```
                     ................10........20........30........40........50......6
ISFV-NC_034204.1     -----GCGACAGAAGGAACCCAAAAAGGGAACC---AGAUGGAGCAAAACG----GAAA   47
ISFV-NC_034017.1     -----UAAAAAUUUCUUUGUGCGACGUGUACGGAAA--AGAUGGAGCAAUCGA---GGAA   45
ISFV-NC_024299.2     -----UUGAAAAACUUUUGCGACGUUGUUGAGA---AGAUGGAGCGAA-A----GAAA   45
ISFV-NC_027817.1     -----CAAACUCUGGUGUCACGUUUGUAUUA--AAUAG--AGUGGUGUUAA---AAGC   47
ISFV-NC_027819.1     -----UCACGGAAUAGUUUAGAGAGAGAGCUUAC--UUAUGUCAAGUUAGG---AAGC   47
ISFV-NC_008604.2     -----UUGUUUAUAAUAGUCGUUUUUCUCGCAG--AAAUGGGAAAGGACG----ACGG   47
ISFV-NC_021069.1     -----CCUUAUUACCUCUGGUACAAAGUGGCG---CUAUGAAGAAGAAGG----A--G   45
ISFV-NC_012671.1     -----CUUUAUUACCUCUGGUACUAAACUGGCG---CUAUGAAGAAGAAGG----A--G   45
ISFV-NC_001564.2     -----GAGAGACCGUGAGGUUGACCGGCAAG---GAAUGAAGAAGAAGG----A--G   44
ISFV-NC_005064.1     -----AGUUGUCUUGGAAAAGUGACCUUUGGUUCUC--UUAUGAAUAAGAAGGAA---U--G   44
ISFV-NC_012932.1     -----AGUUGUUUGUGAAAAGAUUAUCUUGGUUC----GUAUGAAUAGGAUGUC---U----U   44
TBFV-NC_001809.1     AA---AGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAAGACCAUCCUG   51
TBFV-NC_027709.1     AA---AGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAAGGCCAUCCUG   51
TBFV-NC_003690.1     AA---AGACAGCUUAGGAGAACAAGAGCUGGGA-----AUGGCCGGGAAGGCCGUUCUA   51
TBFV-NC_001672.1     AA---AGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGUCAAGAAGGCCAUCCUG   51
TBFV-NC_005062.1     AA---AGACAGCUUAGGAGAACAAGAGCUGGGG-----AUCCGGAAGGCCAUCCUG   51
TBFV-NC_003687.1     -----AGCUUUAGGAGAACAAGAGCUGGGAGUGGGUUUG-----AUGACCACUUCU   45
MBFV-NC_001437.1     UUAAACAGUUUUUUAGAACGGAAGAUAACC------AUGACUAAAAA-----CCAG   46
MBFV-NC_002640.1     -----UGUUUGAGAAGAGA-GCAGA-UCUCUGGAAAAAUG---AA--CCA-----ACGA   43
MBFV-NC_001477.1     AAC--AGUUUUUAUUAGAGA-GCAGA-UCUCUG---AUGAACAA--CCA-----ACGG   45
MBFV-NC_001475.2     GAC--AGUUUUUAUUAGAGA-GCAGA-UCUCUG---AUGAACAA--CCA-----ACGG   45
MBFV-NC_001474.2     ACA--GUUUUUAUUAGAGA-GCAGA-UCUCUG---AUGAAUAA--CCA-----ACGG   45
MBFV-NC_012532.1     -----AGUUUGGAUUUUGGAAAC-GAGAG-UUUCGG---UCAUGAAAAA-CCCA-----AAAA   47
MBFV-NC_035889.1     -----UUUUGGAUUUUGGAAAC-GAGAG-UUUCGG---UCAUGAAAA-CCCA-----A---A   42
MBFV-NC_012533.1     -----GUGUUAGGAGAAAGAAA-ACAGAUUCUUCGG---AAAUGAAAAC--CCA-----A-AA   42
ISFV-NC_017086.1     UGAAUAAGGUUUAUUGUGAAUCGAAA-UACCG--------AUGGCGAAUAAA-----CCCA   46
ISFV-NC_040610.1     AAAGAAGUCAAAACUAGAAAAUUUAA--UUUCA-------AUGGUUAGACAA----CUUG   46
ISFV-NC_024017.1     UGAU--AGAGUUGACUAGAUAACGAAUAAUUCG-------AUGGUGACUAAA----UCAA   46
