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                    ............................................................
NKV-NC_004119.1     -----CCGGCUGGAAACCACGUCGCUCUUCGU----CAAAUGACG----AAAGGCAAAAU 47
NKV-NC_003635.1     -----GUGCGCG----UCACGCCACUUUUGUGGCGGGAAUG-----------UGCGAA 39
ISFV-NC_001564.2    -----GAGAGAC----CGUGAGGUUGAACCCGGCAAGGAAUGAAG----------AGGAA 41
ISFV-NC_005064.1    -----AGUUGUU----CGGAAAAGUGACCUUGGUUCGUUAUGAAU----------AGAGG 41
ISFV-NC_012932.1    -----AGUUGUU----UGGAAAAGAGAUAUCUGGUUCGUAUGAAU----------AGGAU 41
ISFV-NC_034204.1    -----GCGACAG----AAGGAACCCAAAAAGGGAACCAGAUGGAG----------CAAAA 41
ISFV-NC_034017.1    -----UAAAAAU----UUCUUUGGCGACGUACGGAAAGUAUGGAG----------CAAUC 41
ISFV-NC_008604.2    -----UUGUUUA----UAAUAGUCGUUUUUCUCGCAGAAAUGGGA----------AAGGA 41
ISFV-NC_021069.1    -----CCUUAUU----ACCUCUGGUACUAAAGUGGCACUAUGAAG----------AAGAA 41
ISFV-NC_012671.1    -----CUUUAUU----ACCUCUGGUACUAACGUGGCGCUAUGAAG----------AAGAA 41
ISFV-NC_024299.2    -----UUGAAAA----ACUUUUGCGACGUUGUUGAGAAGAUGGAG----------CGAAA 41
ISFV-NC_027819.1    -----UCACGGA----AUAGCUUGAGAAGAAGCUUACUUAUGUCA----------AGUAA 41
ISFV-NC_027817.1    -----CAAACUC----UGGUGUCCAACGUUUGUAUUAAAAUGAGU----------GGAUU 41
TBFV-NC_003690.1    AAAGACAGCUUAGGAGAACAAGAGCUGGGA---------AUGGCCGGGAAGGCCGUUCUA 51
TBFV-NC_001809.1    AAAGACAGCUUAGGAGAACAAGAGCUGGGG---------AUGGGCAGGAAGACCAUCCUG 51
TBFV-NC_027709.1    AAAGACAGCUUAGGAGAACAAGAGCUGGGG---------AUGGGCAGGAAGUCCAUCCUG 51
TBFV-NC_001672.1    AAAGACAGCUUAGGAGAACAAGAGCUGGGG---------AUGGUCAAGAAGGCCAUCCUG 51
TBFV-NC_005062.1    AAAGACAGCUUAGGAGAACAAGAGCUGGGG---------AUGGCCGGGAAGGCCAUUCUG 51
TBFV-NC_003687.1    ------AGCUUAGGAGAACAAGAGCUGGGAGUGGU---UAUG------AUGACCACUUCU 45
ISFV-NC_017086.1    UGAAUAGGUUUA-UUGUGAAUCGAAAUACCG--------AUGGCGAA--UAAACCCAAGA 49
MBFV-NC_002640.1    ------UGUUUG-AAUAGAGAGC-AG-AUCUCUGGAAAAAUG---AA--CCAACGAAAAA 46
MBFV-NC_001474.2    ACAGU-UUUUUA-AUUAGAGAGC-AG-AUCUCUG-----AUGAAUAA--CCAACGGAAAA 49
MBFV-NC_001477.1    AACAG-UUUUUU-AUUAGAGAGC-AG-AUCUCUG-----AUGAACAA--CCAACGGAAAA 49
MBFV-NC_001475.2    GACAG-UUUUUU-AUUAGAGAGC-AG-AUCUCUG-----AUGAACAA--CCAACGGAAGA 49
MBFV-NC_012532.1    ----A-UUUGGA-UUUGGAAACG-AGAGUUUCUGG--UCAUGAAAAACCCCAAAGAAGAA 51
MBFV-NC_035889.1    ----U-UUUGGA-UUUGGAAACG-AGAGUUUCUGG--UCAUGAAAAA-CCCAAAAAAGAA 50
MBFV-NC_012533.1    ----G-UGUUGA-GAAAGAAAAC-AGAUUCUUCGG--AAAUGAAAA--CCCAAAAAAGG- 48
MBFV-NC_001437.1    -------UUAAACAGUUUUUUAGAACGGAAG--AUAACCAUGACU---AAAAAACCAGGA 48
MBFV-NC_009029.2    ACACAGUGUGAACAAGUUUUUUGAACAG---------CAAUGACC---AAAAAACCAGGA 48
MBFV-NC_032088.1    ----AGUGUGAACAAGUUUUUUGAUCAGUGG-----AUAAUGACU---AAAAAACCAGGA 48
MBFV-NC_000943.1    --------UGAAC-AGUUUUUUGGAGCUUUUGAUUUCAAAUGUCU---AAAAAACCAGGA 48
MBFV-NC_034151.1    --------UGAAU-AGUUUUUUGAACAGAGGUGAAAACCAUGUCU---AAAAAACCAGGA 48
MBFV-NC_001563.2    ------UGCGAGCUGUUUCUUGGCACGAAG--A-UCUCGAUGUCU---AAGAAACCAGGA 48
MBFV-NC_009942.1    ------UGCGAGCUGUUUCUUAGCACGAAG--A-UCUCGAUGUCU---AAGAAACCAGGA 48
MBFV-NC_006551.1    ------GCCG-GCAGUUUCUUUGAGCGUUG--AUUUUCAAUGUCU---AAGAAACCAGGA 48
