAGUUUUGCUUCUCU--AUGGCAACAAAGAGGAGCAGCA 52
ISFV-NC_016997.1 -----CAUUAAGAAUAGGUUUUGAAAGUUUGAAUC--AAUGGAAUCAAUGGUUUUCCAG 52
MBFV-NC_001437.1 -----UUAAACAGUUUUUUUAGAACCGGAAGAUAAACCAUGACU--AAAAACCAGGAG 49
NKV-NC_005039.1 -----CGGAAGAACACUCUGCCGGAUCAAAGACGAUG-----AAAAUCGUA 43
NKV-NC_004119.1 -----CCGGUGGAAACACGUCGCUUCUGU-----CAAAUGACGAAAGGCAAAUAGAAAA 52
NKV-NC_003635.1 -----GUGCGCG-----UCACGCCACUUUUUGUGCGGGAAUGUGCGAAAAUAACAGGAAA 52
.....10.....20.....30.....40.....50.....6



ISFV-NC_008604.2 AGAAGAAGGGGC-----CGG-----GCUCCUCCGGAUGGUUGUUAACCA--CCGGGGCGCG 100
ISFV-NC_021069.1 AGUAGUGGAGAG-----AGG-----ACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAA 100
ISFV-NC_012671.1 AGUGUGGAGAG-----AGG-----ACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAG 100
ISFV-NC_024299.2 AUUAUGGGAGAA-----GGG-----UAGC-----UAUCCCGCCCCUCCAUUAAGGAGAAAA 100
ISFV-NC_027819.1 AGAAGAAGGGGAA-----UUU-----CCUUGGGAGGAGGAGGAAAAACGGGGCUUUGAG--A 100
ISFV-NC_001564.2 GGCGCGAGGAAA-----AGC-----GCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAA 100
ISFV-NC_005064.1 GAGGGGGCCUCC-----GGG-----CCCAACAUCCAACCGCGGCGUUGGAGCCGCCG 100
ISFV-NC_012932.1 GAAAGGGCCUUC-----GGG-----CCAAACAUUCUGGUCGGCAUCGACCAGACCGCCGG 100
ISFV-NC_034204.1 UUCCCGACCCGG-----AGG-----UGGUCGGGCCCCGGCCAGAGAGGGGCUUGGGGAAGA 100
ISFV-NC_034017.1 UUCCCGGCCGG-----AGG-----CGGUCGGGCGGCCCCAAAGAGAGGGGCUUGGCGAAGA 100
ISFV-NC_027817.1 UUCCCCUCCGAG-----GGA-----AGAA--AAAGAAAGCCCCUGUGAUACAUCACAGG 100
TBFV-NC_003687.1 AAGGAAAGGGGG-----GCGGUCCCCUAGGCGCAAGCUUAAAGUGACCAGCAAAUAAAGU 100
TBFV-NC_003690.1 AAGGAAAGGGGG-----GGGUCCCCUCCGACGAGCGUCCGAAAGUGGGCCUUGGCGAAGA 100
TBFV-NC_027709.1 AAGGUAAGGGGG-----GCGGUCCCCUCCGACGAGUGUCGAAAGAGACCGCAA----- 100
TBFV-NC_001809.1 AAGGCAAGGGGG-----GCGGUCCCCUCCGACGAGUGUCGAAAGAGACCGCGA----- 100
TBFV-NC_001672.1 AAGGUAAAGGGGG-----GCGGUCCCCUCCGACGAGUGUCGAAAGAGACCGCAA----- 100
TBFV-NC_005062.1 AAGGAAAGGGGG-----GCGGUCCCCUCCGACGAGUGUCGAAAGAGACCGCUA----- 100
MBFV-NC_001563.2 GGCCC---GGUAAAAACCGGGCUGUCAAUUAGCUAAAAACCGGGUAUGCCCCGCG----- 100
MBFV-NC_009942.1 GGCCC---GGCAAGAGCCGGGCGUGUCAAUUAGCUAAAAACCGCGGUAUGCCCCGCG----- 100
MBFV-NC_006551.1 GGCCC---GGAAGAAACCGGGCCAUCAAUUAGCUAAACCGCGGCAUACCCCGCG----- 100
MBFV-NC_000943.1 GACCC---GGGAAGCCCCGGGUCGUCAAUUAGCUAAAAACCGCGGCAUACCCCGCG----- 100
