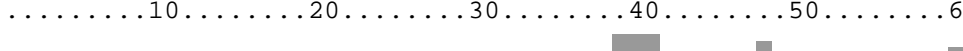


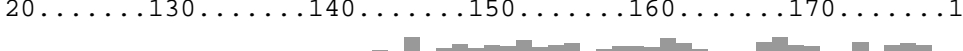
ISFV-NC_008604.2 U--UGUUUAUAAUAGUCGUUUUUUCUCGCAGAA-----AUGGGAAAGGACGACGGUAGG 52
ISFV-NC_021069.1 C--CUUUAUAAUCCUCUGGUACUAAAGUGGCGACU-----AUGAAGAAGAAGGA--GUCGGG 50
ISFV-NC_012671.1 C--UUUAUUUACCUCUGGUACUAACGUGGGCGCU-----AUGAAGAAGAAGGA--GUCUGG 50
ISFV-NC_024299.2 U--UGAAAAACUUUUUGCGACGUUGUUGAGAAG-----AUGGAGCGAAAGAAAGGAGUAA 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-----AUGAAUAGAGGAA--GCUUGAA 50
ISFV-NC_034204.1 G--CGACAGAAGGAACCCAAAAAGGGAACCG-----AUGGAGCAAAACG--GAAAGUU 50
ISFV-NC_034017.1 U--AAAAAUUUUUUUUGGCGACGUACGGAAAGU-----AUGGAGCAAUUGA--GGAGAUU 50
ISFV-NC_027817.1 C--AAACUCUGGUGUCCACGCUUUUGUAUUAUAA-----AUGAGUGGAUUAGGAGCCUUC 52
TBFV-NC_003687.1 -----AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG-----AUGACCACUUCUA 46
TBFV-NC_003690.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCGGGAAGGCCGUUCUAA 52
TBFV-NC_027709.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCAGGAAGUCCAUCCUGA 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 ACACAGUGUGAAACAAGUUUUUGAAGCAG-----CAAUGACC-----AAAAACCAGGAA 49
MBFV-NC_032088.1 -----AGUGUGAAACAAGUUUUUGAUCAGUGGA-----UAAUGACU--AAAAACCAGGAA 49
MBFV-NC_007580.2 -----GCGAACAGUUUUUUUAGCAGGGAAU--UACCCAAUGUCU--AAAAACCAGGAA 49
MBFV-NC_012534.1 -----AGUUUGAACGUUUUUUUGGAGAGAGAUAAU-----AUGUUUAACAAAAACCAGGAA 52
MBFV-NC_015843.2 -----AGUUUGAACGUUUUUUUGGAUAGAGAACU-----AUGUCUAACAAAAACCAGGAA 52
MBFV-NC_009028.2 -----UCUGAAUAGUUUUUUU--AGCAAGGGA--UUUCCCAUGUCU--AAAAACCAGGAA 49
MBFV-NC_040776.1 -----UCUGAAUAGUUUUUUU--AGCAGAGGU--ACUACCAUGUCU--AAAAACCAGGAG 49
MBFV-NC_034151.1 -----UGAAUAGUUUUUUUGAACAGAGGUGAAAAACCAUGUCU--AAAAACCAGGAG 49
MBFV-NC_009026.2 UCGAGGAAAGA-----CGAGAAGCGAAUUCUCUCAAUGGCU-----AGAAACCAGGAA 49
ISFV-NC_040610.1 A-----AAGAAAGUCAAAACUAGAAAUUUAAUUUCAUUGGU-----AGACAACUUGGAA 49
ISFV-NC_017086.1 -----UGAAUAGGUUUUAUUGUGAAUCCGAAUACCGAUGGCG-----AAUAAACCAAGA 49
MBFV-NC_001474.2 ACAGUUUUUUUAA-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_001477.1 AACAGUUUUUUU-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_001475.2 GACAGUUUUUUU-----UUAGAGAGCAGAUUCUCG-----AUGAAU--A--CCAACGAAA 48
MBFV-NC_002640.1 -----UGUUUGA-----AUAGAGAGCAGAUUCUCUGGAAAAAUG-----A--A--CCAACGAAA 45
MBFV-NC_012533.1 -----GUGUUGAG-----AAAGAAAACAGAU--UCUUCGGAUUGAAAA--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--GAGUUUUGCUUCUCAAUGGCAACAAAGAGGAGCAGCA 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 -----CCGGUGGAAACACAGUCGCUUCUGU-----CAAAUGACGAAAGGCAAAUAGAAAA 52