MBFV-NC_012534.1    ----AGUUUGAAC--GUUUUUUGGAGAGAG---AUUAUUAUGUUUAACAAAAAACCAGGA 51
MBFV-NC_015843.2    ----AGUUUGAAC--GUUUUUUGGAUAGAG---ACAACUAUGUCUAACAAAAAACCAGGA 51
MBFV-NC_009026.2    ------UC-GAGGAAAGACGA-GAAGCG-AAUUCUCUGAAUGGCU---AGAAAACCAGGA 48
MBFV-NC_007580.2    ------GC-GAACAGUUUUUU-AGCAGG-GAAUUACCCAAUGUCU---AAAAAACCAGGA 48
MBFV-NC_009028.2    ------UCUGAAUAGUUUUUUAGCAAGG-GAUUUCCC--AUGUCU---AAAAAACCAGGA 48
MBFV-NC_040776.1    ------UCUGAAUAGUUUUUUAGCAGAG-GUACUACC--AUGUCU---AAAAAACCAGGA 48
ISFV-NC_024017.1    --------UGAUAGAGUUGACUAGAUAACGA-AUUAUCGAUGGUG---ACUAAAUCAAGA 48
MBFV-NC_002031.1    --------UUGAGCGAU-UAGCAGAGAACUGACCAGAACAUGUC---------UGGUCGU 42
MBFV-NC_008719.1    --------AAAGGGUAGAGAAAAG-GAGAUUUGCUUCUCAUGGCAACAAGAGGAGCAAGC 51
MBFV-NC_012735.1    --------AAGGAGUAGAAGAAAG-GAGAUUCAUUCCCAAUGGCAACAAAGGGGAUGAAU 51
ISFV-NC_027999.1    --------UUGUUUGAGAGAACAGCGAGAAGUACUACU-AUGGUUGAGAAAACUAGAGUA 51
ISFV-NC_040610.1    --------AAAGAAGUCAAAAC-UAGAAAUUUAAUUUCAAUGGUUAGACAACUUGGAAAG 51
NKV-NC_005039.1     --------CGGAAGAACAACUGUCCG-GAAUCAAAGACGAUG----AAAAAUCGUAGACA 47
ISFV-NC_016997.1    CA----UUAGAAUAGGUUUGAAAGAG-----UUGAAUCAAUGAGAUCAAUGGUCUUUCGA 51
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NKV-NC_004119.1     GAAAAACGCGAAUAAAA-----AGAAAGCAAAAAACUCUAAGACCAAUCCUGGUCGAA-- 100
NKV-NC_003635.1     AAUAACAGGAAAAAGAACCCUGGAAAGGGUUCAAAGAAGAAAAAGAAUGCUGGAAGGAAU 99
ISFV-NC_001564.2    GG--AUUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGA 99
ISFV-NC_005064.1    AA--GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCC 99
ISFV-NC_012932.1    GC--UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCG 99
ISFV-NC_034204.1    CG--GAAAGUUUUCCCGACCGGGAGGUGGUCGGGCCCGGCCCAGAGAGGGGCUGGGGAAG 99
ISFV-NC_034017.1    GA--GGAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGGCGAAG 99
ISFV-NC_008604.2    CGACGGUAAGAAGAAGAAGGGGCCGGGCUCCUCCGGAUGGUUGUUACCA--CCGGGGCGC 99
ISFV-NC_021069.1    GGA--GUCGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGA 99
ISFV-NC_012671.1    GGA--GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGA 99
ISFV-NC_024299.2    GAAAGGAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAA 99
ISFV-NC_027819.1    GGAAGCACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAGGAAAACGGGGCUUUGAG-- 99
ISFV-NC_027817.1    AGGAGGCCUUCUUCCCCUCCGAGGGAAGAA--AAAGAAAGCCCCUGUGAUACAAUCACAG 99
TBFV-NC_003690.1    AAAGGAAAGGGGGGGGUCCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA----------- 100
TBFV-NC_001809.1    AAAGGCAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA----------- 100
TBFV-NC_027709.1    AAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA----------- 100
TBFV-NC_001672.1    AAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA----------- 100
TBFV-NC_005062.1    AAAGGAAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA----------- 100
TBFV-NC_003687.1    AAAGGAAAGGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAAGU----- 100
ISFV-NC_017086.1    AG--CC-----CGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCGCGUCU-- 100
