```
                    ..............................................................
ISFV-NC_034204.1    ---------A--UUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAG--------- 40
ISFV-NC_034017.1    ---------C--UAGUGUUGUUAAAAAUUUCUUUGGCGACGUACGGAAAGU--------- 40
ISFV-NC_024299.2    ---------C--UUUGGUUGUUUGAAAAACUUUUGCGACGUUGUUGAGAAG--------- 40
ISFV-NC_008604.2    ---------AAGGCUUGAGUUGUUUAUAAUAGUCGUUUUUCUCGCAGAA----------- 40
ISFV-NC_021069.1    ---------AA--UUUCGCUACCUUAUUACCUCUGGUACUAAAGUGGCACU--------- 40
ISFV-NC_012671.1    ---------UA--AUUUGCUACUUUAUUACCUCUGGUACUAACGUGGCGCU--------- 40
ISFV-NC_001564.2    ---------AA--UUUUUGCCGAGAGACCGUGAGGUUGAACCCGGCAAGGA--------- 40
ISFV-NC_005064.1    ---------AA--AUCAAAGGAGUUGUUCGGAAAAGUGACCUUGGUUCGUU--------- 40
ISFV-NC_012932.1    ---------AA--AAUCUAGAAGUUGUUUGGAAAAGAGAUAUCUGGUUCGU--------- 40
ISFV-NC_027819.1    ---------UAGACGUUUGUCACGGAAUAGCUU--GAGAAGAAGCUUACUU--------- 40
ISFV-NC_027999.1    --------------GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA-CU----- 40
MBFV-NC_008719.1    --------------AUUUCUACUGAAAGG-GUAGAGAAAAGGAGUUUUGCUU-CUC---- 40
MBFV-NC_012735.1    --------------UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUC-CCA---- 40
MBFV-NC_002031.1    --------------UAAUCGUUCGUUGAGCGAUUAG-CAGAGAACUGACCAG-AAC---- 40
NKV-NC_005039.1     --------------AAAU-UAUUUCCGGAAGAACAACUGUCCGGAAUCAAAG-ACG---- 40
MBFV-NC_009029.2    UGAUUACAUAACACAGUGUGAACAAGUUUU-UUGAACAGCA------------------- 40
MBFV-NC_032088.1    UC----UAUAACUCAGUGUGAACAAGUUUU-UUGAUCAGUGGA-U----A---------- 40
MBFV-NC_001437.1    -U------UAGUGCAGUUUAAACAGUUUUU-UAGAACGGAAGA-UAAC-C---------- 40
MBFV-NC_000943.1    --------UAACGCGGUUUGAACAGUUUUU-UGGAGCUUUUGA-UUUCAA---------- 40
MBFV-NC_006551.1    -U------UAACACAGUGCCGGCAGUUUCU-UUGAGC-GUUGA-UUUUCA---------- 40
MBFV-NC_001563.2    AU------UAACACAGUGCGAGCUGUUUCU-UGGCAC-GAAGA-UCUC-G---------- 40
MBFV-NC_009942.1    AU------UAACACAGUGCGAGCUGUUUCU-UAGCAC-GAAGA-UCUC-G---------- 40
MBFV-NC_009028.2    UU------UAACACAGUCUGAAUAGUUUUU-U--AGCAAGGGA-UUUCCC---------- 40
MBFV-NC_040776.1    UU------UAACACAGUCUGAAUAGUUUUU-U--AGCAGAGGU-ACUACC---------- 40
MBFV-NC_034151.1    --------UAACACAGUGUGAAUAGUUUUU-U-GAACAGAGGUGAAAACC---------- 40
MBFV-NC_012534.1    CCGUA----AACACAGUUUGAAC--GUUUU-UUGGAGAGAG---AUUAUU---------- 40
MBFV-NC_015843.2    ACGUA----AACACAGUUUGAAC--GUUUU-UUGGAUAGAG---ACAACU---------- 40