MBFV-NC_009029.2 GGCCC---GGUGAAAAUCGGGCGGUGUCAAUUAGCUAAAGCGCGGUGGCGUCCCGCG----- 100
MBFV-NC_032088.1 AGUCC---GGCAUCGUCGGACUGUCAAUUAGCUAAAGCGCGGAGCAUCCCGCG----- 100
MBFV-NC_007580.2 AACCC---GGUAGAAACCGGGUUGUCAAUUAGCUAAAAACCGCGGCUAUCCCCGCG----- 100
MBFV-NC_012534.1 AACCC---GGGCAACCCCGGUGUCAAUUAGCUAAAGCGCGGUGUACCCCGCG----- 100
MBFV-NC_015843.2 GACCC---GGCUACAGGCCGGGUUGUCAAUUAGCUAAAGCGCGGGAACGUCCC----- 100
MBFV-NC_009028.2 AGUCC---GCUGCAAAGCGGACUGUCAAUUAGCUAAAAACGCCUGGCUAGCGUGU----- 100
MBFV-NC_040776.1 GACCC---GCGGCAAGGCGGUGUCAAUUAGCUAAAGCGCGGGAUACCGCGUU----- 100
MBFV-NC_034151.1 GACCC---GGAAAGAACCGGGUCGUUAAUAGCUAAGCGGCACAUCCCGCGCUA----- 100
MBFV-NC_009026.2 GACCC---GGAGGAAACCGGGUCGUCAAUUAGCUAAAGCGUACGGCGGCAACG----- 100
ISFV-NC_040610.1 AGACC---GUUGCAAGAGCGGUGUCAAUUAGCUAAAGCGGGAUACCCCGCU----- 100
ISFV-NC_017086.1 AGCCC---GG---AAGACGGGCUAUCGAUUAUGUGAGACGUGCGUUGCCUCGCGUCU--- 100
MBFV-NC_001474.2 AAGGC---G---AAAAACACGCUUUUCAAUUAGCUGAAACCGCGAGAGAAACCGCGUGU--- 100
MBFV-NC_001477.1 AAGAC---G---GUCGACCCGUUUUCAAUUAGCUGAAACCGCGAGAGAAACCGCGUGU--- 100
MBFV-NC_001475.2 AAGAC---G---GGAAACCGUCUAUCAAUUAGCUGAAACCGCGUGAGAGAAACCGUGUGU--- 100
MBFV-NC_002640.1 AAGGU---G---GUUAGACCCACCUUUCAAUUAGCUGAAACCGCGAGAGAAACCGCGUAUCAA 100
MBFV-NC_012533.1 AGGGC---C---GCAUCGCGCCGUGUCAAUUAGCUGAAAGCGCGUUGGUAUCUAGCGCUGGCC 100
MBFV-NC_012532.1 AAUCC---GGAGGA--UCCGGAUUGUCAAUUAGCUAAACCGCGGAGUAGCCCGUG----- 100
MBFV-NC_035889.1 AAUCC---GGAGGAUUCGGAUUGUCAAUUAGCUAAACCGCGGAGUAGCCCGUG----- 100
ISFV-NC_024017.1 AGCCC---GCGGCAAGCGGGCGUGUAAUAGCUAAACCGCAACGCAUACCAAGCGUG----- 100
MBFV-NC_002031.1 AAGCUCAGGGAAAAACCCUGGGCGUCAAUUAGGUACGACGAGGAGUUCGCUCCUUGU--- 100
ISFV-NC_027999.1 GAUCUCGGGUUAAAAACCGGAGGUGUCAAUUAGUUAUAGGCGUACUG-----GAA--- 100
MBFV-NC_012735.1 AGUCUCGGGUCGUAUCCCGAGGCGUGUCAAUUAGGUAGCGGCUAGAG-----UGA--- 100
MBFV-NC_008719.1 AGUCUCGGGUCACGACCCGGGGCGUCAAUUAGGUGACGGCCAAGG-----CCA--- 100
ISFV-NC_016997.1 CUAGGAGGCCGUUAAACCGGGCGGUGUCAAUUAAUAAACGUAAG-----CUGC--- 100
MBFV-NC_001437.1 GGCCC---GGUAAAAACCGGGCUAUCGAUUAUGCUAAAGCGGCUUACCCCGCG----- 100
GACAAAACACCCGAAUUCGCGGUGUCAAUUAAAAAAGCAAAAUCCAAAAAACAAAA--- 100
NKV-NC_005039.1 ACGCGAAUAAAAAG-----AAAGCAAAAAACUCUAAAGACCAUUCUGGUCGAA 100
NKV-NC_004119.1 AGAACCUCGGAAG-----GGUCAAAGAGAAAGAAAGUAGCUGGAAGGAAUG 100
NKV-NC_003635.1 0.....70.....80.....90.....100.....110.....1
.....



