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                   ..............................................................
ISFV-NC_008604.2   U--UGUUUAUAAUAGUCGUUUUUCUCGCAGAA------AUG---------GGAAAGGACG 43
ISFV-NC_021069.1   C--CUUAUUACCUCUGGUACUAAAGUGGCACU------AUG---------AAGAAGAAGG 43
ISFV-NC_012671.1   C--UUUAUUACCUCUGGUACUAACGUGGCGCU------AUG---------AAGAAGAAGG 43
ISFV-NC_024299.2   U--UGAAAAACUUUUGCGACGUUGUUGAGAAG------AUG---------GAGCGAAAGA 43
ISFV-NC_027819.1   U--CACGGAAUAGCUUGAGAAGAAGCUUACUU------AUG---------UCAAGUAAGG 43
ISFV-NC_001564.2   G--AGAGACCGUGAGGUUGAACCCGGCAAGGA------AUG---------AAGAGGAAGG 43
ISFV-NC_005064.1   A--GUUGUUCGGAAAAGUGACCUUGGUUCGUU------AUG---------AAUAGAGGAA 43
ISFV-NC_012932.1   A--GUUGUUUGGAAAAGAGAUAUCUGGUUCGU------AUG---------AAUAGGAUGC 43
ISFV-NC_034204.1   G--CGACAGAAGGAACCCAAAAAGGGAACCAG------AUG---------GAGCAAAACG 43
ISFV-NC_034017.1   U--AAAAAUUUCUUUGGCGACGUACGGAAAGU------AUG---------GAGCAAUCGA 43
ISFV-NC_027817.1   C--AAACUCUGGUGUCCAACGUUUGUAUUAAA------AUG---------AGUGGAUUAG 43
TBFV-NC_003687.1   --------AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG--------------AUGA 37
TBFV-NC_001672.1   A--AAGACAGCUUAGGAGAACAAGAGCUGGGG------AUG---------GUCAAGAAGG 43
TBFV-NC_005062.1   A--AAGACAGCUUAGGAGAACAAGAGCUGGGG------AUG---------GCCGGGAAGG 43
TBFV-NC_001809.1   A--AAGACAGCUUAGGAGAACAAGAGCUGGGG------AUG---------GGCAGGAAGA 43
TBFV-NC_027709.1   A--AAGACAGCUUAGGAGAACAAGAGCUGGGG------AUG---------GGCAGGAAGU 43
TBFV-NC_003690.1   A--AAGACAGCUUAGGAGAACAAGAGCUGGGA------AUG---------GCCGGGAAGG 43
MBFV-NC_000943.1   ------U--GAACAGUUUUUUGGAGCUUUUGAUUUCAAAUG------UCU---AAAAAAC 43
MBFV-NC_006551.1   ------GCCG-GCAGUUUCUUUGAGC-GUUGAUUUUCAAUG------UCU---AAGAAAC 43
MBFV-NC_001563.2   ------UGCGAGCUGUUUCUUGGCAC-GAAGA-UCUCGAUG------UCU---AAGAAAC 43
MBFV-NC_009942.1   ------UGCGAGCUGUUUCUUAGCAC-GAAGA-UCUCGAUG------UCU---AAGAAAC 43
MBFV-NC_009029.2   ACACAGUGUGAACAAGUUUUUUGAACAG--------CAAUG------ACC---AAAAAAC 43
MBFV-NC_032088.1   ----AGUGUGAACAAGUUUUUUGAUCAGUGGA----UAAUG------ACU---AAAAAAC 43
MBFV-NC_007580.2   -------GCGAACAGUUUUUAGCAGGGAAU-UACCCAAUG------UCU---AAAAAAC 43
MBFV-NC_012534.1   ----AGUUUGAACGUUUUUUGGAGAGAGAUUAUU----AUG------UUUAACAAAAAAC 46
