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                   ............................................................
ISFV-NC_027817.1   ---------CAAACUCUGGUGUCCAACGUUUGU-AUUAA---AAUGAGUG--GA-UUAGG 44
NKV-NC_004119.1    ---------CCGGCUGGAAACCACGUCGCUCUU-CGUCA---AAUGACGA--AAGGCAAA 45
NKV-NC_003635.1    ---------GUGCGCGUCACGCCACUUUUUGUG-GCGGG---AAUGUGCG--AAAAUAAC 45
ISFV-NC_001564.2   ---------GAGAGACCGUGAGGUUGAACCCGG-CAAGG---AAUGAAGAGGAAGGAU-U 46
ISFV-NC_005064.1   ---------AGUUGUUCGGAAAAGUGACCUUGG-UUCGU---UAUGAAUAGAGGAAGC-U 46
ISFV-NC_012932.1   ---------AGUUGUUUGGAAAAGAGAUAUCUG-GUUCG---UAUGAAUAGGAUGCUC-U 46
ISFV-NC_034204.1   --------GCGACAGAAGGAACCCAAAAAGGGA-ACCAG----AUGGAGCAAAACGGA-A 46
ISFV-NC_034017.1   --------UAAAAAUUUCUUUGGCGACGUACGG-AAAGU----AUGGAGCAAUCGAGG-A 46
ISFV-NC_024299.2   ---------UUGAAAAACUUUUGCGACGUUGUU-GAGAA---GAUGGAGCGAAAGAAAGG 47
ISFV-NC_008604.2   ---------UUGUUUAUAAUAGUCGUUUUUCUC-GCAGA---AAUGGGAAAGGACGACGG 47
ISFV-NC_021069.1   ---------CCUUAUUACCUCUGGUACUAAAGU-GGCAC---UAUGAAGAAGAAGGA--G 45
ISFV-NC_012671.1   ---------CUUUAUUACCUCUGGUACUAACGU-GGCGC---UAUGAAGAAGAAGGA--G 45
ISFV-NC_027819.1   ---------UCACGGAAUAGCUUGAGAAGAAGC-UUACU---UAUGUCAAGUAAGGAAGC 47
TBFV-NC_003687.1   --------AGCUAGGAGAACAAGAGCUGGGAG--UGGU---UAUG------AUGACCAC 41
TBFV-NC_005062.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGG---GAUGGCCGGGAAGGCCAU 47
TBFV-NC_001672.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGG---GAUGGUCAAGAAGGCCAU 47
TBFV-NC_001809.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGG---GAUGGGCAGGAAGACCAU 47
TBFV-NC_027709.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGG---GAUGGGCAGGAAGUCCAU 47
TBFV-NC_003690.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGG---AAUGGCCGGGAAGGCCGU 47
MBFV-NC_002031.1   -------------UUGAGCGAUUAGCAGAGAACUGACCAGAACAUGUCUG--------GU 39
NKV-NC_005039.1    -------------CGGAAGAACAACUGUCCGGAAUCAAAGACGAUGAAAA--------AU 39
ISFV-NC_017086.1   ---------UGAAUAGGUUUAUUGUGAAUCGAAAUACC----GAUGGCGA---AUA-AAC 43
ISFV-NC_024017.1   ---------UGAUAGAGUUGACUAGAUAACGAAUUAUC----GAUGGUGA---CUA-AAU 43
MBFV-NC_009028.2   ------UCUGAAU-AGUUUUUU--AGCAAGGGA-UUUCC---CAUGUCUA---AAA-AAC 43
MBFV-NC_040776.1   ------UCUGAAU-AGUUUUUU--AGCAGAGGU-ACUAC---CAUGUCUA---AAA-AAC 43