ISFV-NC_008604.2 NNNNNNNN-CUGACGCCACCUCGCCG-AAAUGUGUAUAGUAAACAGAGCAUGCCUGCAGCAGC 158
ISFV-NC_021069.1 NNNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGC 157
ISFV-NC_012671.1 NNNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAGC 157
ISFV-NC_024299.2 NNNNNNNN-----ACGACACACCCCCCG-AAAUGUGUAUUAUGCCCAUGUAGUUAUACAAC 155
ISFV-NC_027819.1 NNNNNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUUAUAGUGACAGAAUACAGUAGGA 158
ISFV-NC_001564.2 NNNNNNNN--AGCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAUCUAUCUU 157
ISFV-NC_005064.1 NNNNNNNN-----CGUACCCCUCCUG-AAAUGUGAAAAAGUAUGGGGAAAAAGAACCCAGCUA 155
ISFV-NC_012932.1 NNNNNNNN-UUGCCAAACCCCCCG-AAAGUGUAAAAAUUAGGGGAAAAAGUACCCAGCUA 158
ISFV-NC_034204.1 NNNNNNNN--ACGACCACCCUCUG-AAAUGUUAUAAAAACUGUGCGGAAAAAUCCAAAGA 156
ISFV-NC_034017.1 NNNNNNNN-----GCGACACCCUCUG-UAAUGUUUAUUAAGUUGCGGAAAAAUCCAAAGA 156
ISFV-NC_027817.1 NNNNNNNN-NAGUAGCGUACCCUCCG-UGUGAAACAGAUUGACGCGGUAAGACUAAAGCUG 158
TBFV-NC_003687.1 NNNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUCCCCCAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_003690.1 NNNNNNNNNGAUUUUCCUCCUCCUAUACCAA--UUCCCCUCGUCAGAGGGGGGGCGGUUC 159
TBFV-NC_027709.1 NNNNNNNNNGAUUUUCCUCCUCCUAUACCAA--UUCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_001809.1 NNNNNNNNG-UUUUUUCCUCCUCCUAUACCAA--UUCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_001672.1 NNNNNNNN-----UAUACAAA--UUCCCCUCGUGAGAGGGGGGGCGGUUC 144
TBFV-NC_005062.1 NNNNNNNN-----UAUACAAA--UUCCCCUCUACUGAGAGGGGGGGCGGUUC 144
MBFV-NC_001563.2 NNNNNNNNNAACAC--CA-AAAGAAACAGCAUUAUGACACCUGGGA--UAGACUAGGGGAUC 156
MBFV-NC_009942.1 NNNNNNNNNAACACC--AC-AACAAAACAGCAUUAUGACACCUGGGA--UAGACUAGGGAGUC 156
MBFV-NC_006551.1 NNNNNNNNNAUAGCA--UC-AAAAAACAGCAUUAUGACACCUGGGAUUAUGACUAAAGAU 158
MBFV-NC_000943.1 NNNNNNNNNAAAAGC-AU-CAAAACACAGCAUUAUGACACCUGGGA--AGACUAGGAGAU 157
MBFV-NC_009029.2 NNNNNNNNAGCGCA-AA-CACAAAACAGCAUUAUGACACCUGGGA--AAGACUAGGGAGU 156
MBFV-NC_032088.1 NNNNNNNNNAACGCA-AA-AACAAAACAGCAUUAUGACACCUGGGA--UAGACUAGGGAGU 156