MBFV-NC_009029.2     ACAC--AGUUUGUAACAAGGUUUUUGAUCGAGCA---AUGACC-AAAAAA---C---CAG   46
MBFV-NC_032088.1     -----AGUUGUGAACAAGGUUUUUGAUCGUGGAU----AUGACU-AAAAAA--C---CAG   46
MBFV-NC_012534.1     -----AGUUUGAACGUUUUUUUGGAUAGAGACAAC---UAUGUUUGAACAAAA----AACCAG   49
MBFV-NC_015843.1     -----AGUUUGAACGUUUUUUUGGAUAGAGACAAC---UAUGUCUAACAAAA----AACCAG   49
MBFV-NC_007580.2     GC---GAACAGUUUUUU--AGC-AGGGAAUAUACCC----AAUGUCUAAAAAA----CCAG   46
MBFV-NC_009028.2     UC--U-GAAUAGUUUUUUAGCA-AGGGAUUUCCC----AAUGUCUAAAAAA----CCAG   46
MBFV-NC_040776.1     UC--U-GAACAGUUUUUUAGCA-GAGGUAUCUUCAU----AAUGUCUAAAAAA----CCAG   46
MBFV-NC_000943.1     -U-G-GACAGUUUUUUUGGAGUCUUUUGAUUUCAA----AAUGUCUAAAAAA----CCAG   46
MBFV-NC_006551.1     GC-C-GGCAGUUUUCUUUGAGC--UGUUGAUUUCA----AAUGUCUAAGAAA----CCAG   46
MBFV-NC_001563.2     UG-C-GAGCUGUUUUCUUUGCUGCGAAGAUCUC-G----AAUGUCUAAGAAA----CCAG   46
MBFV-NC_009942.1     UG--C-GAGCUGUUUUCUUUGCUGCGAAGAUCUC-G----AAUGUCUAAGAAA----CCAG   46
MBFV-NC_009026.2     U----CGAGGA--AAGACGAGAAGCGAAUUCUCUG-----AAUGGCUAGAAAA----CCAG   46
MBFV-NC_034151.1     U----GAAUAG-UUUUUUGAACGAGAGGUGAAAAC-----CAUGUCUAGAAAA----CCAG   46
MBFV-NC_002031.1     AGUGCGGAUUGAGCAAAGAGUUUGAAAAGACCAGA---AAUUGA-GAUCAAUGGGUCUUUCG   50
ISFV-NC_016997.1     -----CAUUAGAAAGUUUGAAAAGAUUGAAAUC-----AAUUGA-GAAAUGGGUCUUUUCG   50
MBFV-NC_027999.1     -----UUGUUUGAGAAGACAGCGAGAAGUACUAC---U-AUGGUUGAGAAAACUAGAGUA   51
MBFV-NC_008719.1     -----AAAGGGAUAGAAGAAAG-GAGUUUUGCUUC---UCAUGG-CAACAAGAGGAGCAAG   50
MBFV-NC_012735.1     -----AAGGAGUAGAAGAAAG-GAGAUUCAUUCC---CAAUGG-CAACAAAGGGGAUGAA   50
NKV-NC_005039.1      CGGAAGAACAACUGUCCGGAACUCAAAGACG-------AUGAAAACGUA----GACA   47
NKV-NC_004119.1      C----CGGCUGGAAACCACGUCGCUCUUCGUCA-----AUGACGAAAGGCA---AAAUG   48
NKV-NC_003635.1      G----UGCGCGUCACGCCACUUUUUGUGGGCGGG---AAGUGCAGAAAUA---ACAGG   48
                     0........70........80........90.......100.......110.......1
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(secondary-structure track: `(((((((((((`)

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ISFV-NC_034204.1     GUUUUCCCGACCGGGAG-GUGGUCGG--GCCCGGCCCAGAGAGGGGCUGGGGAAGA----   100
ISFV-NC_034017.1     GUUUUCCCGGCGGGAG-GCGGUCGG--GCGGGCCCCAAAGAGGGGCUGGCGAAGA----   100
ISFV-NC_024299.2     GGAGUAAAUUAUGGAG-AAGGGGAAGCUAUCCGCCCCUCCCUUCAUUGAGGGGAGAAAA----   100
ISFV-NC_027817.1     GCCUU--CUUCCCCCG-AGGUUGGCUU-GGGAGGAGGAAGGAAAAACGGGCUUUGAG--A--   100
ISFV-NC_027819.1     ACUGGAAGAAGAAGGGAGAAUUUCCU--GGGAGGAGAGGAAUGGGCUAGAGUGA----   100
ISFV-NC_008604.2     UAAGAAGAAGAAGGGGCCGGGCUC--CUCCGGAUGGUUGUUUACCACCG--GGGCGCG--   100