MBFV-NC_002640.1    AG--GU---G--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUAUCA 99
MBFV-NC_001474.2    AG--GC---G--AAAAACACGCCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGUGU-- 100
MBFV-NC_001477.1    AG--AC---G--GGUCGACCGUCUUUCAAUAUGCUGAAACGCGCGAGAAACCGCGUGU-- 100
MBFV-NC_001475.2    AG--AC---G--GGAAACCGUCUAUCAAUAUGCUGAAACGCGUGAGAAACCGUGUGU-- 100
MBFV-NC_012532.1    AU--CC---GGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG----- 100
MBFV-NC_035889.1    AU--CC---GGAGGAUUCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG----- 100
MBFV-NC_012533.1    ---------GCCGCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGAUCUAGCGCUGGC 99
MBFV-NC_001437.1    GGGCCC---GGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCCUACCCCGCG----- 100
MBFV-NC_009029.2    AGGCCC---GGUAGAAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUGCGUCCCGCG----- 100
MBFV-NC_032088.1    AAGUCC---GGCAAUCGUCGGACUGUCAAUAUGCUAAAGCGCGGAGCAUCCCGCG----- 100
MBFV-NC_000943.1    GGACCC---GGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGGCAUACCCCGCG----- 100
MBFV-NC_034151.1    GGACCC---GGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGC-ACAUCCCGCGCUA---- 100
MBFV-NC_001563.2    GGGCCC---GGUAAAAACCGGGCUGUCAAUAUGCUAAAACGCGGUAUGCCCCGCG----- 100
MBFV-NC_009942.1    GGGCCC---GGCAAGAGCCGGGCUGUCAAUAUGCUAAAACGCGGAAUGCCCCGCG----- 100
MBFV-NC_006551.1    GGGCCC---GGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGGCAUACCCCGCG----- 100
MBFV-NC_012534.1    AAACCC---GGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUGCAUCCC-------- 100
MBFV-NC_015843.2    AGACCC---GGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAACGUCCC-------- 100
MBFV-NC_009026.2    AGACCC---GGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGGCGGCCAACG----- 100
MBFV-NC_007580.2    AAACCC---GGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGGCGUAUCCCGCG----- 100
MBFV-NC_009028.2    AAGUCC---GCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCUAGCGUGU----- 100
MBFV-NC_040776.1    GGACCC---GCCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAGCCAGCGUUU----- 100
ISFV-NC_024017.1    AAGCCC---GCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAGCGUG----- 100
MBFV-NC_002031.1    AAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUACGACGAGGAGUUCGCUCCUUGU-- 100
MBFV-NC_008719.1    AAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGUGACGGCCAAGGC---CA-------- 100
MBFV-NC_012735.1    AAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAGCGGCUAGAGU---GA-------- 100
ISFV-NC_027999.1    GGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUACUGG---AA-------- 100
ISFV-NC_040610.1    A------CCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUUAC---CACCCGCU-- 100
NKV-NC_005039.1     AAACACCCG-----AAUUUCGGGUGUCAAUAUAAAAAAGCAAAAUCCAAAAAACAAAA-- 100
ISFV-NC_016997.1    GCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC----------- 100
                    0........70........80........90........100.......110.......1

                    ..........................))))))))))))....))................
