```
                   ............................................................
ISFV-NC_001564.2   ------GAGAGACCGUGAGGUUGAACCCGGCAA----GGAAUG-------------AAGA 37
ISFV-NC_005064.1   ------AGUUGUUCGGAAAAGUGACCUUGGUUC----GUUAUG-------------AAUA 37
ISFV-NC_012932.1   ------AGUUGUUUGGAAAAGAGAUAUCUGGUU----CGUAUG-------------AAUA 37
ISFV-NC_034204.1   ------GCGACAGAAGGAACCCAAAAAGGGAAC----CAGAUG-------------GAGC 37
ISFV-NC_034017.1   ------UAAAAAUUUCUUUGGCGACGUACGGAA----AGUAUG-------------GAGC 37
ISFV-NC_008604.2   ------UUGUUUAUAAUAGUCGUUUUUCUCGCA----GAAAUG-------------GGAA 37
ISFV-NC_021069.1   ------CCUUAUUACCUCUGGUACUAAAGUGGC----ACUAUG-------------AAGA 37
ISFV-NC_012671.1   ------CUUUAUUACCUCUGGUACUAACGUGGC----GCUAUG-------------AAGA 37
ISFV-NC_024299.2   ------UUGAAAAACUUUUGCGACGUUGUUGAG----AAGAUG-------------GAGC 37
ISFV-NC_027819.1   ------UCACGGAAUAGCUUGAGAAGAAGCUUA----CUUAUG-------------UCAA 37
ISFV-NC_027817.1   ------CAAACUCUGGUGUCCAACGUUUGUAUU----AAAAUG-------------AGUG 37
NKV-NC_004119.1    ------CCGGCUGGAAACCACGUCGCUCUUCGU----CAAAUG-------------ACGA 37
NKV-NC_003635.1    ----------GUGCGCGUCACGCCACUUUUGUGGCGGGAAUG-------------UGCG 37
MBFV-NC_001437.1   -------UUAAACAGUUUUUAGAACGGAAGAUAACC---AUG---------ACUAAAAA 41
MBFV-NC_012534.1   ----AGUUUGAACGUUUUUUGGAGAGAGAUUAUU------AUGUUU------AACAAAAA 44
MBFV-NC_015843.2   ----AGUUUGAACGUUUUUUGGAUAGAGACAACU------AUGUCU------AACAAAAA 44
MBFV-NC_009029.2   ACACAGUGUGAACAAGUUUUUUGAACAG----CA------AUG---------ACCAAAAA 41
MBFV-NC_032088.1   ----AGUGUGAACAAGUUUUUUGAUCAGUGGAUA------AUG---------ACUAAAAA 41
MBFV-NC_007580.2   -------GCGAACAGUUUUUUAG--CAGGGAAUUACCC-AAUGU---------CUAAAAA 41
MBFV-NC_000943.1   ------U--GAACAGUUUUUUGGAGCUUUUGAUU-UCA-AAUGU---------CUAAAAA 41
MBFV-NC_001563.2   ------UGCGAGCUGUUUCUUGGCAC-GAAGA-U-CUC-GAUGU---------CUAAGAA 41
MBFV-NC_009942.1   ------UGCGAGCUGUUUCUUAGCAC-GAAGA-U-CUC-GAUGU---------CUAAGAA 41
MBFV-NC_006551.1   ------GCCG-GCAGUUUCUUUGAGC-GUUGAUU-UUC-AAUGU---------CUAAGAA 41
MBFV-NC_009026.2   ------UCGAGGAAA--GACGAG--AAGCGAAUUCUCUGAAUGG---------CUAGAAA 41
MBFV-NC_009028.2   ------UCUGAAUAGUUUUUUAG--CAAGGGAUUUCCC--AUGU---------CUAAAAA 41
MBFV-NC_040776.1   ------UCUGAAUAGUUUUUUAG--CAGAGGUACUACC--AUGU---------CUAAAAA 41
MBFV-NC_034151.1   ------U--GAAUAGUUUUUU-GAACAGAGGUGAAAACC-AUGU---------CUAAAAA 41
MBFV-NC_001477.1   ------AACAGUUUUUUAUUAGAGAGCAG-AUCUCU---GAUGAAC------AA--CCAA 42
MBFV-NC_001475.2   ------GACAGUUUUUUAUUAGAGAGCAG-AUCUCU---GAUGAAC------AA--CCAA 42
MBFV-NC_001474.2   ------ACAGUUUUUUAAUUAGAGAGCAG-AUCUCU---GAUGAAU------AA--CCAA 42
MBFV-NC_002640.1   ---------UGUUUGAAUAGAGAGCAGAU-CUCUGGAAAAAUG---------AA--CCAA 39
MBFV-NC_012533.1   ----------GUGUUGAGAAAGAAAACAGAUUCUUCGGAAAUGAAA------AC--CCAA 42
MBFV-NC_012532.1   ------A----UUUGGAUUUGGAAACGAGAGUUUCUGGUCAUGAAA------AACCCCAA 44
MBFV-NC_035889.1   ------U----UUUGGAUUUGGAAACGAGAGUUUCUGGUCAUGAAA------AA-CCCAA 43
ISFV-NC_040610.1   ------AAAGAAGUCAAAACUAGAAAUUUAAUU-UC---AAUGG---------UUAGACA 41
ISFV-NC_024017.1   ------UGAU-AGAGUUGACUAGAUAACGAAUUAUC---GAUGG---------UGACUAA 41