MBFV-NC_015843.2   ----AGUUUGAACGUUUUUUGGAUAGAGACAACU----AUG------UCUAACAAAAAAC 46
MBFV-NC_009028.2   ------UCUGAAUAGUUUUUUAGCAAGG--GAUUUCCCAUG------UCU---AAAAAAC 43
MBFV-NC_040776.1   ------UCUGAAUAGUUUUUUAGCAGAG--GUACUACCAUG------UCU---AAAAAAC 43
MBFV-NC_009026.2   UCGAGGAAAGA--------CGAGAAGCGAAUUCUCUGAAUG------GCU---AGAAAAC 43
MBFV-NC_034151.1   --------UGAAUAGUUUUUUGAACAGAGGUGAAAACCAUG------UCU---AAAAAAC 43
ISFV-NC_040610.1   A--------AAGAAGUCAAAACUAGAAAUUUAAUUUCAAUG------GUU---AGACAAC 43
ISFV-NC_017086.1   --------UGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG---------GCGAAUAAAC 43
ISFV-NC_016997.1   --------CAUUAGAAUAGGUUUGAAAGAGUUGAAUCAAUGAGAUCAAUGGUCUUUCGAG 52
ISFV-NC_024017.1   --------UGAUAGAGUUGACUAGAUAACGAAUUAUCGAUG---------GUGACUAAAU 43
MBFV-NC_001437.1   -------UUAAACAGUUUUUUAGAACGGAAGAUAACC-AUG---------ACUAAAAAAC 43
ISFV-NC_027999.1   -------UUGUUUGAGAGAACAGCGAGAAGUACUACU-AUG---GUUGAGAAAACUAGAG 49
MBFV-NC_008719.1   -------AAAGGGUAGAGAAAAG-GAGUUUUGCUUCUCAUG---GCAACAAGAGGAGCAA 49
MBFV-NC_012735.1   -------AAGGAGUAGAAGAAAG-GAGAUUCAUUCCCAAUG---GCAACAAAGGGGAUGA 49
MBFV-NC_002031.1   -------UUGAGCGAU-UAGCAGAGAACUGACCAGAACAUG---UC---------UGGUC 40
MBFV-NC_001474.2   ACAGUUUUUUAA---UUAGAGAGCAGAUCUCUG-----AUG--------AAUAA--CCAA 42
MBFV-NC_001477.1   AACAGUUUUUUA---UUAGAGAGCAGAUCUCUG-----AUG--------AACAA--CCAA 42
MBFV-NC_001475.2   GACAGUUUUUUA---UUAGAGAGCAGAUCUCUG-----AUG--------AACAA--CCAA 42
MBFV-NC_002640.1   -----UGUUUGA---AUAGAGAGCAGAUCUCUGGAAAAAUG-----------AA--CCAA 39
MBFV-NC_012532.1   -----AUUUGGA---UUUGGAAACGAGAGUUUCUGGUCAUG--------AAAAACCCCAA 44
MBFV-NC_035889.1   -----UUUUGGA---UUUGGAAACGAGAGUUUCUGGUCAUG--------AAAAA-CCCAA 43
MBFV-NC_012533.1   ----GUGUUGAG---AAAGAAAACAGAU-UCUUCGGAAAUG--------AA-------AA 37
NKV-NC_005039.1    --------CGGAAGAACAACUGUCCGGAAUCAAAGACGAUG------------AAAAAUC 40
NKV-NC_004119.1    ----CCGGCUGGAAACCACGUCGCUCUUCGU----CAAAUG--------ACGAAAGGCAA 44
NKV-NC_003635.1    ----GUGCGCG----UCACGCCACUUUUUGUGGCGGGAAUG--------UGCGAAAAUAA 44
                   .........10........20........30........40........50........6
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                   ...................(((((...(((((((((.......................