NKV-NC_004119.1     -NNNNNNNUUG-----CCUCAUCUUGAAUGGGGGGCGGCGUGGAUAUAUACUCCAGCCAG 154
NKV-NC_003635.1     GNNNNNNNUUU-----UCCAAAUUCCCGUUACCGGGGGUGGGGUGAUUAGCCAUGGUCGC 154
ISFV-NC_001564.2    ANNNNNNN--AGCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAACUCAUCU 156
ISFV-NC_005064.1    GNNNNNNN----CGUACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCCAGCU 154
ISFV-NC_012932.1    GNNNNNNN-UUGCGAACCCCCUCCG-AAAUGUGAAAAAUUAUGGGGAAAAGUACCCAUCU 157
ISFV-NC_034204.1    ANNNNNNN---ACGACCACCCUCUG-AAAUGUUAAUAAAACUGUGCGGAAAAAUCCAAAG 155
ISFV-NC_034017.1    ANNNNNNN---GCGACCACCCUCUG-UAAUGUUUAAUAAGUUGUGCGGAAAAAUCCAAAG 155
ISFV-NC_008604.2    GNNNNNNN-CUGACGCCACCUCCCG-AAAUGUGUAUAGUAACAGAGCAUGCCUGCAGCAG 157
ISFV-NC_021069.1    ANNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAG 156
ISFV-NC_012671.1    GNNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCAGCAG 156
ISFV-NC_024299.2    ANNNNNNN----ACGACACCCCCCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAUACAA 154
ISFV-NC_027819.1    ANNNNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAGAUGG 157
ISFV-NC_027817.1    GNNNNNNNAGUAGCGUCACCCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAAAGUC 157
TBFV-NC_003690.1    -NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGUCAGAGGGGGGGCGGUU 158
TBFV-NC_001809.1    -NNNNNNNGU-UUUUCCUCCUCCUAUACCAAAUUUCCCCCUCGACAGAGGGGGGGCGGUU 158
TBFV-NC_027709.1    -NNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGACAGAGGGGGGGCGGUU 158
TBFV-NC_001672.1    -NNNNNNN---------------UAUACAAAA-UUCCCCCUCGGUAGAGGGGGGGCGGUU 143
TBFV-NC_005062.1    -NNNNNNN---------------UAUACCAAA-UUCCCCCUCAUCUGAGGGGGGGCGGUU 143
TBFV-NC_003687.1    -NNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGAAACUGGGGGGGCGGUU 158
ISFV-NC_017086.1    -NNNNNNNCCC--AAAAAUGAAGGGCACUAUAUCGACACUUGGGAA--AGACCAGAGGUA 155
MBFV-NC_002640.1    ANNNNNNNCCCCCCAACACA-AAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGAUC 155
MBFV-NC_001474.2    -NNNNNNNCCCCCGAAACAA-AAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGAUC 155
MBFV-NC_001477.1    -NNNNNNNCCCGCACAACAA-CAAACAGCAUAUUGACG-CUGGGAG--AGACCAGAGAUC 155
MBFV-NC_001475.2    -NNNNNNNCCCCCGCAAAUA-AAAACAGCAUAUUGACG-CUGGGAG--AGACCAGAGAUC 155
MBFV-NC_012532.1    -NNNNNNNCCCCGGAAAACGCAAAACAGCAUAUUGACG--UGGGAA--AGACCAGAGACU 155
MBFV-NC_035889.1    -NNNNNNN-CCCGGAAAACGCAAAACAGCAUAUUGACG-CUGGGAA--AGACCAGAGACU 155