MBFV-NC_034151.1   --------UGAAU-AGUUUUUUG-AACAGAGGUGAAAAC---CAUGUCUA---AAA-AAC 43
MBFV-NC_000943.1   ------UG--AAC-AGUUUUUUGGAGCUUUUGA-UUUCA---AAUGUCUA---AAA-AAC 43
MBFV-NC_006551.1   -------GCCGGC-AGUUUCUUUGAGC-GUUGA-UUUUC---AAUGUCUA---AGA-AAC 43
MBFV-NC_001563.2   ------UGCGAGC-UGUUUCUUGGCAC-GAAGA--UCUC---GAUGUCUA---AGA-AAC 43
MBFV-NC_009942.1   ------UGCGAGC-UGUUUCUUAGCAC-GAAGA--UCUC---GAUGUCUA---AGA-AAC 43
MBFV-NC_009029.2   ACACAGUGUGAACAAGUUUUUUG-AACAG---------C---AAUGACCA---AAA-AAC 43
MBFV-NC_032088.1   ----AGUGUGAACAAGUUUUUUG-AUCAGUGGA-----U---AAUGACUA---AAA-AAC 43
MBFV-NC_007580.2   -------GCGAAC-AGUUUUUU--AGCAGGGAAUUACCC---AAUGUCUA---AAA-AAC 43
MBFV-NC_009026.2   -------UCGAG---GAAAGACGAGAAGCGAAUUCUCUG---AAUGGCUA---GAA-AAC 43
MBFV-NC_012534.1   ----AGUUUGAAC--GUUUUUUG-GAGAGAGA---UUAU---UAUGUUUAACAAAA-AAC 46
MBFV-NC_015843.2   ----AGUUUGAAC--GUUUUUUG-GAUAGAGA---CAAC---UAUGUCUAACAAAA-AAC 46
ISFV-NC_040610.1   ---------AAAGAAGUCAAAACUAGAAAUUUA-AUUUC---AAUGGUUA---GAC-AAC 43
MBFV-NC_001437.1   --------UUAAACAGUUUUUUAGAACGGAAGA--UAAC---CAUGACUA---AAA-AAC 43
MBFV-NC_002640.1   ------------UGUUUGAAUAGAGAGCAGAUC-UCUGGAAAAAUG---A--A--C-CAA 39
MBFV-NC_001474.2   -------ACAGUUUUUAAUUAGAGAGCAGAUC-UCUG-----AUGAAUA--A--C-CAA 42
MBFV-NC_001477.1   -------AACAGUUUUUUAUUAGAGAGCAGAUC-UCUG-----AUGAACA--A--C-CAA 42
MBFV-NC_001475.2   -------GACAGUUUUUUAUUAGAGAGCAGAUC-UCUG-----AUGAACA--A--C-CAA 42
MBFV-NC_012532.1   ----------AUUUGGAUUUGGAAACGAGAGUU-UCUGG--UCAUGAAAA--ACCC-CAA 44
MBFV-NC_035889.1   ----------UUUUGGAUUUGGAAACGAGAGUU-UCUGG--UCAUGAAAA--A-CC-CAA 43
MBFV-NC_012533.1   ----------GUGUUGAGAAAGAAAACAGAUUC-UUCGG--AAAUGAAAA--CCCA-AAA 44
ISFV-NC_027999.1   ------------UUGUUUGAGAGAACAGCGAGAAGUACUACU-AUGGUUGAGAAAACUAG 47
MBFV-NC_008719.1   ------------AAAGGGUAGAGAAAAG-GAGUUUUGCUUCUCAUGGCAACAAGAGGAGC 47
MBFV-NC_012735.1   ------------AAGGAGUAGAAGAAAG-GAGAUUCAUUCCCAAUGGCAACAAAGGGGAU 47
ISFV-NC_016997.1   ------------CAUUAGAAUAGGUUUGAAAGAGUUGAAUC-AAUGAGAUCAAUGGUCUU 47
                   .........10........20........30........40........50........6
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                   ...............................((((((.((((...................