ISFV-NC_008604.2   ACGGUAAGAAGAAGAAGGGGCCGG----GCUCCUCCGGAUGGUUGUUACCA--CCGGGGC 97
ISFV-NC_021069.1   A--GUCGGGAGUAGUGGAGAGAGG----ACGCGGUACGCGCCCUCCGACCAACCGUGGAG 97
ISFV-NC_012671.1   A--GUCUGGAGUGGUGGAGAGAGG----ACGUGGAACGCGUCCUCCUGCCAAUCGUGGAG 97
ISFV-NC_024299.2   AAGGAGUAAAUUAUGGGAGAAGGG----AAGC--UAUCCCGCCCCCUCCCAUUAAGGAGA 97
ISFV-NC_027819.1   AAGCACUGAAGAAGAAGGGAAUUU----CCCUGGGAGGAGGAGGAAAACGGGGCUUUGAG 99
ISFV-NC_001564.2   --AUUUGGAGGCGCGAGGAAAAGC----GCCAGGGAGGGAUUUUUCCACCCCGUUCUGGG 97
ISFV-NC_005064.1   --GCUUGAAGAAGGGGCCUCCGGG----CCCAACAUCCAACCGUCCGUCGGUUGGACCGC 97
ISFV-NC_012932.1   --UCUUGAAGAAAGGGCCUUCGGG----CCAAACAUCUGGUCGGCCAUCGACCAGACCGC 97
ISFV-NC_034204.1   --GAAAGUUUUCCCGACCGGGAGG----UGGUCGGGCCCGGCCCAGAGAGGGGCUGGGGA 97
ISFV-NC_034017.1   --GGAAGUUUUCCCGGCCGGGAGG----CGGUCGGGCGGGCCCCAAAGAGGGGCUGGCGA 97
ISFV-NC_027817.1   GAGGCCUUCUUCCCCUCCGAGGGA----AGAA--AAAGAAAGCCCCUGUGAUACAAUCAC 97
TBFV-NC_003687.1   CCACUUCUAAAGGAAAGGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUA 97
TBFV-NC_001672.1   CCAUCCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA--- 100
TBFV-NC_005062.1   CCAUUCUGAAAGGAAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA--- 100
TBFV-NC_001809.1   CCAUCCUGAAAGGCAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA--- 100
TBFV-NC_027709.1   CCAUCCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA--- 100
TBFV-NC_003690.1   CCGUUCUAAAAGGAAAGGGGGGGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA--- 100
MBFV-NC_000943.1   CAGGAGGACCCGGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGGCAUACCCCGCG--- 100
MBFV-NC_006551.1   CAGGAGGGCCCGGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGGCAUACCCCGCG--- 100
MBFV-NC_001563.2   CAGGAGGGCCCGGUAAAAACCGGGCUGUCAAUAUGCUAAAACGCGGUAUGCCCCGCG--- 100
MBFV-NC_009942.1   CAGGAGGGCCCGGCAAGAGCCGGGCUGUCAAUAUGCUAAAACGCGGAAUGCCCCGCG--- 100
MBFV-NC_009029.2   CAGGAAGGCCCGGUAGAAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUGCGUCCCGCG--- 100
MBFV-NC_032088.1   CAGGAAAGUCCGGCAAUCGUCGGACUGUCAAUAUGCUAAAGCGCGGAGCAUCCCGCG--- 100
MBFV-NC_007580.2   CAGGAAAACCCGGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGGCGUAUCCCGCG--- 100
MBFV-NC_012534.1   CAGGAAAACCCGGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUGCAUCCC------ 100
MBFV-NC_015843.2   CAGGAAGACCCGGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAACGUCCC------ 100
MBFV-NC_009028.2   CAGGAAAGUCCGCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCUAGCGUGU--- 100
MBFV-NC_040776.1   CAGGAGGACCCGCCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAGCCAGCGUUU--- 100
MBFV-NC_009026.2   CAGGAAGACCCGGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGGCGGCCAACG--- 100
MBFV-NC_034151.1   CAGGAGGACCCGGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGCACAUCCCGCGCUA--- 100
ISFV-NC_040610.1   UUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUUACCACCCGCU--- 100
ISFV-NC_017086.1   CCAAGAAGCCCGGAAGA---CGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCGCGUCU 100
ISFV-NC_016997.1   CUAGGAGGCCCGUUAAA---CGGGCCGUCGAUAUAAUAAAACGUAAGCUGC--------- 100