ISFV-NC_017086.1   ------UGAAUAGGUUUAUUGUGAAUCGAAAUACC----GAUG---------GCGAAUAA 41
ISFV-NC_016997.1   ------CAUUAGAAUAGGUUUGAAAGAGUUGAAUC----AAUGAGAUCAAUGGUCUUUCG 50
MBFV-NC_002031.1   ------UUG----AGCGAUUAGCAGAGAACUGACCAGAACAUGUC------------UGG 38
ISFV-NC_027999.1   ------UUGUUUGAGAGAACAGCGAGAAGUACUAC---U-AUGGUUGAG---AAAACUAG 47
MBFV-NC_012735.1   ------AAGGAGUAGAAGAAAG-GAGAUUCAUUCC---CAAUGGCAACA---AAGGGGAU 47
MBFV-NC_008719.1   ------AAAGGGUAGAGAAAAG-GAGUUUUGCUUC---UCAUGGCAACA---AGAGGAGC 47
NKV-NC_005039.1    C----GGAAGAACAACUGUCCGGAAUCAAAGAC------GAUG------------AAAAA 38
TBFV-NC_003687.1   ----------AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG------AUGACCACUUC 44
TBFV-NC_001672.1   ----AAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGGCCAUCCU 50
TBFV-NC_005062.1   ----AAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGCCAUUCU 50
TBFV-NC_027709.1   ----AAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGUCCAUCCU 50
TBFV-NC_001809.1   ----AAAGACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGACCAUCCU 50
TBFV-NC_003690.1   ----AAAGACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGCCGUUCU 50
                   .........10........20........30........40........50........6
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                   ..............................(((((((((((....................
ISFV-NC_001564.2   GGAAGG--AUUUGGAGGCGCGAGGAAAAGCGCCAGGGAGGGAUUUUUCCACCCCGUUCUG 95
ISFV-NC_005064.1   GAGGAA--GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGGUUGGACC 95
ISFV-NC_012932.1   GGAUGC--UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACC 95
ISFV-NC_034204.1   AAAACG--GAAAGUUUUCCCGACCGGGAGGUGGUCGGGCCCGGCCCAGAGAGGGGCUGGG 95
ISFV-NC_034017.1   AAUCGA--GGAAGUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGGC 95
ISFV-NC_008604.2   AGGACGACGGUAAGAAGAAGAAGGGGCCGGGCUCCUCCGGAUGGUUGUUACCA--CCGGG 95
ISFV-NC_021069.1   AGAAGGA--GUCGGGAGUAGUGGAGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGG 95
ISFV-NC_012671.1   AGAAGGA--GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGG 95
ISFV-NC_024299.2   GAAAGAAAGGAGUAAAUUAUGGGAGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGA 95
ISFV-NC_027819.1   GUAAGGAAGCACUGAAGAAGAAGGGAAUUUCCCUGGGAGGAGGAGGAAAACGGGGCUUUG 97
ISFV-NC_027817.1   GAUUAGGAGGCCUUCUUCCCCUCCGAGGGAAGAA--AAAGAAAGCCCCUGUGAUACAAUC 95
NKV-NC_004119.1    AAGGCAAAAUGAAAAACGCGAAUAAAAAGAAAGC--AAAAAACUCUAAGACCAAUCCUGG 95
NKV-NC_003635.1    AAAAUAACAGGAAAAAGAACCCUGGAAAGGGUUC--AAAGAAGAAAAAGAAUGCUGGAAG 95
MBFV-NC_001437.1   ACCAGGAGGGCCCGGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCCUACCCCGCG- 100
MBFV-NC_012534.1   ACCAGGAAAACCCGGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUGCAUCCC---- 100
MBFV-NC_015843.2   ACCAGGAAGACCCGGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAACGUCCC---- 100
MBFV-NC_009029.2   ACCAGGAAGGCCCGGUAGAAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUGCGUCCCGCG- 100
MBFV-NC_032088.1   ACCAGGAAAGUCCGGCAAUCGUCGGACUGUCAAUAUGCUAAAGCGCGGAGCAUCCCGCG- 100