NKV-NC_003635.1 -----GUGCGCG-----UCACGCCACUUUUUGUGCGGGAAUGUGCGAAAAUAACAGGAAA 52
.....10.....20.....30.....40.....50.....6



ISFV-NC_008604.2 AGAAGAAGGGGC-----CGG-----GCUCCUCCGGAUGGUUGUUACCA--CCGGGGCGCG 100
ISFV-NC_021069.1 AGUAGUGGAGAG-----AGG-----ACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAA 100
ISFV-NC_012671.1 AGUGUGGAGAG-----AGG-----ACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAG 100
ISFV-NC_024299.2 AUUAUGGGAGAA-----GGG-----AAGC--UAUCCCGCCCCUCCAUUAAGGAGAAAA 100
ISFV-NC_027819.1 AGAAGAAGGGGAA-----UUU-----CCUUGGGAGGAGGAGGAAAAACGGGGCUUUGAG--A 100
ISFV-NC_001564.2 GGCGCGAGGAAA-----AGC-----GCCAGGGAGGGAUUUUCCACCCCGUUCUGGGGAA 100
ISFV-NC_005064.1 GAAAGGGGCCUCC-----GGG-----CCCAACAUCCAACCGCGGCGUUGGACCGCCG 100
ISFV-NC_012932.1 GAAAGGGGCCUUC-----GGG-----CCAAACAUUCGUGCGGCCAUCCAGACCCGCCG 100
ISFV-NC_034204.1 UUCCCGACCCGG-----AGG-----UGGUCGGGCGCCCGCCAGAGAGGGGCUUGGGGAAGA 100
ISFV-NC_034017.1 UUCCCGCGCCGG-----AGG-----CGGUCGGGCGGCCCCAAAGAGAGGGGCUUGGCGAAGA 100
ISFV-NC_027817.1 UUCCCCUCCGAG-----GGA-----AGAA--AAAGAAAGCCCCUGUGAUACAUCACAGG 100
TBFV-NC_003687.1 AAGGAAAGGGGG-----GCGGUCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUUAAGU 100
TBFV-NC_003690.1 AAGGAAAGGGGG-----GGGUCCCCUCCGACGAGCGUCCGAAAGUGGCGUUGGCCAA 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---GGUAAAAACCGGGCUGUCAAUUAGCUAAAAACCGGUAUGCCCCGCG-----100
MBFV-NC_009942.1 GGCCC---GGCAAGAGCCGGGCGUGUCAAUUAGCUAAAAACCGGCGUAGCCCCGCG-----100
MBFV-NC_006551.1 GGCCC---GGAAGAAACCGGGCCAUCAAUUAGCUAAACCGGCAUACCCCGCG-----100
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MBFV-NC_032088.1 AGUCC---GGCAUCGUGCGGACUGUCAAUUAGCUAAAGCGCGGAGCAUCCCGCG-----100
MBFV-NC_007580.2 AACCC---GGUAGAAACCGGGUUGUCAAUUAGCUAAAAACCGGCGUAUCCCGCG-----100
MBFV-NC_012534.1 AACCC---GGGCAACACCGGGUUGUCAAUUAGCUAAAGCGCGGUGUAUCCCGCG-----100
MBFV-NC_015843.2 GACCC---GGCUACAGGCCGGGUUGUCAAUUAGCUAAAGCGCGGAACGUCCC-----100
MBFV-NC_009028.2 AGUCC---GCUGCAAAGCGGACUGUCAAUUAGCUAAAAACGCCUGGCUAGCGUGU-----100
MBFV-NC_040776.1 GACCC---GCGGCAAGGCGGUGUGUCAAUUAGCUAAAGCGCGGAGUCCAGCGUU-----100
MBFV-NC_034151.1 GACCC---GGAAAGAACCGGGUGCGUAAUUAUGCUAAGCGGCACAUCGCCGCGCUA-----100
MBFV-NC_009026.2 GACCC---GGAGGAAACCGGGUGCGUCAAUUAGCUAAAGCGUACGGCGGCCAACG-----100
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ISFV-NC_017086.1 AGCCC---GG---AAGACGGGCUAUCGAUUAUGUGAGACGUGCGUUGCCUCGCGUCU---100
MBFV-NC_001474.2 AAGGC---G---AAAAACACGCUUUUCAAUUAGCUGAAACCGCGAGAGAAACCGCGUGU---100