ISFV-NC_024017.1   CAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAGCGUG--- 100
MBFV-NC_001437.1   CAGGAGGGCCCGGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCCUACCCCGCG--- 100
ISFV-NC_027999.1   UAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUACUG---------GAA 100
MBFV-NC_008719.1   GCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGUGACGGCCAAGG---------CCA 100
MBFV-NC_012735.1   AUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAGCGGCUAGAG---------UGA 100
MBFV-NC_002031.1   GUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUACGACGAGGAGUUCGCUCCUUGU 100
MBFV-NC_001474.2   CGGAAAAAGGCGAAAAACAC--GCCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUGU 100
MBFV-NC_001477.1   CGGAAAAAGACGGGUCGACC--GUCUUUCAAUAUGCUGAAACGCGCGAGAAACCGCGUGU 100
MBFV-NC_001475.2   CGGAAGAAGACGGGAAAACC--GUCUAUCAAUAUGCUGAAACGCGUGAGAAACCGUGUGU 100
MBFV-NC_002640.1   CGAAAAAAGGUGGUUAGACC--ACCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUAU 97
MBFV-NC_012532.1   AGAAGAAAUCCGGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG--- 100
MBFV-NC_035889.1   AAAAGAAAUCCGGAGGAUUCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG--- 100
MBFV-NC_012533.1   CCCAAAAAAGGGCCGCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGAUCUAGCGCUG 97
NKV-NC_005039.1    GUAGACAAAACACCCGAAUUUCGGGUGUCAAUAUAAAAAAGCAAAAUCCAAAAAACAAAA 100
NKV-NC_004119.1    AAUGAAAAACGCGAAUAAAAAGAAAGCAAAAAACUCUAAGACC-------AAUCCUGGUC 97
NKV-NC_003635.1    CAGGAAAAAGAACCCUGGAAAGGGUUCAAAGAAGAAAAAGAAU-------GCUGGAAGGA 97
                   0........70........80........90........100.......110.......1
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                   ..............................)))))))))).)))))................
ISFV-NC_008604.2   GCGNNNNNNN-CUGACGCCACCUCCCG-AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGC 155
ISFV-NC_021069.1   GAANNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGC 154
ISFV-NC_012671.1   GAGNNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGC 154
ISFV-NC_024299.2   AAANNNNNNN----ACGACACCCCCCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUAC 152
ISFV-NC_027819.1   --ANNNNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAU 155
ISFV-NC_001564.2   GAANNNNNNN--AGCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAACUCAU 154
ISFV-NC_005064.1   CCGNNNNNNN----CGUACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCCAG 152
ISFV-NC_012932.1   CGGNNNNNNN-UUGCGAACCCCCUCCG-AAAUGUGAAAAAUUAUGGGGAAAAGUACCCAU 155
ISFV-NC_034204.1   AGANNNNNNN---ACGACCACCCUCUG-AAAUGUUAAUAAAACUGUGCGGAAAAAUCCAA 153
ISFV-NC_034017.1   AGANNNNNNN---GCGACCACCCUCUG-UAAUGUUUAAUAAGUUGUGCGGAAAAAUCCAA 153
ISFV-NC_027817.1   AGGNNNNNNNAGUAGCGUCACCCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAAAG 155
TBFV-NC_003687.1   AGUNNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGAAACUGGGGGGGCGG 156
TBFV-NC_001672.1   ---NNNNNNN---------------UAUACAAAA-UUCCCCCUCGGUAGAGGGGGGGCGG 141
TBFV-NC_005062.1   ---NNNNNNN---------------UAUACCAAA-UUCCCCCUCAUCUGAGGGGGGGCGG 141