MBFV-NC_007580.2   ACCAGGAAAACCCGGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGGCGUAUCCCGCG- 100
MBFV-NC_000943.1   ACCAGGAGACCCGGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGGCAUACCCCGCG- 100
MBFV-NC_001563.2   ACCAGGAGGGCCCGGUAAAAACCGGGCUGUCAAUAUGCUAAAACGCGGUAUGCCCCGCG- 100
MBFV-NC_009942.1   ACCAGGAGGGCCCGGCAAGAGCCGGGCUGUCAAUAUGCUAAAACGCGGAAUGCCCCGCG- 100
MBFV-NC_006551.1   ACCAGGAGGGCCCGGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGGCAUACCCCGCG- 100
MBFV-NC_009026.2   ACCAGGAAGACCCGGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGGCGGCCAACG- 100
MBFV-NC_009028.2   ACCAGGAAAGUCCGCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCUAGCGUGU- 100
MBFV-NC_040776.1   ACCAGGAGGACCCGCCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAGCCAGCGUUU- 100
MBFV-NC_034151.1   ACCAGGAGGACCCGGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGCACAUCCCGCGCUA- 100
MBFV-NC_001477.1   CGGAAAAAGA--CG--GGUCGACCGUCUUUCAAUAUGCUGAAACGCGCGAGAAACCGCGU 98
MBFV-NC_001475.2   CGGAAGAAGA--CG--GGAAAACCGUCUAUCAAUAUGCUGAAACGCGUGAGAAACCGUGU 98
MBFV-NC_001474.2   CGGAAAAAGG--CG--AAAAACACGCCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGU 98
MBFV-NC_002640.1   CGAAAAAAGG--UG--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGAGAAACCGCGU 95
MBFV-NC_012533.1   ---AAAAGGG--CC--GCAUCGCGGCCUAUCAAUAUGCUGAAGCGCUUUGGAUCUAGCGC 95
MBFV-NC_012532.1   AGAAGAAAUC--CGGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG- 100
MBFV-NC_035889.1   AAAAGAAAUC--CGGAGGAUUCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCCCGUG- 100
ISFV-NC_040610.1   ACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUACCACCCGCU- 100
ISFV-NC_024017.1   AUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAGCGUG- 100
ISFV-NC_017086.1   ACCCAAGAAGCCCGGAAGA---CGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCGCGU 98
ISFV-NC_016997.1   AGCUAGGAGGCCCGUUAAA---CGGGCCGUCGAUAUAAUAAAACGUAAGCUGC------- 100
MBFV-NC_002031.1   UCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUACGACGAGGAGUUCGCUCCUU 98
ISFV-NC_027999.1   AGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCG--UACUGG------- 98
MBFV-NC_012735.1   GAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAG--CGGCUAGAGU------- 98
MBFV-NC_008719.1   AAGCAAGUCUCGGG-UC-ACGACCCGGGGCGUCAAUAUGGUGACGGCCAAGGC------- 98
NKV-NC_005039.1    UCGUAGACAAAACACCCGAAUUUCGG--GUGUCAAUAUAAAAAAGCAAAAU-CCAAAAAA 95
TBFV-NC_003687.1   UAAAGGAAAGGGGGGCGGUCCCCC--UAGGCGCAAGCUUAAAGUGACCGCAAAUAAGU-- 100
TBFV-NC_001672.1   GAAAGGUAAGGGGGGCGGUCCCCC--UCGACGAGUGUCGAAAGAGACCGCAA-------- 100
TBFV-NC_005062.1   GAAAGGAAAGGGGGGCGGUCCCCC--UCGACGAGUGUCGAAAGAGACCGCUA-------- 100
TBFV-NC_027709.1   GAAAGGUAAGGGGGGCGGUCCCCC--UCGACGAGUGUCGAAAGAGACCGCAA-------- 100
TBFV-NC_001809.1   GAAAGGCAAGGGGGGCGGUCCCCC--UCGACGAGUGUCGAAAGAGACCGCGA-------- 100
TBFV-NC_003690.1   AAAAGGAAAGGGGGGGGGUCCCCC--UCGACGAGCGUCGAAAGUGGCCCCAA-------- 100
                   0........70........80........90........100.......110.......1
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                   ...............................))))))))))).................