MBFV-NC_009026.2    AG-------AAAAGUUUUCGAGGAAAGACG-A---GAAGCGAA-UUCUCUGA-------- 40
MBFV-NC_007580.2    UA-------ACUUGACUGCGAACAGUUUUU-U---AGCAGGGA-AUUACCCA-------- 40
ISFV-NC_040610.1    --G------AACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAA-UUUC---A-------- 40
ISFV-NC_024017.1    UCG------CGCAUUUUGAUAGAGUU---GACUAGAUAACGAAUUAUC---G-------- 40
ISFV-NC_017086.1    AU-------AACACGUUUGAAUAGGUUUAU-UGUGAAUCGAAA-UACC---G-------- 40
MBFV-NC_012532.1    ---------AACAGGUUUA-----AUUUGGAUUUGGAAACGAGAGUUUCUGG--UC---- 40
MBFV-NC_035889.1    ---------ACAGGUUUUA-----UUUUGGAUUUGGAAACGAGAGUUUCUGG--UC---- 40
MBFV-NC_012533.1    ---------AACAGUUUUC-----GUGUUGAGAAAGAAAACAGAUUCUUCGG--AA---- 40
MBFV-NC_002640.1    ----------------UCUAACAGUUUGUUUGAAUAGAGAGCAGAUCUCUGGAAAA---- 40
MBFV-NC_001474.2    ----------CGUAGUUCUAACAGUUU-UUUAAUUAGAGAGCAGAUCUCUG--------- 40
MBFV-NC_001477.1    ---------ACGUAGUUCUAACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG--------- 40
MBFV-NC_001475.2    ---------ACGUAGUGCUGACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG--------- 40
ISFV-NC_016997.1    A---AACAUAACGCAUUAGAAUAGGUUUGAAAGAGUUGAAUCA----------------- 40
TBFV-NC_005062.1    GUGUUGAGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGG--- 40
TBFV-NC_027709.1    GUGUUGAGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGG--- 40
TBFV-NC_001672.1    GUGUUGAGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGG--- 40
TBFV-NC_001809.1    GUGUUGAGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGG--- 40
TBFV-NC_003690.1    GUGUUGAGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGA--- 40
TBFV-NC_003687.1    ------AGAAAAAGAC-----------------AGCUUAGGAGAACAAGAGCUGGGAGUG 37
ISFV-NC_027817.1    UUGAAAGGUUCAAACUCUGGUGUC-CAA-CGUUUGUAU---UAAA--------------- 40
NKV-NC_003635.1     ---AAUUUGUU--CUGUGCGCGUCACGCCACUUUUUGUGGCGGGA--------------- 40
NKV-NC_004119.1     ---ACCUUGGU--UUCCGGCUGGAAACCACGUCGCUCUUCGUCAA--------------- 40
                    .........10........20........30........40........50........6
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                    ...............................................((((.((((((((((
ISFV-NC_034204.1    ---AUGGAGC-----AAAACGGAAAG----------UUUUCCCGACCGGGAGGUGGUCGG 82
ISFV-NC_034017.1    ---AUGGAGC-----AAUCGAGGAAG----------UUUUCCCGGCCGGGAGGCGGUCGG 82
ISFV-NC_024299.2    ---AUGGAGCGAAAGAAAGGAGUAAA----------UUAUGGGAGAAGGGAAGCUAUCCC 87