TBFV-NC_001809.1   ---NNNNNNNG-UUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGACAGAGGGGGGGCGG 156
TBFV-NC_027709.1   ---NNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGACAGAGGGGGGGCGG 156
TBFV-NC_003690.1   ---NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUCAGAGGGGGGGCGG 156
MBFV-NC_000943.1   ---NNNNNNAAAAGCAU-C-AAACAACAGCAUAUUGACACCUGGGAAA-AGACUAGGAG 154
MBFV-NC_006551.1   ---NNNNNNUAAGCAUC-A-AAAAAACAGCAUAUUGACACCUGGGAAUUAGACUAGGAG 155
MBFV-NC_001563.2   ---NNNNNNAAACACCA-A-AAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGG 153
MBFV-NC_009942.1   ---NNNNNNAACACCAC-A-ACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAG 153
MBFV-NC_009029.2   ---NNNNNNAGCGCAAA-C-ACAAAACAGCAUAUUGACACCUGGGA--AAGACUAGGGG 153
MBFV-NC_032088.1   ---NNNNNNUACCGCAA-A-ACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAG 153
MBFV-NC_007580.2   ---NNNNNNCCGUUAACGC-AAACAACAGCAUAUUGACACCUGGAA---AGAC-AGGAG 152
MBFV-NC_012534.1   ---NNNNNNAACACCAA-A-ACAAAACAGCAUAUUGACACCUGGGA--GAGACUAGGAG 153
MBFV-NC_015843.2   ---NNNNNNGAACACCA-C-AAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAG 153
MBFV-NC_009028.2   ---NNNNNNCAAGCGCGAA--ACAAACAGCAUAUUGACGCCUGGGAA--AGACCGGGAG 153
MBFV-NC_040776.1   ---NNNNNNN-AAGCGCAAA-GACAAACAGCAUAUUGACACCUGGGAA--AGACCGGGAG 153
MBFV-NC_009026.2   ---NNNNNNN-ACGCACCAA-AAGAAACAGCAUAUUGACACCUGGGA--UAGACCGGAGA 153
MBFV-NC_034151.1   ---NNNNNNNGAA-CGCAAA-UAAAAACAGCAUAUUGACACCUGGGA--UAGACUGGGAG 153
ISFV-NC_040610.1   ---NNNNNNNCUCAUUAUUU-AUAUAACAGCAUAUUGACACCAGGGA--UAGACCGGAGA 154
ISFV-NC_017086.1   ---NNNNNNNCCCA-AAAAU-GAAGGGCACUAUAUCGACACUUGGGAA--AGACCAG-AG 152
ISFV-NC_016997.1   ---NNNNNNNCAAA-AAUGU-AUAGGCUAUUAUAUCGACACCUAGGAA--AGACUGG-AG 152
ISFV-NC_024017.1   ---NNNNNNNC-UGCUGAGA-GCAAAAUAGCAUAUUGACACUUGGGAA-GAGACCAG-AG 153
MBFV-NC_001437.1   ---NNNNNNN----------------CAUAUUGA---CACCUGGGAA-UAGACUGGGAG 136
ISFV-NC_027999.1   ---NNNNNNNCC-AUUGGAA-AUUUAGCUUCAUAUUGACACCAGGGAAAAGACCGGAGAA 155
MBFV-NC_008719.1   ---NNNNNNNCCGAGCAC-U-AUAGCGGACCAUAUUGACGCCUGGG-AAAGACCGGAGAC 154
MBFV-NC_012735.1   ---NNNNNNNCCGAGCACAC-AUAGCGGACCAUAUUGACGCCAGGG-AAAGACCGGAGAC 155
MBFV-NC_002031.1   ---NNNNNNAGGGAACAAA-UAGUGGGACCAUAUUGACGCCAGGG-AAAGACCGGAGUG 155
MBFV-NC_001474.2   ---NNNNNNCCCCCGAAACAA-AAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGA 153
MBFV-NC_001477.1   ---NNNNNNCCCGCACAACAA-CAAACAGCAUAUUGACG-CUGGGAG--AGACCAGAGA 153
MBFV-NC_001475.2   ---NNNNNNCCCCCGCAAAUA-AAAACAGCAUAUUGACG-CUGGGAG--AGACCAGAGA 153
MBFV-NC_002640.1   CAANNNNNNNCCCCCCAACACA-AAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGA 153
MBFV-NC_012532.1   ---NNNNNNCCCCGGAAAACGCAAAACAGCAUAUUGACG--UGGGAA--AGACCAGAGA 153
MBFV-NC_035889.1   ---NNNNNN-CCCGGAAAACGCAAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGA 153
MBFV-NC_012533.1   GCCNNNNNNNAUAACCAA--CAUAAAACAGCAUAUUGACACCUGGGAA--AGACCGGAGA 153
NKV-NC_005039.1    ---NNNNNNUACGC--AUGGAUUUUGCAAUAUGUUGACAUCAGGGAA--AGACCGGGUG 153
NKV-NC_004119.1    GAANNNNNNNUUGCCUCAUCUUGAAU------GGGGGGCGGCGUGGAUAUAUACUCCAGC 151