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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---GCGGCAAGCGGGGCGUGUAAUUAUGCUAACACGCAUCACCAAGCGUG-----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 CUAGGAGGCCGUUAAACCGGGCGGCGUCAAUUAAUAAACGUAAG-----CUGC---100
MBFV-NC_001437.1 GGCCC---GGUAAAAACCGGGCUAUCGAUUAUGCUAAAGCGGCCUACCCCGCG-----100
NKV-NC_005039.1 GACAAAACACCCGAAUUCGCGGUGUCAAUUAAAAAAGCAAAAUCCAAAAAACAAAA---100
NKV-NC_004119.1 ACGCGAAUAAAAAG-----AAAGCAAAAAACUCUAAAGACCAUUCUGGUCGAA 100
NKV-NC_003635.1 AGAACCUCGGAAG-----GGUCAAAGAGAAAGAAAGAUUGCUGGAAGGAAUG 100
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-----ACGACACCCCCCG-AAAUGUGUAUUAUGCCCAUGUAGUUAUACAAC 155
ISFV-NC_027819.1 NNNNNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUUAUAGUGACAGAAUACAGUAGA 158
ISFV-NC_001564.2 NNNNNNNN--AGCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAUCUAUCUU 157
ISFV-NC_005064.1 NNNNNNNN-----CGUACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAAGAACCCAGCUA 155
ISFV-NC_012932.1 NNNNNNNN-UUGCCAAACCCCCCG-AAAGUGUAAAAAUUAGGGGAAAAAGUACCCAGCUA 158
ISFV-NC_034204.1 NNNNNNNN--ACGACCACCCUCUG-AAAUGUUAUUAUAAACUGUGCGGAAAAAUCCAAAGA 156
ISFV-NC_034017.1 NNNNNNNN-----GCGACACCCUCUG-UAAUGUUUAUUAAGUUGCGGAAAAAUCCAAAGA 156
ISFV-NC_027817.1 NNNNNNNN-NAGUAGCGUCACCCUCCGU--UGUGAAACAGAUUGCGACUAGAAACUAAUCUG 158
TBFV-NC_003687.1 NNNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUCCCCCAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_003690.1 NNNNNNNNNGAUUUUCCUCCUCCUAUACCAA--UUCCCCCUCGUCAGAGGGGGGGCGGUUC 159
TBFV-NC_027709.1 NNNNNNNNNGAUUUUCCUCCUCCUAUACCAA--UUCCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_001809.1 NNNNNNNNG-UUUUUUCCUCCUCCUAUACCAAUCCCCCUCGACAGAGGGGGGGCGGUUC 159
TBFV-NC_001672.1 NNNNNNNN-----UAUACAAA--UUCCCCCUCGUAAGAGGGGGGGCGGUUC 144
TBFV-NC_005062.1 NNNNNNNN-----UAUACAAA--UUCCCCCUCUACUGAGAGGGGGGGCGGUUC 144
MBFV-NC_001563.2 NNNNNNNNAAACAC--CA-AAAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGGAUC 156
MBFV-NC_009942.1 NNNNNNNNAACACC--AC-AACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGAUC 156
MBFV-NC_006551.1 NNNNNNNNUAAGCA--UC-AAAAAACAGCAUAUUGACACCUGGGAUUAAGACUAGGAGU 158
MBFV-NC_000943.1 NNNNNNNNAAAGC--AU-CAAACAACAGCAUAUUGACACCUGGGA--AGACUAGGGAUC 157
MBFV-NC_009029.2 NNNNNNNNAGCGCA--AA-CACAAAACAGCAUAUUGACACCUGGGA--AAGACUAGGGAUC 156
MBFV-NC_032088.1 NNNNNNNNUACGCG--AA-AACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGAUC 156
MBFV-NC_007580.2 NNNNNNNNCCGUUA--ACGCAAAACAACAGCAUAUUGACACCUGGAA--AGAC--AGGAGUUC 155
MBFV-NC_012534.1 NNNNNNNNAACACC--AA-AACAAAACAGCAUAUUGACACCUGGGA--GAGACUAGGAGAC 156