MBFV-NC_012533.1    CNNNNNNNA--UAACCAACAUAAAACAGCAUAUUGACACCUGGGAA--AGACCGGAGACU 155
MBFV-NC_001437.1    -NNNNNNN--------------------CAUAUUGACACCUGGGAA-UAGACUGGGAGAU 138
MBFV-NC_009029.2    -NNNNNNN-AGCGCAAAC-ACAAAACAGCAUAUUGACACCUGGG-A-AAGACUAGGGGAU 155
MBFV-NC_032088.1    -NNNNNNN-UACCGCAAA-ACAAAACAGCAUAUUGACACCUGGG-A-UAGACUAGGAGAU 155
MBFV-NC_000943.1    -NNNNNNN-AAAAGCAUC-AAACAACAGCAUAUUGACACCUGGGAA-AAGACUAGGAGAU 156
MBFV-NC_034151.1    -NNNNNNN-GAACGCAAA-UAAAAACAGCAUAUUGACACCUGGG-A-UAGACUGGGAGAC 155
MBFV-NC_001563.2    -NNNNNNN-AAACACCAA-AAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGGGAU 155
MBFV-NC_009942.1    -NNNNNNN-AACACCACA-ACAAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGAU 155
MBFV-NC_006551.1    -NNNNNNN-UAAGCAUCA-AAAAAACAGCAUAUUGACACCUGGGAAUUAGACUAGGAGAU 157
MBFV-NC_012534.1    -NNNNNNN-AACACCAAA-ACAAAACAGCAUAUUGACACCUGGGA--GAGACUAGGAGAC 155
MBFV-NC_015843.2    -NNNNNNN-GAACACCAC-AAGAAACAGCAUAUUGACACCUGGGA--UAGACUAGGAGAC 155
MBFV-NC_009026.2    -NNNNNNNACG-CACCAA-AAGAAACAGCAUAUUGACACCUGGGA--UAGACCGGAGAAU 155
MBFV-NC_007580.2    -NNNNNNNCCGUUAACGC-AAACAACAGCAUAUUGACACCUGGAA---AGACAGGAGAUC 155
MBFV-NC_009028.2    -NNNNNNNCAAGCGCGAA--ACAAACAGCAUAUUGACGCCUGGGA--AAGACCGGGAGAU 155
MBFV-NC_040776.1    -NNNNNNN-AAGCGCAAA-GACAAACAGCAUAUUGACACCUGGGA--AAGACCGGGAGAC 155
ISFV-NC_024017.1    -NNNNNNNCUGCUG-AGA-GCAAAAUAGCAUAUUGACACUUGGGAA-GAGACCAG-AGAU 155
MBFV-NC_002031.1    -NNNNNNN-AGGGAACAAAUAGUGGGACCAUAUUGACGCCAGGG-A-AAGACCGGAGUGG 156
MBFV-NC_008719.1    -NNNNNNN-CCGAGCAC-UAUAGCGGACCAUAUUGACGCCUGGG-A-AAGACCGGAGACA 155
MBFV-NC_012735.1    -NNNNNNN-CCGAGCACACAUAGCGGACCAUAUUGACGCCAGGG-A-AAGACCGGAGACA 156
ISFV-NC_027999.1    -NNNNNNN-CC-AUUGGAAAUUUAGCUUCAUAUUGACACCAGGGAA-AAGACCGGAGAAA 156
ISFV-NC_040610.1    -NNNNNNN-CUCAUUAUUUAUAUAACAGCAUAUUGACACCAGGG-A-UAGACCGGAGAAA 156
NKV-NC_005039.1     -NNNNNNN-UAC-GCAUGGAUUUUGCAAUAUGUUGACAUCAGGG-A-AAGACCGGGUGUU 155
ISFV-NC_016997.1    -NNNNNNNCAA--AAAUGUAUAGGCUAUUAUAUCGACACCUAGGAA--AGACUGGAGAUA 155
                    20.......130.......140.......150.......160.......170.......1

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NKV-NC_004119.1     AAAAGACUC----------------AGAU 167
NKV-NC_003635.1     ACAGAUCAA----------------GCUC 167
ISFV-NC_001564.2    UAACCUCCC----------------AC-- 167
ISFV-NC_005064.1    AAAUCUCCC----------------ACCG 167
ISFV-NC_012932.1    AAGCCUCCC----------------A--- 167
ISFV-NC_034204.1    ACUGUCCCC----------------ACC- 167