ISFV-NC_027817.1   AGGCCUUCUUC-CCCU--CCGAGGGAAGAAAAAGAAAGCCCCUGUGAUACAAUCACAGG- 100
NKV-NC_004119.1    AUGAAAAACGC-GAAU--AAAAAGAAAGCAAAAAACUCUAAGACCAAUCCUGGU-CGAA- 100
NKV-NC_003635.1    AGGAAAAAGAA-CCCU--GGAAAGGGUUCAAAGAAGAAAAAGAAUGCUGGAAGG-AAUG- 100
ISFV-NC_001564.2   UGGAGG--CGCGAGG--AAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUGGGGAA-- 100
ISFV-NC_005064.1   UGAAGA--AGGGGCC--UCCGGGCCCAACAUCCAACCGUCCGUCGGUUGGACCGCCCG-- 100
ISFV-NC_012932.1   UGAAGA--AAGGGCC--UUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCGG-- 100
ISFV-NC_034204.1   AGUUUU--CCCGACCG-GGAGGUGGUCGGGCCCGGCCCAGAGAGGGGCUGG-GGAAGA-- 100
ISFV-NC_034017.1   AGUUUU--CCCGGCCG-GGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGG-CGAAGA-- 100
ISFV-NC_024299.2   AGUAAA--UUAUGGGA-GAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAAA-- 100
ISFV-NC_008604.2   UAAGAA--GAAGAAGG-GGCCGGGCUC-CUCCGGAUGGUUGUUACCACCGGGG-CGCG-- 100
ISFV-NC_021069.1   UCGGGA--GUAGUGGA-GAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAA-- 100
ISFV-NC_012671.1   UCUGGA--GUGGUGGA-GAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAG-- 100
ISFV-NC_027819.1   ACUGAA--GAAGAAGG-GAAUUUCCCUGGGAGGAGGAGGAAAACGGGGCUUUGAG--A-- 100
TBFV-NC_003687.1   UUCUAAAGGAAAGGGG-GGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAAGU 100
TBFV-NC_005062.1   UCUGAAAGGAAAGGGG-GGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA------ 100
TBFV-NC_001672.1   CCUGAAAGGUAAGGGG-GGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA------ 100
TBFV-NC_001809.1   CCUGAAAGGCAAGGGG-GGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA------ 100
TBFV-NC_027709.1   CCUGAAAGGUAAGGGG-GGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA------ 100
TBFV-NC_003690.1   UCUAAAAGGAAAGGGG-GGGGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCCAA------ 100
MBFV-NC_002031.1   CGUAAA--GCUCAGGG-AAAAACCCUGGGCGUCAAU-AUGGUACGACGAGGAGUUC--GC 93
NKV-NC_005039.1    CGUAGA--CAAAACAC-CCGAAUUUCGG--GUGUCA-AUAUAAAAAAGCAAAAUCCAAAA 93
ISFV-NC_017086.1   CCAAGA--AGCCCG-G--AAGACGGGCUAUCGAUAU-AGUGAGACGUGCGUUGCCUCGCG 97
ISFV-NC_024017.1   CAAGAA--AGCCCGCC--GCAAAGCGGGCUGUUAAU-AUGCUAACACGCAUCACCAAGCG 98
MBFV-NC_009028.2   CAGGAA--AGUCCGCU--GCAAAGCGGACUGUCAAU-AUGCUAAAACGCCUGGCUAGCGU 98