MBFV-NC_015843.2 NNNNNNNNUAACAC--CA-CAAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGACC 156
MBFV-NC_009028.2 NNNNNNNNCAAGCG--CGAA-ACAAACAGCAUAUUGACGCCUGGGA--AAGACCGGGAGUUC 156
MBFV-NC_040776.1 NNNNNNNN--AAGCG--CAAAGCAAAACAGCAUAUUGACACCUGGGA--AAGACCGGGAGACC 156
MBFV-NC_034151.1 NNNNNNNNGAA--CG-CAAAUAAAAACAGCAUAUUGACACCUGGGA--UAGACUGGGAGACC 156
MBFV-NC_009026.2 NNNNNNNN--ACGCA--CCAAAGAAACAGCAUAUUGACACCUGGGA--UAGACCGGAGAAU 156
ISFV-NC_040610.1 NNNNNNNNCUCAU--AUUUUAUUAACAGCAUAUUGACACCAGGGA--UAGACCGGAGAAA--156
ISFV-NC_017086.1 NNNNNNNNC-CCAA--AAUUGAAGGCAUUAUUGACACCUGGGA--AAGACCAG--AGUA 155
MBFV-NC_001474.2 NNNNNNNNCCCCCG--AAACAAAAACAGCAUAUUGACG--CUGGGA--AAGACCAG--AGAUC 155
MBFV-NC_001477.1 NNNNNNNNCCCCCG--CAACAACAAACAGCAUAUUGACG--CUGGGA--GAGACCAG--AGAUC 155
MBFV-NC_001475.2 NNNNNNNNCCCCCG--CAAAUAAAAACAGCAUAUUGACG--CUGGGA--GAGACCAG--AGAUC 155
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MBFV-NC_012533.1 NNNNNNNNUAUAAC--CAACAUAAAAACAGCAUAUUGACACCUGGGA--AAGACCGG--AGACU 155
MBFV-NC_012532.1 NNNNNNNNCCCCGAAAACGCAAAACAGCAUAUUGACG--UGGGA--AAGACCAG--AGACU 155
MBFV-NC_035889.1 NNNNNNNN--CCCGGAAAACGCAAAACAGCAUAUUGACG--CUGGGA--AAGACCAG--AGACU 155
ISFV-NC_024017.1 NNNNNNNNCUGCUG--AGAGCAAAAAGCAUAUUGACACCUGGGA--GAGACCAG--AGAU 156
MBFV-NC_002031.1 NNNNNNNNA--GGGACAAAUAUGGGGACCAUAUUGACGCCAGG--A-AAGACC--GGAGUGG 156
ISFV-NC_027999.1 NNNNNNNNC--C-AUUGGAAAUUAGCUUCAUAUUGACACCAGGGA--AAGACC--GGAGAAA 156
MBFV-NC_012735.1 NNNNNNNNC-CGAGCACACAUAGCGGACCAUAUUGACGCCAGG--A-AAGACC--GGAGACA 156
MBFV-NC_008719.1 NNNNNNNNC-CGAGCAC-UAUAGCGGACCAUAUUGACGCCAGG--A-AAGACC--GGAGACA 155
ISFV-NC_016997.1 NNNNNNNN--CAAAAAUGUAUAGGCUAUAUAUAGCACCUCUAGG--A-AAGACU--GGAGUA 155
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NKV-NC_005039.1 NNNNNNNNUACGC--AUGGAUUUUGCAUAUUGAUUGACUAGG--AAGACCGGGUGUUC 156
NKV-NC_004119.1 NNNNNNNNUUGCCUCAUCUUGAAUGGGGGGCGGCGUGGAUAUAUACUCCAGCCAGAAAAAGA 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 CUCUGU-UC-----UCC 167
MBFV-NC_040776.1 CUCUGC-UC-----UGC 167
MBFV-NC_034151.1 CUCUGC-UU-----UCC 167
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ISFV-NC_040610.1 -UACACAU-----UUC 167
ISFV-NC_017086.1 CUCGCUGAU-----UCA 167
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MBFV-NC_012533.1 CUGUUGCU-----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 UUCUCUGCU-----U-U 167
ISFV-NC_027999.1 CUUCUUGCU-----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 UUCUGGCUAUCUCAACAUCAGCUACUA 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.....