NKV-NC_003635.1    AUGNNNNNNNUUUUCCAAAUUCCCGU------UACCGGGGGUGGGGUGAUUAGCCAUGGU 151
                   20.......130.......140.......150.......160.......170.......1
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ISFV-NC_008604.2   AGCAGGUCUCC----------------A--- 167
ISFV-NC_021069.1   AGCAAAUCUCC----------------AC-- 167
ISFV-NC_012671.1   AGCAAAUCUCC----------------AC-- 167
ISFV-NC_024299.2   AACAGGCUUCC----------------ACCG 167
ISFV-NC_027819.1   GGAAGAUCCCC----------------A--- 167
ISFV-NC_001564.2   CUUAACCUCCC----------------AC-- 167
ISFV-NC_005064.1   CUAAAUCUCCC----------------ACCG 167
ISFV-NC_012932.1   CUAAGCCUCCC----------------A--- 167
ISFV-NC_034204.1   AGACUGUCCCC----------------ACC- 167
ISFV-NC_034017.1   AGAAUGUCCCC----------------ACC- 167
ISFV-NC_027817.1   UCGAGAGUCUC-------------------C 167
TBFV-NC_003687.1   UUCUUGUUCUC--------------------- 167
TBFV-NC_001672.1   UUCUUGUUCUCCCU-----GAGCCACCAUCA 167
TBFV-NC_005062.1   UUCUUGUUCUCCCU-----GAGCCACCAUCA 167
TBFV-NC_001809.1   UUCUUGUUCUC--------------------- 167
TBFV-NC_027709.1   UUCUUGUUCUC--------------------- 167
TBFV-NC_003690.1   UUCUUGUUCUC--------------------- 167
MBFV-NC_000943.1   AUCUUCUGC-UC----------------UA- 167
MBFV-NC_006551.1   AUCUUCUGC-UC----------------U-- 167
MBFV-NC_001563.2   AUCUUCUGC-UC----------------UGC 167
MBFV-NC_009942.1   AUCUUCUGC-UC----------------UGC 167
MBFV-NC_009029.2   AUUUGCUGC-UC----------------UGG 167
MBFV-NC_032088.1   AUCCGCUGC-UC----------------UGG 167
MBFV-NC_007580.2   AUCCCCUGCUUU----------------UUC 167
MBFV-NC_012534.1   ACCCUCUGU-CC----------------UAA 167
MBFV-NC_015843.2   ACCCUCUGU-CC----------------UAA 167
MBFV-NC_009028.2   AUCCUCUGC-UU----------------UCC 167
MBFV-NC_040776.1   ACCCUCUGC-UC----------------UGC 167
MBFV-NC_009026.2   AUUUGCUGCCUC-----------------GC 167
MBFV-NC_034151.1   ACCCUCUGCUUA----------------U-U 167
ISFV-NC_040610.1   AA--UACACAAU----------------UUC 167
ISFV-NC_017086.1   GUACUCGCUGAU----------------UCA 167
ISFV-NC_016997.1   AUACCUUUGGUU----------------CUU 167
ISFV-NC_024017.1   AUAUGAUGU-UG----------------UUU 167
MBFV-NC_001437.1   AUCUUCUGCUCUAUCUCAACAUCAGCUACUA 167
ISFV-NC_027999.1   ACUUCUUGCUU-------------------A 167
MBFV-NC_008719.1   ACUCCUUGAUUC------------------U 167
MBFV-NC_012735.1   ACUCCUUGAUUC--------------------- 167
MBFV-NC_002031.1   GUUCUCUGCUU------------------U 167
MBFV-NC_001474.2   UCCUGCUGUCUC-----------------CU 167
MBFV-NC_001477.1   UCCUGCUGUCUC-----------------UA 167
MBFV-NC_001475.2   UCCUGCUGUCUC-----------------CU 167
MBFV-NC_002640.1   UCCUGCUGUCUC-----------------UG 167
MBFV-NC_012532.1   CUCCAUGAGUUU-----------------CC 167
MBFV-NC_035889.1   CUCCAUGAGUUU-----------------CC 167
MBFV-NC_012533.1   CUCUGUUGGCUU-----------------UC 167
NKV-NC_005039.1    UUUGUCGGUUCC-----------------GG 167
NKV-NC_004119.1    CAGAAAAGACUC----------------AGAU 167
NKV-NC_003635.1    CGCACAGAUCAA----------------GCUC 167
                   80.......190.......200.........
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