MBFV-NC_040776.1   CAGGAG--GACCCGCC--GGAAGGCGGGUUGUCAAU-AUGCUGAAACGCCCAGCCAGCGU 98
MBFV-NC_034151.1   CAGGAG--GACCCGGA--AAGAACCGGGUCGUUAAU-AUGCUGAAGCGCACAUCCCGCGC 98
MBFV-NC_000943.1   CAGGAG--GACCCGGG--AAGCCCCGGGUCGUCAAU-AUGCUAAAACGCGGCAUACCCCG 98
MBFV-NC_006551.1   CAGGAG--GGCCCGGA--AGAAACCGGGCCAUCAAU-AUGCUGAAACGCGGCAUACCCCG 98
MBFV-NC_001563.2   CAGGAG--GGCCCGGU--AAAAACCGGGCUGUCAAU-AUGCUAAAACGCGGUAUGCCCCG 98
MBFV-NC_009942.1   CAGGAG--GGCCCGGC--AAGAGCCGGGCUGUCAAU-AUGCUAAAACGCGGAAUGCCCCG 98
MBFV-NC_009029.2   CAGGAA--GGCCCGGU--AGAAAUCGGGCCGUCAAU-AUGCUAAAGCGCGGUGCGUCCCG 98
MBFV-NC_032088.1   CAGGAA--AGUCCGGC--AAUCGUCGGACUGUCAAU-AUGCUAAAGCGCGGAGCAUCCCG 98
MBFV-NC_007580.2   CAGGAA--AACCCGGU--AGAAACCGGGUUGUCAAU-AUGCUAAAACGCGGCGUAUCCCG 98
MBFV-NC_009026.2   CAGGAA--GACCCGGA--GGAAACCGGGUCGUCAAU-AUGCUAAAGCGUACGGCGGCCAA 98
MBFV-NC_012534.1   CAGGAA--AACCCGGG--GCAACCCGGGUUGUCAAU-AUGCUAAAGCGCGGUGCAUCCC- 100
MBFV-NC_015843.2   CAGGAA--GACCCGGC--UCAGGCCGGGUUGUCAAU-AUGCUAAAGCGCGGAACGUCCC- 100
ISFV-NC_040610.1   UUGGAA--AGACCGUU--GCAAGACGGUCUGUCAAU-AUGCGAAAGCGGAUUACCACCCG 98
MBFV-NC_001437.1   CAGGAG--GGCCCGGU--AAAAACCGGGCUAUCAAU-AUGCUGAAACGCGGCCUACCCCG 98
MBFV-NC_002640.1   CGAAAA--AAGGUGG---UUAGACCACCUUUCAAUA-UGCUGAAACGCGAGAGAAACCGC 93
MBFV-NC_001474.2   CGGAAA--AAGGCGA---AAAACACGCCUUUCAAUA-UGCUGAAACGCGAGAGAAACCGC 96
MBFV-NC_001477.1   CGGAAA--AAGACGG---GUCGACCGUCUUUCAAUA-UGCUGAAACGCGCGAGAAACCGC 96
MBFV-NC_001475.2   CGGAAG--AAGACGG---GAAAACCGUCUAUCAAUA-UGCUGAAACGCGUGAGAAACCGU 96
MBFV-NC_012532.1   AGAAGA--AAUCCGGA--GGA-UCCGGAUUGUCAAU-AUGCUAAAACGCGGAGUAGCCCG 98
MBFV-NC_035889.1   AAAAGA--AAUCCGGA--GGAUUCCGGAUUGUCAAU-AUGCUAAAACGCGGAGUAGCCCG 98
MBFV-NC_012533.1   AAGGGC--CGCAUCGC--GGCCUAUCAAUAUGCUGAAGCGCUUUGGAUC-UAGC-GCUGG 98
ISFV-NC_027999.1   AGUAGG--AUCUCGGGUUAAAACCCGAGGUGUCAAU-AUGAUUAGGCGUACUGGAA---- 100
MBFV-NC_008719.1   AAGCAA--GUCUCGGGUCACGACCCGGGGCGUCAAU-AUGGUGACGGCCAAGGCCA---- 100
MBFV-NC_012735.1   GAAUAA--GUCUCGGGCUCGAUCCCGAGGCGUCAAU-AUGGUAGCGGCUAGAGUGA---- 100
ISFV-NC_016997.1   UCGAGC--UAGGAGGCCCGUUAAACGGGCCGUCGAU-AUAAUAAAACGUAAGCUGC---- 100
                   0........70........80........90........100.......110.......1
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                   ..................................)))))))))).................