ISFV-NC_008604.2    ---AUGGGAAAGGACGACGUAAGAA----------GAAGAAGGGGCCGGGCUCCUCCGG 87
ISFV-NC_021069.1    ---AUGAAGAAGAAGGA--GUCGGGA----------GUAGUGGAGAGAGGACGCGGUACG 85
ISFV-NC_012671.1    ---AUGAAGAAGAAGGA--GUCUGGA----------GUGGUGGAGAGAGGACGUGGAACG 85
ISFV-NC_001564.2    ---AUGAAGAGGAAGGAUUUGGAGGC----------G--CGAGGAAAAGCGCCAGGGAGG 85
ISFV-NC_005064.1    ---AUGAAUAGAGGAAGCUUGAAGAA----------G--GGGCCUCCGGGCCCAACAUCC 85
ISFV-NC_012932.1    ---AUGAAUAGGAUGCUCUUGAAGAA----------A--GGGCCUUCGGGCCAAACAUCU 85
ISFV-NC_027819.1    ---AUGUCAAGUAAGGAAGCACUGAA----------GAAGAAGGGAAUUUCCCUGGGAGG 87
ISFV-NC_027999.1    ---AUGGUUGAGAAAACUAGAGUAGG---AUCUCGGGUUAAAACCCGAGGUGUCAAUAUG 94
MBFV-NC_008719.1    ---AUGGC---AACAAGAGGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUG 94
MBFV-NC_012735.1    ---AUGGC---AACAAAGGGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUG 94
MBFV-NC_002031.1    ---AUGUCU-----------GGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUG 85
NKV-NC_005039.1     ---AUG--------AAAAAUCGUAGACAAAACACCCG----AAUUUCGGGUGUCAAUAUA 85
MBFV-NC_009029.2    ---AUGACCA---AAAAACCAGGAAG------GCCCGGUAGAAAUCGGGCCGUCAAUAUG 88
MBFV-NC_032088.1    ---AUGACUA---AAAAACCAGGAAA------GUCCGGCAAUCGUCGGACUGUCAAUAUG 88
MBFV-NC_001437.1    ---AUGACUA---AAAAACCAGGAGG------GCCCGGUAAAAACCGGGCUAUCAAUAUG 88
MBFV-NC_000943.1    ---AUGUCUA---AAAAACCAGGAGG------ACCCGGGAAGCCCCGGGUCGUCAAUAUG 88
MBFV-NC_006551.1    ---AUGUCUA---AGAAACCAGGAGG------GCCCGGAAGAAACCGGGCCAUCAAUAUG 88
MBFV-NC_001563.2    ---AUGUCUA---AGAAACCAGGAGG------GCCCGGUAAAAACCGGGCUGUCAAUAUG 88
MBFV-NC_009942.1    ---AUGUCUA---AGAAACCAGGAGG------GCCCGGCAAGAGCCGGGCUGUCAAUAUG 88
MBFV-NC_009028.2    ---AUGUCUA---AAAAACCAGGAAA------GUCCGCUGCAAAGCGGACUGUCAAUAUG 88
MBFV-NC_040776.1    ---AUGUCUA---AAAAACCAGGAGG------ACCCGCCGGAAGGCGGGUUGUCAAUAUG 88
MBFV-NC_034151.1    ---AUGUCUA---AAAAACCAGGAGG------ACCCGGAAAGAACCGGGUCGUUAAUAUG 88
MBFV-NC_012534.1    ---AUGUUUAACAAAAAACCAGGAAA------ACCCGGGGCAACCCGGGUUGUCAAUAUG 91
MBFV-NC_015843.2    ---AUGUCUAACAAAAAACCAGGAAG------ACCCGGCUCAGGCCGGGUUGUCAAUAUG 91
MBFV-NC_009026.2    ---AUGGCUA---GAAAACCAGGAAG------ACCCGGAGGAAACCGGGUCGUCAAUAUG 88
MBFV-NC_007580.2    ---AUGUCUA---AAAAACCAGGAAA------ACCCGGUAGAAACCGGGUUGUCAAUAUG 88
ISFV-NC_040610.1    ---AUGGUUA---GACAACUUGGAAA------GACCGUUGCAAGACGGUCUGUCAAUAUG 88
ISFV-NC_024017.1    ---AUGGUGA---CUAAAUCAAGAAA------GCCCGCCGCAAAGCGGGCUGUUAAUAUG 88
ISFV-NC_017086.1    ---AUGGCGA---AUAAACCCAAGAA------GCCCGG---AAGACGGGCUAUCGAUAUA 85