MBFV-NC_007580.2 NNNNNNNNCCGUUA-ACGCAAAACACAGCAUUAUGACACCUGGAA--AGAC--AGGAGAU 155
MBFV-NC_012534.1 NNNNNNNNNAACACC-AA-AACAAAACAGCAUUAUGACACCUGGGA--GAGACUAGGGAGC 156
MBFV-NC_015843.2 NNNNNNNNNAACAC--CA-CAAGAAACAGCAUUAUGACACCUGGGA--UAGACUAGGGAGC 156
MBFV-NC_009028.2 NNNNNNNNCAAGCG--CGAA-ACAAACAGCAUUAUGACGCCUGGGA--AAGACCGGGAGAU 156
MBFV-NC_040776.1 NNNNNNNN--AAGCG--CAAAAGAAACAGCAUUAUGACACCUGGGA--AAGACCGGGAGC 156
MBFV-NC_034151.1 NNNNNNNNNGAA--CG-CAAAUAAAAACAGCAUUAUGACACCUGGGA--UAGACUGGGAGAU 156
MBFV-NC_009026.2 NNNNNNNN--ACGCA--CCAAAGAAACAGCAUUAUGACACCUGGGA--UAGACCGGAGAAU 156
ISFV-NC_040610.1 NNNNNNNNCUCAU--AUUUUAUUAACAGCAUUAUGACACCAGGGA--UAGACCGGAGAAA-- 156
ISFV-NC_017086.1 NNNNNNNNC-CCAA-AAUUGAAGGCAUUAUUGACACCUGGGA--AAGACCAG--AGUA 155
MBFV-NC_001474.2 NNNNNNNNCCCCCG-AAACAAAAACAGCAUUAUGACG--CUGGGA--AAGACCAG--AGAU 155
MBFV-NC_001477.1 NNNNNNNNCCCCCG-AAACAAAAACAGCAUUAUGACG--CUGGGA--GAGACCAG--AGAU 155
MBFV-NC_001475.2 NNNNNNNNCCCCCG-CAAAUAAAAACAGCAUUAUGACG--CUGGGA--GAGACCAG--AGAU 155
MBFV-NC_002640.1 NNNNNNNNCCCCCG-AAACAAAAACAGCAUUAUGACG--CUGGGA--AAGACCAG--AGAU 155
MBFV-NC_012533.1 NNNNNNNNNAUAAC--CAACAAAAACAGCAUUAUGACACCUGGGA--AAGACCGG--AGAU 155
MBFV-NC_012532.1 NNNNNNNNCCCGGAAACGCAAAACAGCAUUAUGACG--UGGGA--AAGACCAG--AGAU 155
MBFV-NC_035889.1 NNNNNNNN-CCCGGAAACGCAAAACAGCAUUAUGACG--CUGGGA--AAGACCAG--AGAU 155
ISFV-NC_024017.1 NNNNNNNNCUGCUG--AGAGCAAAUAGCAUUAUGACACCUGGGA--GAGACCAG--AGAU 156
MBFV-NC_002031.1 NNNNNNNNA-GGGACAAAUAGUGGGACCAUUAUGACGCCAGGGA-AAGACC-GGAGUGG 156
ISFV-NC_027999.1 NNNNNNNNC-C-AUUGGAAAUUAGCUUCAUUAUGACACCAGGGA-AAGACC-GGAGAAA 156
MBFV-NC_012735.1 NNNNNNNNC-CGAGCACACUAGCGGACCAUUAUGACGCCAGGGA-AAGACC-GGAGACA 156
MBFV-NC_008719.1 NNNNNNNNC-CGAGCAC-UUAGCGGACCAUUAUGACGCCAGGGA-AAGACC-GGAGACA 155
ISFV-NC_016997.1 NNNNNNNN--CAAAAAUGUAUAGGCUAUAUUAUCGACACCUAGG-A-AAGACU--GGAGUA 155
MBFV-NC_001437.1 NNNNNNNN-----CAUAUUGA-----CACCUGGGA--AAGACCGGGAGU 139
NKV-NC_005039.1 NNNNNNNNUACGC--AUGGAUUUUGCAUAUUGUUGACUACAGGGA--AAGACCGGGUGU 156
NKV-NC_004119.1 NNNNNNNNUUGCCUCAUCUUGAAUGGGGGGCGGCGUGGAUUAUUAUCUCCAGCCAGAAAAA 160