ISFV-NC_027817.1   -------NNNNNNN-AGUAGCGUCACCCCC---CGGUUGUGAAAACGAUU---GCGACUA 146
NKV-NC_004119.1    -------NNNNNNNUUGCCUCAUCUUGAAU---GGGGGGCGGCGUGGAUA---UAUACUC 147
NKV-NC_003635.1    -------NNNNNNNUUUUCCAAAUUCCCGU---UACCGGGGGUGGGGUGA---U-UAGCC 146
ISFV-NC_001564.2   -------NNNNNNN---AGCGAACCUCCUC---CGUUAAUG-UGAAAAAGUAUGGGG-AA 145
ISFV-NC_005064.1   -------NNNNNNN-----CGUACCCCCUC---CG-UAAUG-UGAAAAAGUAUGGGGAAA 143
ISFV-NC_012932.1   -------NNNNNNN--UUGCGAACCCCCUC---CG-AAAUG-UGAAAAAUUAUGGGGAAA 146
ISFV-NC_034204.1   -------NNNNNNN----ACGACCACCCUC---UGAAAUGU-U-AAUAAAACUGUGCGGA 144
ISFV-NC_034017.1   -------NNNNNNN----GCGACCACCCUC---UGUAAUGU-U-UAAUAAGUUGUGCGGA 144
ISFV-NC_024299.2   -------NNNNNNN-----ACGACACCCCC---CGAAAUGU-GUAUAUAGCCCAUGUUGA 144
ISFV-NC_008604.2   -------NNNNNNN--CUGACGCCACCUCC---CGAAAUGU-GUAUAGUAACAGAGCAUG 147
ISFV-NC_021069.1   -------NNNNNNN--GUGACGCCACCCCC---CGAAAUGU-GUAUUGUGACAG-GCUUG 146
ISFV-NC_012671.1   -------NNNNNNN--GUGACGCCACCCCC---CGAAAUGU-GUAUUGUGACAG-GCUUG 146
ISFV-NC_027819.1   -------NNNNNNN--UUAGCCACACCCCC---CGAAACGU-GAUUUAUAUGAUGACAAG 147
TBFV-NC_003687.1   -------NNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUU-CCCCCAGGAAACUGGGGG 151
TBFV-NC_005062.1   -------NNNNNNN---------------UAUACCAAA-UU-CCCCCUCAUCUGAGGGGG 136
TBFV-NC_001672.1   -------NNNNNNN---------------UAUACAAAA-UU-CCCCCUCGGUAGAGGGGG 136
TBFV-NC_001809.1   -------NNNNNNNG-UUUUUCCUCCUCCUAUACCAAAUUU-CCCCCUCGACAGAGGGGG 151
TBFV-NC_027709.1   -------NNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UU-CCCCCUCGACAGAGGGGG 151
TBFV-NC_003690.1   -------NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UU-CCCCCUCGUCAGAGGGGG 151
MBFV-NC_002031.1   UCCUUGUNNNNNNNAGGGAACAAAUAGUGG-GACCAUAUUGACGCCAGGGA--AAGACCG 150
NKV-NC_005039.1    AACAAAANNNNNNNUACGCAUGGAU-UUUG-CAAUAUGUUGACAUCAGGGA--AAGACC- 148
ISFV-NC_017086.1   UCU----NNNNNNN-CCCAAAAAUGAAGGG-CACUAUAUCGACACUUGGGAA-AGACCAG 150
ISFV-NC_024017.1   U-G----NNNNNNN-CUGCUGAGAGCAAAA-UAGCAUAUUGACACUUGGGAAGAGACCAG 151
MBFV-NC_009028.2   G-U----NNNNNNNCAAGCGCGAA-ACAAA-CAGCAUAUUGACGCCUGGGA--AAGACCG 149
MBFV-NC_040776.1   U-U----NNNNNNN-AAGCGCAAAGACAAA-CAGCAUAUUGACACCUGGGA--AAGACCG 149