MBFV-NC_012532.1    ---AUGAAAAACCCCAAAGAAGAAAU--------CCGGAGGA-UCCGGAUUGUCAAUAUG 88
MBFV-NC_035889.1    ---AUGAAAAA-CCCAAAAAAGAAAU--------CCGGAGGAUUCCGGAUUGUCAAUAUG 88
MBFV-NC_012533.1    ---AUGAAAA--CCCAAAAAAGG-----------GCCGCAUCGCGGCCUAUCAAUAUG 82
MBFV-NC_002640.1    ---AUG---AA--CCAACGAAAAAAG--------GUG--GUUAGACCACCUUUCAAUAUG 82
MBFV-NC_001474.2    ---AUGAAUAA--CCAACGGAAAAAG--------GCG--AAAAACACGCCUUUCAAUAUG 85
MBFV-NC_001477.1    ---AUGAACAA--CCAACGGAAAAAG--------ACG--GGUCGACCGUCUUUCAAUAUG 85
MBFV-NC_001475.2    ---AUGAACAA--CCAACGGAAGAAG--------ACG--GGAAAACCGUCUAUCAAUAUG 85
ISFV-NC_016997.1    ---AUGAGAUCAAUGGUCUUUCGAGCUAGGAGGCCCGUUAAA---CGGGCCGUCGAUAUA 94
TBFV-NC_005062.1    ---AUGGCCGGGAAGGCCAUUCUGAAA---------GGAAAGGGGGGCGGUCCCCCUCGA 88
TBFV-NC_027709.1    ---AUGGGCAGGAAGUCCAUCCUGAAA---------GGUAAGGGGGGCGGUCCCCCUCGA 88
TBFV-NC_001672.1    ---AUGGUCAAGAAGGCCAUCCUGAAA---------GGUAAGGGGGGCGGUCCCCCUCGA 88
TBFV-NC_001809.1    ---AUGGGCAGGAAGACCAUCCUGAAA---------GGCAAGGGGGGCGGUCCCCCUCGA 88
TBFV-NC_003690.1    ---AUGGCCGGGAAGGCCGUUCUAAAA---------GGAAAGGGGGGGGGUCCCCCUCGA 88
TBFV-NC_003687.1    GUUAUG------AUGACCACUUCUAAA---------GGAAAGGGGGGCGGUCCCCCUAGG 82
ISFV-NC_027817.1    ---AUGAGUGGAUUAGGAGGCCUUCU----------------UCCCCUCCGAGGGAAGAAA 82
NKV-NC_003635.1     ---AUGUGCGAAAAUAACAGGAAAAA----------------GAACCCUGGAAAGGGUUCA 82
NKV-NC_004119.1     ---AUGACGAAAGGCAAAAUGAAAAA----------------CGCGAAUAAAAAGAAAGCA 82
                    0........70........80........90........100.......110.......1
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                    ((.............................................................))))))
ISFV-NC_034204.1    GCCCGGCCCAGAGAGGGGCUGGGGAAGANNNNNNNACG---ACCACCCUCUGAAAUGUUA 139
ISFV-NC_034017.1    GCGGGCCCCAAAGAGGGGCUGGCGAAGANNNNNNNGCG---ACCACCCUCUGUAAUGUUU 139
ISFV-NC_024299.2    GCCCCCUCCCAUUAAGGA-----GAAAANNNNNNNACG---A-CACCCCCCGAAAUGUGU 138
ISFV-NC_008604.2    AUGGUUGUUACCA--CCGGGGCG---CGNNNNNNNCUG-ACGCCACCCCCCGAAAUGUGU 141
ISFV-NC_021069.1    CGCCCUCCGACCAACCGUGGAGG---AANNNNNNNGUG-ACGCCACCCCCCGAAAUGUGU 141
ISFV-NC_012671.1    CGUCCUCCUGCCAAUCGUGGAGG---AGNNNNNNNGUG-ACGCCACCCCCCGAAAUGUGU 141
ISFV-NC_001564.2    GAUUUUUCCACCCCGUUCUGGGG---AANNNNNNNAG-CGAACCUCCUCCGUUAAUGUGA 141
ISFV-NC_005064.1    AACCGUCCGUCGGUUGGACCGCC---CGNNNNNNN---CGUACCCCCUCCG-UAAUGUGA 138
ISFV-NC_012932.1    GGUCGGCCAUCGACCAGACCGCC---GGNNNNNNNUUGCGAACCCCUCCG-AAAUGUGA 141