NKV-NC_003635.1 NNNNNNNNUUUUCCAAAUUCCCGUUAACCGGGGUGGGGUGAUUAGCCAUGGUCGCACAGAU 160
20.....130.....140.....150.....160.....170.....1



ISFV-NC_008604.2 AGGUCUCC-----A--- 167
ISFV-NC_021069.1 AAAAUCUCC-----AC-- 167
ISFV-NC_012671.1 AAAAUCUCC-----AC-- 167
ISFV-NC_024299.2 AGGCUUCC-----ACCG 167
ISFV-NC_027819.1 AGAUCCCC-----A--- 167
ISFV-NC_001564.2 AACCUCUCC-----AC-- 167
ISFV-NC_005064.1 AAUCUCC-----ACCG 167
ISFV-NC_012932.1 AGCCUCC-----A--- 167
ISFV-NC_034204.1 CUGUCC-----ACC 167
ISFV-NC_034017.1 AUGUCC-----ACC 167
ISFV-NC_027817.1 AGAGUC-----C 167
TBFV-NC_003687.1 UUGUUC----- 167
TBFV-NC_003690.1 UUGUUC----- 167
TBFV-NC_027709.1 UUGUUC----- 167
TBFV-NC_001809.1 UUGUUC----- 167
TBFV-NC_001672.1 UUGUUCUCCU-----GAGCCACCAUCA 167
TBFV-NC_005062.1 UUGUUCUCCU-----GAGCCACCAUCA 167
MBFV-NC_001563.2 UUCUGC-UC-----UGC 167
MBFV-NC_009942.1 UUCUGC-UC-----UGC 167
MBFV-NC_006551.1 UUCUGC-UC-----U-- 167
MBFV-NC_000943.1 UUCUGC-UC-----UA- 167
MBFV-NC_009029.2 UUCUGC-UC-----UGG 167
MBFV-NC_032088.1 CGCUGC-UC-----UGG 167
MBFV-NC_007580.2 CCCUGCUU-----UUC 167
MBFV-NC_012534.1 CUCUGU-CC-----UAA 167
MBFV-NC_015843.2 CUCUGU-CC-----UAA 167
MBFV-NC_009028.2 CUCUGC-UU-----UCC 167
MBFV-NC_040776.1 CUCUGC-UC-----UGC 167
MBFV-NC_034151.1 CUCUGC-UU-----UCC 167
MBFV-NC_009026.2 UGCUGCCU-----GC 167
ISFV-NC_040610.1 -UACACAAU-----UUC 167
ISFV-NC_017086.1 CUCGCUAG-----UCA 167
MBFV-NC_001474.2 CUCGCUAG-----CCU 167
MBFV-NC_001477.1 CUCGUGUC-----CUA 167
MBFV-NC_001475.2 CUCGUGUC-----CCU 167
MBFV-NC_002640.1 CUCUGC-UC-----CUG 167
MBFV-NC_012533.1 CUGUUGGU-----UUC 167
MBFV-NC_012532.1 CCAUGAGUU-----UCC 167
MBFV-NC_035889.1 CCAUGAGUU-----UCC 167
ISFV-NC_024017.1 UGAUGU-UG-----UUU 167
MBFV-NC_002031.1 UUCUCUGC-----U-U 167
ISFV-NC_027999.1 CUUCUUGC-----U-A 167
MBFV-NC_012735.1 CUUCUUGAU-----UC- 167
MBFV-NC_008719.1 CUUCUUGAU-----UCU 167
ISFV-NC_016997.1 CCUUUGGU-----CUU 167
MBFV-NC_001437.1 UUCUGGCUAUCUCAACAUACGACUACUA 167
NKV-NC_005039.1 GUCGUUCC-----AGG 167
NKV-NC_004119.1 CAA-----AGU 167
NKV-NC_003635.1 CAA-----GCUC 167
80.....190.....200.....