ISFV-NC_001564.2   GGGAANNNNNNN--AGCGAACCUCCUCCGUUAAUGUGAAAAAGUAUGGGG-AAAGAACUC 152
ISFV-NC_005064.1   GCCCGNNNNNNN----CGUACCCCCUCCG-UAAUGUGAAAAAGUAUGGGGAAAAGAACCC 150
ISFV-NC_012932.1   GCCGGNNNNNNN-UUGCGAACCCCCUCCG-AAAUGUGAAAAAUUAUGGGGAAAAGUACCC 153
ISFV-NC_034204.1   GAAGANNNNNNN---ACGACCACCCUCUG-AAAUGUUAAUAAAACUGUGCGGAAAAAUCC 151
ISFV-NC_034017.1   GAAGANNNNNNN---GCGACCACCCUCUG-UAAUGUUUAAUAAGUUGUGCGGAAAAAUCC 151
ISFV-NC_008604.2   GCGCGNNNNNNN-CUGACGCCACCUCCCG-AAAUGUGUAUAGUAACAGAGCAUGCCUGCA 153
ISFV-NC_021069.1   AGGAANNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCA 152
ISFV-NC_012671.1   AGGAGNNNNNNN-GUGACGCCACCCCCCG-AAAUGUGUAUUGUGACAG-GCUUGCUCGCA 152
ISFV-NC_024299.2   GAAAANNNNNNN----ACGACACCCCCCG-AAAUGUGUAUAUAGCCCAUGUUGAGUCUAU 150
ISFV-NC_027819.1   AG--ANNNNNNN-UUAGCCACACCCCCCG-AAACGUGAUUUAUAUGAUGACAAGAAUCAG 153
ISFV-NC_027817.1   ACAGGNNNNNNNAGUAGCGUCACCCCCCGGU--UGUGAAAACGAUUGCGACUAGAACUAA 153
NKV-NC_004119.1    UCGAANNNNNNNUUGCCUCA---UCUUGAAUGGGGGGCGGCGUGGAUAUAUACUCCAGC- 151
NKV-NC_003635.1    GAAUGNNNNNNNUUUUCCAAAUUCCCGUUACCGGGGGUGGGGUGAUUAGCCAUGGUCGC- 154
MBFV-NC_001437.1   -----NNNNNNN-----------------CAUAU--UGACAC-CUGGGA-AUAGACUGG 133
MBFV-NC_012534.1   -----NNNNNNNAAC-ACCAAAACAAAA-CAGCAUAUUGACAC-CUGGGA--GAGACUAG 150
MBFV-NC_015843.2   -----NNNNNNNGAA-CACCACAAGAAA-CAGCAUAUUGACAC-CUGGGA--UAGACUAG 150
MBFV-NC_009029.2   -----NNNNNNNAGC-GCAAACACAAAA-CAGCAUAUUGACAC-CUGGGA--AAGACUAG 150
MBFV-NC_032088.1   -----NNNNNNNUAC-CGCAAAACAAAA-CAGCAUAUUGACAC-CUGGGA--UAGACUAG 150
MBFV-NC_007580.2   -----NNNNNNNCCGUUAACGCAAACAA-CAGCAUAUUGACAC-CUGG---AAAGAC-AG 149
MBFV-NC_000943.1   -----NNNNNNNAAAAGCAU-CAAACAA-CAGCAUAUUGACAC-CUGGGAAA-AGACUAG 151
MBFV-NC_001563.2   -----NNNNNNNAAACACCA-AAAGAAA-CAGCAUAUUGACAC-CUGGGA--UAGACUAG 150
MBFV-NC_009942.1   -----NNNNNNNAACACCAC-AACAAAA-CAGCAUAUUGACAC-CUGGGA--UAGACUAG 150
MBFV-NC_006551.1   -----NNNNNNNUAAGCAUC-AAAAAAA-CAGCAUAUUGACAC-CUGGGAAUUAGACUAG 152
MBFV-NC_009026.2   -----NNNNNNN-ACGCACCAAAAGAAA-CAGCAUAUUGACAC-CUGGGA--UAGACCGG 150
MBFV-NC_009028.2   -----NNNNNNNCAAGCGCGAA-ACAAA-CAGCAUAUUGACGC-CUGGGA--AAGACCGG 150
MBFV-NC_040776.1   -----NNNNNNN-AAGCGCAAAGACAAA-CAGCAUAUUGACAC-CUGGGA--AAGACCGG 150