ISFV-NC_027819.1    AGGAGGAAAACGGGGCUUUGAG-----ANNNNNNNUUA-GCCACACCCCCCGAAACGUGA 141
ISFV-NC_027999.1    AU---------UAGGCGUACUGGAA---NNNNNNNC-CAUUGGAAAUUUAGCU--UCAUA 139
MBFV-NC_008719.1    GU---------GACGGCCAAGGCCA---NNNNNNNC-C-GAGCAC-UAUAGCGGACCAUA 139
MBFV-NC_012735.1    GU---------AGCGGCUAGAGUGA---NNNNNNNC-C-GAGCACACAUAGCGGACCAUA 140
MBFV-NC_002031.1    GUACGACGAGGAGUUCGCUCCUUGU---NNNNNNNA-GGGAACAAAU-AGUGGGACCAUA 140
NKV-NC_005039.1     AAAAAGCAAAAUCCAAAAAACAAAA---NNNNNNNU-ACGCAUGGAUUUUGC--AAUAUG 139
MBFV-NC_009029.2    CUAAAGCGCGGUGCGUCCCGCG------NNNNNNNA-GCGCAAACAC-A-AAACAGCAUA 139
MBFV-NC_032088.1    CUAAAGCGCGGAGCAUCCCGCG------NNNNNNNU-ACCGCAAAAC-A-AAACAGCAUA 139
MBFV-NC_001437.1    CUGAAACGCGGCCUACCCCGCG------NNNNNNNC-GCAUUUGCAU-C-AAACAGCAUA 139
MBFV-NC_000943.1    CUAAAACGCGGCAUACCCCGCG------NNNNNNNA-AAAGCAUCAA-A-CAACAGCAUA 139
MBFV-NC_006551.1    CUGAAACGCGGCAUACCCCGCG------NNNNNNNU-AAGCAUCAAA-A-AAACAGCAUA 139
MBFV-NC_001563.2    CUAAAACGCGGUAUGCCCCGCG------NNNNNNNA-AACACCAAAA-G-AAACAGCAUA 139
MBFV-NC_009942.1    CUAAAACGCGGAAUGCCCCGCG------NNNNNNNA-ACACCACAAC-A-AAACAGCAUA 139
MBFV-NC_009028.2    CUAAAACGCCUGGCUAGCGUGU------NNNNNNNC-AAGCGCGAA--ACAAACAGCAUA 139
MBFV-NC_040776.1    CUGAAACGCCCAGCCAGCGUUU------NNNNNNN--AAGCGCAAAG-ACAAACAGCAUA 139
MBFV-NC_034151.1    CUGAAGCGCACAUCCCGCGCUA------NNNNNNNG-AA-CGCAAAU-AAAAACAGCAUA 139
MBFV-NC_012534.1    CUAAAGCGCGGUGCAUCCC---------NNNNNNN--AACACCAAAA-CAAAACAGCAUA 139
MBFV-NC_015843.2    CUAAAGCGCGGAACGUCCC---------NNNNNNNG-AACACCACAA--GAAACAGCAUA 139
MBFV-NC_009026.2    CUAAAGCGUACGGCGGCCAACG------NNNNNNNACGC--ACCAAA-AGAAACAGCAUA 139
MBFV-NC_007580.2    CUAAAACGCGGCGUAUCCCGCG------NNNNNNNCCGUUAACGCAA-AC-AACAGCAUA 140
ISFV-NC_040610.1    CGAAAGC--GGAUUACCACCCG-CU---NNNNNNNCUCAUUAUUUAU-A-UAACAGCAUA 140
ISFV-NC_024017.1    CUAACAC--GCAUCACCAAGCG-UG---NNNNNNNCU-GCUGAGAGC-A-AAAUAGCAUA 139
ISFV-NC_017086.1    GUGAGACGUGCGUUGCCUCGCGUCU---NNNNNNNCCCAAAAAUGA--A-GGGCACUAUA 139
MBFV-NC_012532.1    CUAAAACGCGGAGUAGCCCGUG------NNNNNNNCCCCGGA-AAACGCAAAACAGCAUA 141
MBFV-NC_035889.1    CUAAAACGCGGAGUAGCCCGUG------NNNNNNN-CCCGGA-AAACGCAAAACAGCAUA 140
MBFV-NC_012533.1    CUGAAGCGCUUUGGAUCUAGCGCUGGCCNNNNNNNA---UAACCAACAUAAAACAGCAUA 139
MBFV-NC_002640.1    CUGAAACGCGAGAGAAACCGCGUAUCAANNNNNNNCCCCCC--AACACAAAAACAGCAUA 140
MBFV-NC_001474.2    CUGAAACGCGAGAGAAACCGCGUGU---NNNNNNNCCCCCG--AAACAAAAAACAGCAUA 140