MBFV-NC_034151.1   U-A----NNNNNNNGAA-CGCAAAUAAAAA-CAGCAUAUUGACACCUGGGA--UAGACUG 149
MBFV-NC_000943.1   C-G----NNNNNNN-AAAAGCAUCAAACAA-CAGCAUAUUGACACCUGGGAA-AAGACUA 150
MBFV-NC_006551.1   C-G----NNNNNNN-UAAGCAUCAAAAAAA-CAGCAUAUUGACACCUGGGAAUUAGACUA 151
MBFV-NC_001563.2   C-G----NNNNNNN-AAACACCAAAAGAAA-CAGCAUAUUGACACCUGGGA--UAGACUA 149
MBFV-NC_009942.1   C-G----NNNNNNN-AACACCACAACAAAA-CAGCAUAUUGACACCUGGGA--UAGACUA 149
MBFV-NC_009029.2   C-G----NNNNNNN-AGCGCAAACACAAAA-CAGCAUAUUGACACCUGGGA--AAGACUA 149
MBFV-NC_032088.1   C-G----NNNNNNN-UACCGCAAAACAAAA-CAGCAUAUUGACACCUGGGA--UAGACUA 149
MBFV-NC_007580.2   C-G----NNNNNNNCCGUUAACGCAAACAA-CAGCAUAUUGACACCUGGAA---AGAC-A 148
MBFV-NC_009026.2   C-G----NNNNNNNACGCACCAAAAGAAAC-AGCAUAUUGAC-ACCUGGGA--UAGA--- 146
MBFV-NC_012534.1   -------NNNNNNN-AACACCAAAACAAAA-CAGCAUAUUGACACCUGGGA--GAGACUA 149
MBFV-NC_015843.2   -------NNNNNNN-GAACACCACAAGAAA-CAGCAUAUUGACACCUGGGA--UAGACUA 149
ISFV-NC_040610.1   C-U----NNNNNNNCUCAUUAUUUAUAUAA-CAGCAUAUUGACACCAGGGA--UAGACC- 149
MBFV-NC_001437.1   C-G----NNNNNNN-CAUAUUGACACCUGG-GAAUAGACUGGGAGAUCUUC--UGCUCUA 149
MBFV-NC_002640.1   GUAUCAANNNNNNNCCCCCCAACACAAAAA--CAGCAUAUUGACGCUGGGA-----AAGA 146
MBFV-NC_001474.2   GUGU---NNNNNNNCCCCCGAAACAAAAAA--CAGCAUAUUGACGCUGGGA-----AAGA 146
MBFV-NC_001477.1   GUGU---NNNNNNNCCCGCACAACAACAAA--CAGCAUAUUGACGCUGGGA-----GAGA 146
MBFV-NC_001475.2   GUGU---NNNNNNNCCCCCGCAAAUAAAAA--CAGCAUAUUGACGCUGGGA-----GAGA 146
MBFV-NC_012532.1   UG-----NNNNNNNCCCCGGAAAACGCAAAACAGCAUAUUGAC-G-UGGGA-----AAGA 146
MBFV-NC_035889.1   UG-----NNNNNNN-CCCGGAAAACGCAAAACAGCAUAUUGAC-GCUGGGA-----AAGA 146
MBFV-NC_012533.1   CC-----NNNNNNNAUAACCAACAUAAAACAGCAUAUUGACAC--CUGGGA-----AAGA 146
ISFV-NC_027999.1   -------NNNNNNN-CC-AUUGGAAAUUUAGCUUCAUAUUGACACCAGGGAA-AAGACCG 150
MBFV-NC_008719.1   -------NNNNNNN-CCGAGCAC-UAUAGCGGACCAUAUUGACGCCUGGG-A-AAGACCG 149
MBFV-NC_012735.1   -------NNNNNNN-CCGAGCACACAUAGCGGACCAUAUUGACGCCAGGG-A-AAGACCG 150
ISFV-NC_016997.1   -------NNNNNNN-CAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGG--A-AAGACUG 149
                   20.......130.......140.......150.......160.......170.......1
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                   ...............................