MBFV-NC_034151.1   -----NNNNNNNGAA-CGCAAAUAAAAA-CAGCAUAUUGACAC-CUGGGA--UAGACUGG 150
MBFV-NC_001477.1   GU---NNNNNNNCCCGCACAACAACAAA-CAGCAUAUUGA--CGCUGGGA--GAGACCAG 150
MBFV-NC_001475.2   GU---NNNNNNNCCCCCGCAAAUAAAAA-CAGCAUAUUGA--CGCUGGGA--GAGACCAG 150
MBFV-NC_001474.2   GU---NNNNNNNCCCCCGAAACAAAAAA-CAGCAUAUUGA--CGCUGGGA--AAGACCAG 150
MBFV-NC_002640.1   AUCAANNNNNNNCCCCCC--AACACAAA-AACAGCAUAUUGACGCUGGGA--AAGACCAG 150
MBFV-NC_012533.1   UGGCCNNNNNNNAUAAC-CAACAUAAAA-CAGCAUAUUGACAC-CUGGGA--AAGACCGG 150
MBFV-NC_012532.1   -----NNNNNNNCCCCGGAAAACGCAAAACAGCAUAUUGACG---UGGGA--AAGACCAG 150
MBFV-NC_035889.1   -----NNNNNNN-CCCGGAAAACGCAAA--ACAGCAUAUUGACGCUGGGA--AAGACCAG 150
ISFV-NC_040610.1   -----NNNNNNNCUCAUUAUUUAUAUAA-CAGCAUAUUGACACCAGGG-A--UAGACCGG 151
ISFV-NC_024017.1   -----NNNNNNNCUGCUGAGAG-CAAAA-UAGCAUAUUGACACUUGGGAA--GAGACCAG 151
ISFV-NC_017086.1   CU---NNNNNNNCCC--AAAAAUGAAGGGCACUAUAUCGACACUUGGGAA---AGACCAG 150
ISFV-NC_016997.1   -----NNNNNNNCAA--AAAUGUAUAGGCUAUUAUAUCGACACCUAGGAA---AGACUGG 150
MBFV-NC_002031.1   GU---NNNNNNNA-GGGAACAAAUAGUGGGACCAUAUUGACGCCAGGG-A--AAGACCGG 151
ISFV-NC_027999.1   AA---NNNNNNNC-C-AUUGGAAAUUUAGCUUCAUAUUGACACCAGGGAA--AAGACCGG 151
MBFV-NC_012735.1   GA---NNNNNNNC-CGAGCACACAUAGCGGACCAUAUUGACGCCAGGG-A--AAGACCGG 151
MBFV-NC_008719.1   CA---NNNNNNNC-CGAGCAC-UAUAGCGGACCAUAUUGACGCCUGGG-A--AAGACCGG 150
NKV-NC_005039.1    CAAAANNNNNNNUACGCAUGGAUUUU--GCAAUAUGUUGACAUCAGGGAA---AGACCGG 150
TBFV-NC_003687.1   -----NNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGA----AACUGGGG 150
TBFV-NC_001672.1   -----NNNNNNN---------------UAUACAAAA-UUCCCCCUCGG----UAGAGGGG 135
TBFV-NC_005062.1   -----NNNNNNN---------------UAUACCAAA-UUCCCCCUCAU----CUGAGGGG 135
TBFV-NC_027709.1   -----NNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGA----CAGAGGGG 150
TBFV-NC_001809.1   -----NNNNNNNG-UUUUUCCUCCUCCUAUACCAAAUUUCCCCCUCGA----CAGAGGGG 150
TBFV-NC_003690.1   -----NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCCUCGU----CAGAGGGG 150
                   20.......130.......140.......150.......160.......170.......1
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                   ..................................