MBFV-NC_001477.1    CUGAAACGCGCGAGAAACCGCGUGU---NNNNNNNCCCCGCA--CAACAACAAACAGCAUA 140
MBFV-NC_001475.2    CUGAAACGCGUGAGAAACCGUGUGU---NNNNNNNCCCCCG--CAAAUAAAAACAGCAUA 140
ISFV-NC_016997.1    AUAAAACGUAAGCUGC------------NNNNNNNC---AAAAAUGUAUAGGCUAUUAUA 139
TBFV-NC_005062.1    CGAGUGUCGAAAGAGACCGCUA------NNNNNNNUAUACCA-AAUUCCCCCUCAUCUGA 141
TBFV-NC_027709.1    CGAGUGUCGAAAGAGACCGCAA------NNNNNNNUAUACCA-AAUUCCCCCUCGACAGA 141
TBFV-NC_001672.1    CGAGUGUCGAAAGAGACCGCAA------NNNNNNNUAUACAA-AAUUCCCCCUCGGUAGA 141
TBFV-NC_001809.1    CGAGUGUCGAAAGAGACCGCGA------NNNNNNNAUACCAA-AUUUCCCCCUCGACAGA 141
TBFV-NC_003690.1    CGAGCGUCGAAAGUGGCCCCAA------NNNNNNNUAUACCA-AAUUCCCCCUCGUCAGA 141
TBFV-NC_003687.1    CGCAAGCUUAAAGUGACCGCAAAUAAGUNNNNNNNNAUACUAA-AUUUCCCCCAGGAAACU 141
ISFV-NC_027817.1    AAGAAAGCCCCUGUGAUACAAUCACAGGNNNNNNNAGUAGCGUCACCCC-------CCGG 135
NKV-NC_003635.1     AAGAAGAAAAAGAAUGCUGGAAGGAAUGNNNNNNN--UUUUCCAAAUUC-------C-CG 132
NKV-NC_004119.1     AAAAACUCUAAGACCAAUCCUGGUCGAANNNNNNN--UUGCCUCAUCUU-------GAAU 133
                    20.......130.......140.......150.......160.......170.......1
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                    ))))))..))))...................................
ISFV-NC_034204.1    AUAAAA-CUGUGCGGAAAAAUCCAAAGACUGUCCCCACC-------- 177
ISFV-NC_034017.1    AAUAAG-UUGUGCGGAAAAAUCCAAAGAAUGUCCCCACC-------- 177
ISFV-NC_024299.2    AUAUAGCCCAUGUUGAGUCUAUACAACAGGCUUCCACCG-------- 177
ISFV-NC_008604.2    AUAGUAACAGAGC-AUGCCUGCAGCAGCAGGUCUCCA---------- 177
ISFV-NC_021069.1    AUUGUGACAG-GC-UUGCUCGCAGCAGCAAAUCUCCAC--------- 177
ISFV-NC_012671.1    AUUGUGACAG-GC-UUGCUCGCAGCAGCAAAUCUCCAC--------- 177
ISFV-NC_001564.2    AAAAGUAUGGGG-AAAGAACUCAUCUUAA--CCUCCCAC-------- 177
ISFV-NC_005064.1    AAAAGUAUGGGGAAAAGAACCCAGCUAAAUCUCCCACCG-------- 177
ISFV-NC_012932.1    AAAAUUAUGGGGAAAAGUACCCAUCUA---AGCCUCCCA-------- 177
ISFV-NC_027819.1    UUUAU-AUGAUGACAAGAAUCAGAUGGAAGAUCCCCA---------- 177
ISFV-NC_027999.1    UUGACACCAGGGAA--AAGACCGGAGAAACUUCUUG-CUU----A-- 177
MBFV-NC_008719.1    UUGACGCCUGGG-A--AAGACCGGAGACACUCCUUGAUUC----U-- 177
MBFV-NC_012735.1    UUGACGCCAGGG-A--AAGACCGGAGACACUCCUUGAUUC------- 177
MBFV-NC_002031.1    UUGACGCCAGGG-A--AAGACCGGAGUGGUUCUCUG-CUU----U-- 177
NKV-NC_005039.1     UUGACAUCAGGGAA--AGACCGGGUGUUUGUCGGUU-CCG----G-- 177
MBFV-NC_009029.2    UUGACA-CCUGGGA--AAGACUAGGGGAUU--UGCUGCUC----UGG 177
MBFV-NC_032088.1    UUGACA-CCUGGGA--UAGACUAGGAGAUC--CGCUGCUC----UGG 177