ISFV-NC_034017.1    AAUGUCCCC----------------ACC- 167
ISFV-NC_008604.2    CAGGUCUCC----------------A--- 167
ISFV-NC_021069.1    CAAAUCUCC----------------AC-- 167
ISFV-NC_012671.1    CAAAUCUCC----------------AC-- 167
ISFV-NC_024299.2    CAGGCUUCC----------------ACCG 167
ISFV-NC_027819.1    AAGAUCCCC----------------A--- 167
ISFV-NC_027817.1    GAGAGUCUC-------------------C 167
TBFV-NC_003690.1    CUUGUUCUC--------------------- 167
TBFV-NC_001809.1    CUUGUUCUC--------------------- 167
TBFV-NC_027709.1    CUUGUUCUC--------------------- 167
TBFV-NC_001672.1    CUUGUUCUCCCCU-----GAGCCACCAUCA 167
TBFV-NC_005062.1    CUUGUUCUCCCCU-----GAGCCACCAUCA 167
TBFV-NC_003687.1    CUUGUUCUC--------------------- 167
ISFV-NC_017086.1    CUCGCUGAUU-----------------CA 167
MBFV-NC_002640.1    CUGCUGUCUC-----------------UG 167
MBFV-NC_001474.2    CUGCUGUCUC-----------------CU 167
MBFV-NC_001477.1    CUGCUGUCUC-----------------UA 167
MBFV-NC_001475.2    CUGCUGUCUC-----------------CU 167
MBFV-NC_012532.1    CCAUGAGUUU-----------------CC 167
MBFV-NC_035889.1    CCAUGAGUUU-----------------CC 167
MBFV-NC_012533.1    CUGUUGGCUU-----------------UC 167
MBFV-NC_001437.1    CUUCUGCUCUAUCUCAACAUCAGCUACUA 167
MBFV-NC_009029.2    UUGCUGCUCU-----------------GG 167
MBFV-NC_032088.1    CCGCUGCUCU-----------------GG 167
MBFV-NC_000943.1    CUUCUGCUC------------------UA 167
MBFV-NC_034151.1    CCUCUGCUUA-----------------UU 167
MBFV-NC_001563.2    CUUCUGCUCU-----------------GC 167
MBFV-NC_009942.1    CUUCUGCUCU-----------------GC 167
MBFV-NC_006551.1    CUUCUGCUCU--------------------- 167
MBFV-NC_012534.1    CCUCUGUCCU-----------------AA 167
MBFV-NC_015843.2    CCUCUGUCCU-----------------AA 167
MBFV-NC_009026.2    UUGCUGCCUC-----------------GC 167
MBFV-NC_007580.2    CCCUGCUUUU-----------------UC 167
MBFV-NC_009028.2    CCUCUGCUUU-----------------CC 167
MBFV-NC_040776.1    CCUCUGCUCU-----------------GC 167
ISFV-NC_024017.1    AUGAUGUUGU-----------------UU 167
MBFV-NC_002031.1    UUCUCUGCUU-----------------U- 167
MBFV-NC_008719.1    CUCCUUGAUU-----------------CU 167
MBFV-NC_012735.1    CUCCUUGAUU-----------------C- 167
ISFV-NC_027999.1    CUUCUUGCUU-----------------A 167
ISFV-NC_040610.1    -UACACAAUU-----------------UC 167
NKV-NC_005039.1     UGUCGGUUCC-----------------GG 167
ISFV-NC_016997.1    CCUUUGGUUC-----------------UU 167
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