ISFV-NC_027817.1   GAACUAAAGUCGAGAGU----------CUCC 167
NKV-NC_004119.1    CAGCCAGAAAAGACUCA----------GAU- 167
NKV-NC_003635.1    AUGGUCGCACAGAUCAA----------GCUC 167
ISFV-NC_001564.2   AGAACUCAUCUUAACCUC-------CCAC-- 167
ISFV-NC_005064.1   AGAACCCAGCUAAAUCUC-------CCACCG 167
ISFV-NC_012932.1   AGUACCCAUCUAAGCCUC-------CCA--- 167
ISFV-NC_034204.1   AAAAUCCAAAGACUGUCC-------CCACC- 167
ISFV-NC_034017.1   AAAAUCCAAAGAAUGUCC-------CCACC- 167
ISFV-NC_024299.2   GUCUAUACAACAGGCUU--------CCACCG 167
ISFV-NC_008604.2   CCUGCAGCAGCAGGUCU--------CCA--- 167
ISFV-NC_021069.1   CUCGCAGCAGCAAAUCU--------CCAC-- 167
ISFV-NC_012671.1   CUCGCAGCAGCAAAUCU--------CCAC-- 167
ISFV-NC_027819.1   AAUCAGAUGGAAGAUCC--------CCA--- 167
TBFV-NC_003687.1   GGCGGUUCUUGUUCUC--------------- 167
TBFV-NC_005062.1   GGCGGUUCUUGUUCUCCCUGAGCCACCAUCA 167
TBFV-NC_001672.1   GGCGGUUCUUGUUCUCCCUGAGCCACCAUCA 167
TBFV-NC_001809.1   GGCGGUUCUUGUUCUC--------------- 167
TBFV-NC_027709.1   GGCGGUUCUUGUUCUC--------------- 167
TBFV-NC_003690.1   GGCGGUUCUUGUUCUC--------------- 167
MBFV-NC_002031.1   GAGUGGUUCUCUGCUUU--------------- 167
NKV-NC_005039.1    --GGGUGUUUGUCGGUU----------CCGG 167
ISFV-NC_017086.1   AGGUACUCGCUGAUUCA--------------- 167
ISFV-NC_024017.1   AGAUAUGAUGUUGUUU--------------- 167
MBFV-NC_009028.2   GGAGAUCCUCUGC-UUU----------C--C 167
MBFV-NC_040776.1   GGAGACCCUCUGC-UCU----------G--C 167
MBFV-NC_034151.1   GGAGACCCUCUGC-UUA----------U--U 167
MBFV-NC_000943.1   GGAGAUCUUCUGC-UCU----------A--- 167
MBFV-NC_006551.1   GGAGAUCUUCUG---CU----------C--U 167
MBFV-NC_001563.2   GGGGAUCUUCUGC-UCU----------G--C 167
MBFV-NC_009942.1   GGAGAUCUUCUGC-UCU----------G--C 167
MBFV-NC_009029.2   GGGGAUUUGCUGC-UCU----------G--G 167
MBFV-NC_032088.1   GGAGAUCCGCUGC-UCU----------G--G 167
MBFV-NC_007580.2   GGAGAUCCCCUGCUUUU----------U--C 167
MBFV-NC_009026.2   CCGGAGAAUUUGCUGCC----------UCGC 167
MBFV-NC_012534.1   GGAGACCCUCUG---UC----------CUAA 167
MBFV-NC_015843.2   GGAGACCCUCUG---UC----------CUAA 167
ISFV-NC_040610.1   GGAGAAAUACACAAUUU-------------C 167
MBFV-NC_001437.1   UCUCAACAUCAGCUACU-------------A 167
MBFV-NC_002640.1   CCAGAGAUCCUGCUGUC----------UCUG 167
MBFV-NC_001474.2   CCAGAGAUCCUGCUGUC----------UCCU 167
MBFV-NC_001477.1   CCAGAGAUCCUGCUGUC----------UCUA 167
MBFV-NC_001475.2   CCAGAGAUCCUGCUGUC----------UCCU 167
MBFV-NC_012532.1   CCAGAGACUCCAUGAGU----------UUCC 167
MBFV-NC_035889.1   CCAGAGACUCCAUGAGU----------UUCC 167
MBFV-NC_012533.1   CCGGAGACUCUGUUGGC----------UUUC 167
ISFV-NC_027999.1   GAGAAACUUCUUGCUUA--------------- 167
MBFV-NC_008719.1   GAGACACUCCUUGAUUC----------U--- 167
MBFV-NC_012735.1   GAGACACUCCUUGAUUC--------------- 167
ISFV-NC_016997.1   GAGAUACCUUUGGUUCU----------U--- 167
                   80.......190.......200.........
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