MBFV-NC_001437.1    UUGACA-CCUGGGAA-UAGACUGGGAGAUC--UUCUGCUC----UA- 177
MBFV-NC_000943.1    UUGACA-CCUGGGAA-AAGACUAGGAGAUC--UUCUGCUC----UA- 177
MBFV-NC_006551.1    UUGACA-CCUGGGAAUUAGACUAGGAGAUC--UUCUGCUC----U-- 177
MBFV-NC_001563.2    UUGACA-CCUGGGA--UAGACUAGGGGAUC--UUCUGCUC----UGC 177
MBFV-NC_009942.1    UUGACA-CCUGGGA--UAGACUAGGAGAUC--UUCUGCUC----UGC 177
MBFV-NC_009028.2    UUGACG-CCUGGGA--AAGACCGGGAGAUC--CUCUGCUU----UCC 177
MBFV-NC_040776.1    UUGACA-CCUGGGA--AAGACCGGGAGACC--CUCUGCUC----UGC 177
MBFV-NC_034151.1    UUGACA-CCUGGGA--UAGACUGGGAGACC--CUCUGCUU----AUU 177
MBFV-NC_012534.1    UUGACA-CCUGGGA--GAGACUAGGAGACC--CUCUGUCC----UAA 177
MBFV-NC_015843.2    UUGACA-CCUGGGA--UAGACUAGGAGACC--CUCUGUCC----UAA 177
MBFV-NC_009026.2    UUGACA-CCUGGGA--UAGACCGGAGAAUU--UGCUGCCU----CGC 177
MBFV-NC_007580.2    UUGACA-CCUGGAA---AGACAGGAGAUCC--CCUGCUUU----UUC 177
ISFV-NC_040610.1    UUGACA-CCAGGG-AUAGACCGGAGAAAUACACAAUUUC-------- 177
ISFV-NC_024017.1    UUGACA-CUUGGGAAGAGACCAGAGAUAUG-AUGUUGUUU------- 177
ISFV-NC_017086.1    UCGACACUUGGG--AAAGACCAGAGGUACUCGCUGAUUCA------- 177
MBFV-NC_012532.1    UUGACG--UGGGAA---AGACCAGAGACUCCAUGAGUUUC----C-- 177
MBFV-NC_035889.1    UUGACG-CUGGGAA---AGACCAGAGACUCCAUGAGUUUC----C-- 177
MBFV-NC_012533.1    UUGACACCUGGGAA---AGACCGGAGACUCUGUUGGCUUU----C-- 177
MBFV-NC_002640.1    UUGACG-CUGGGAA---AGACCAGAGAUCCUGCUGUCUCU----G-- 177
MBFV-NC_001474.2    UUGACG-CUGGGAA---AGACCAGAGAUCCUGCUGUCUCC----U-- 177
MBFV-NC_001477.1    UUGACG-CUGGGAG---AGACCAGAGAUCCUGCUGUCUCU----A-- 177
MBFV-NC_001475.2    UUGACG-CUGGGAG---AGACCAGAGAUCCUGCUGUCUCC----U-- 177
ISFV-NC_016997.1    UCGACACCUAGGAA---AGACUGGAGAUAC--CUUUGGUU----CUU 177
TBFV-NC_005062.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCAUCA 177
TBFV-NC_027709.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCAUCA 177
TBFV-NC_001672.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCAUCA 177
TBFV-NC_001809.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCAUCA 177
TBFV-NC_003690.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCAUCA 177
TBFV-NC_003687.1    GGGGGGGCGG-----------UUCUUGUUCUCCCUGAGCCACCACCA 177
ISFV-NC_027817.1    UUGUGAAAACGAUUGCGACUA-----GAACUAAAGUCGAGAGUCUCC 177
NKV-NC_003635.1     UUACCGGGGUGGGGUGAUUAGCCAUGGUCGCACAGAUCAAGC--UC 177
NKV-NC_004119.1     GGGGGGCGGCGUGGAU-AUAUACUCCA--GCCAGAAAAGACUCAGAU 177
                    80.......190.......200.......210.......220.....
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