ISFV-NC_001564.2   AUCUUAACCUCCCAC------------------ 167
ISFV-NC_005064.1   AGCUAAAUCUCCCAC-----------------CG 167
ISFV-NC_012932.1   AUCUAAGCCUCCCA-------------------- 167
ISFV-NC_034204.1   AAAGACUGUCCCCAC-----------------C- 167
ISFV-NC_034017.1   AAAGAAUGUCCCCAC-----------------C- 167
ISFV-NC_008604.2   GCAGCAGGUCUCCA-------------------- 167
ISFV-NC_021069.1   GCAGCAAAUCUCCAC------------------- 167
ISFV-NC_012671.1   GCAGCAAAUCUCCAC------------------- 167
ISFV-NC_024299.2   ACAACAGGCUUCCAC-----------------CG 167
ISFV-NC_027819.1   AUGGAAGAUCCCCA-------------------- 167
ISFV-NC_027817.1   AGUCGAGAGUCUC------------------C 167
NKV-NC_004119.1    -CAGAAAAGACUCAG-----------------AU 167
NKV-NC_003635.1    ACAGAUCAAGCUC----------------------- 167
MBFV-NC_001437.1   GAGAUCUUCUGCUCUAUCUCAACAUCAGCUACUA 167
MBFV-NC_012534.1   GAGACCCUCUGU-CC----------------UAA 167
MBFV-NC_015843.2   GAGACCCUCUGU-CC----------------UAA 167
MBFV-NC_009029.2   GGGAUUUGCUGC-UC----------------UGG 167
MBFV-NC_032088.1   GAGAUCCGCUGC-UC----------------UGG 167
MBFV-NC_007580.2   GAGAUCCCCUGCUUU----------------UUC 167
MBFV-NC_000943.1   GAGAUCUUCUGC-UC----------------UA- 167
MBFV-NC_001563.2   GGGAUCUUCUGC-UC----------------UGC 167
MBFV-NC_009942.1   GAGAUCUUCUGC-UC----------------UGC 167
MBFV-NC_006551.1   GAGAUCUUCUGC-UC----------------U-- 167
MBFV-NC_009026.2   AGAAUUUGCUGC-CU----------------CGC 167
MBFV-NC_009028.2   GAGAUCCUCUGC-UU----------------UCC 167
MBFV-NC_040776.1   GAGACCCUCUGC-UC----------------UGC 167
MBFV-NC_034151.1   GAGACCCUCUGCUUA----------------U-U 167
MBFV-NC_001477.1   AGAUCCUGCUGUCUC-----------------UA 167
MBFV-NC_001475.2   AGAUCCUGCUGUCUC-----------------CU 167
MBFV-NC_001474.2   AGAUCCUGCUGUCUC-----------------CU 167
MBFV-NC_002640.1   AGAUCCUGCUGUCUC-----------------UG 167
MBFV-NC_012533.1   AGACUCUGUUGGCUU-----------------UC 167
MBFV-NC_012532.1   AGACUCCAUGAGUUU-----------------CC 167
MBFV-NC_035889.1   AGACUCCAUGAGUUU-----------------CC 167
ISFV-NC_040610.1   AGAAAUACACAAUUU------------------C 167
ISFV-NC_024017.1   AGAUAUGAUGUUGUU------------------U 167
ISFV-NC_017086.1   AGGUACUCGCUGAUU-----------------CA 167
ISFV-NC_016997.1   AGAUACCUUUGGUUC-----------------UU 167
MBFV-NC_002031.1   AGUGGUUCUCUGCUU------------------U 167
ISFV-NC_027999.1   AGAAACUUCUUGCUU------------------A 167
MBFV-NC_012735.1   AGACACUCCUUGAUU-----------------C- 167
MBFV-NC_008719.1   AGACACUCCUUGAUU-----------------CU 167
NKV-NC_005039.1    GUGUUUGUCGGUUCC-----------------GG 167
TBFV-NC_003687.1   GGGCGGUUCUUGUUC--UC--------------- 167
TBFV-NC_001672.1   GGGCGGUUCUUGUUC--UCCCUGAGCCACCAUCA 167
TBFV-NC_005062.1   GGGCGGUUCUUGUUC--UCCCUGAGCCACCAUCA 167
TBFV-NC_027709.1   GGGCGGUUCUUGUUC--UC--------------- 167
TBFV-NC_001809.1   GGGCGGUUCUUGUUC--UC--------------- 167
TBFV-NC_003690.1   GGGCGGUUCUUGUUC--UC--------------- 167
                   80.......190.......200.......210..
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