ISFV-NC_021069.1     UCGGGAGUAGUGGAGAGAGGACGG--GGUACGCGCCCUCGACCAACCGUGGAGGAA--   100
ISFV-NC_012671.1     UCUGGAGUGGUGGAGAGAGGACGU--GGAACGCGUCCUCCGCAUCGUGGAGGAA--   100
ISFV-NC_001564.2     UUUGGAGGCGCGAGGAAAAGUGCCAGGAGAUUCUGG--UCAUGGAGGAAGGAA--   100
ISFV-NC_005064.1     CUUGAAGAAGAAGGCCUCCUCGGGCCCAACAUCCAAC-CGGUCUGUCAGUCGUGGAGG--   100
ISFV-NC_012932.1     CUUGAAGAAGAAGGCCUCCUCGGCCCAAAACUUGGGUCGUCAUCAUGGAACCGCGCCG--   100
TBFV-NC_001809.1     AAAGGCAAGGGGGGGCGGUCCCCCUCGACGAGUGUCAAAAGAGACCGCGA--------   100
TBFV-NC_027709.1     AAAGGUAAGGGGGGGGUGUCCCCCUCGACGAGUGUCGAAAGAGACCCCAA--------   100
TBFV-NC_003690.1     AAAGGAAAGGGGGGCGGUCCCCCUCGACGAGUGUCAAAAGAGACCCGAA--------   100
TBFV-NC_001672.1     AAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCAAAAGUGGCCCAA--------   100
TBFV-NC_005062.1     AAAGGUAAGGGGGGCGGUUCCCCCUCGACGAGUGUCGAAAGAGACCGUCA--------   100
TBFV-NC_003687.1     AAAGGAAAGGGGGGCGGUCCCCCUCUAGCGCCGAGACGUCAAUAUAGG--------   100
MBFV-NC_001437.1     GAGGGCCCGU--AAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCGUAACCCCGC----   100
MBFV-NC_002640.1     AAAAAGGUAG---UAGACCACUUCCUAUUAUGCUGAAACGCGAGAGAAACC--CGC-GUA   96
MBFV-NC_001477.1     AAAAGACGG---GUCGACCGUCUUUCAAUAUGCUGAAACGCGCGAGAAACC--CGC-GUG   99
MBFV-NC_001475.2     AAGAAGACGG---GA--AAACGCGUCUUCAAUAUGCUGAAACGCGUGAGAAAC--CGU-GUG   99
MBFV-NC_001474.2     AAAAAGACGG---GA-AAACCGCUGUCAAUAUGCUGAAACGCGUGAGAAACC--GU-GUG   99
MBFV-NC_012532.1     AAGAAUCCGGA--GUAUCC--GGAUUGUCAAUAUGCUGAAACGCGGAGUAGUGC--CCG-UG-   96
MBFV-NC_035889.1     AAAAAUCCGGA--GUAUCC--GGAUUGUCAAUAUGCUGAAACGCGGGUAGUAGC--CCG-UG-   96
MBFV-NC_012533.1     AAAAGGGCCGC--AUCG-CGGCCUAUCAAUAUGCUGAAGCGCGUUUGGAUCU---AGC-GCU   96
ISFV-NC_017086.1     AGAAGCCCCGG---AAGACCGGUUCUAUCAAUAUAGUGAAGACGUGCGUUGCCUCGCGUCU-   100
ISFV-NC_040610.1     GAAAGACCGUU--GCAAGACGGUCUGUCAAUAUGCUGAAAGC--GGAUUACCACCCG-CU-   100
ISFV-NC_024017.1     GAAGCCCCGCC--GCAAAGCGGGCGCGUCAAUAUGCUAACAC--GCAUCACCAAGCG-UG-UU   100
MBFV-NC_009029.2     GAAGUCCCCGGU--AGUAAUCGGCCGUCAUAAUGCUGUAAGCGCGGUGUCGUC--CCG-CU-   100
MBFV-NC_032088.1     GAAGUUCCGGC--AAUCUUCGACUGUCAAUAUGCUGUAGCGCGGCGGUCGUC--CCG-CU-   100
MBFV-NC_012534.1     GAAACCCCGGU--CAACCCGGGGUCUGUCAAUAUGCUGUAGCGCGGUCGUC--CC-----   100
MBFV-NC_015843.1     GAAGACCCGGC---UCAGCCCGGGGUUGUCAAUAUGCUGUAAGCGCGGGAACGUC--CC-----   100
MBFV-NC_007580.2     AAAACCCGGG---CUGACAGGGGUUGUCAAUAUGCUGAAACGCGGGAUCGUC--CCG-UG-   100
MBFV-NC_009028.2     GAAAGUCGCU--GCAAAGCGGACGUGUCAAUAUGCUGAAACGCGCGGUAGAGC--CGU-GU-   100
MBFV-NC_040776.1     GAGGACCGCC---GGAAGACGGGGUGUCAAUAUGCUGAAACGCGCGGUAGGU--UU----   100
MBFV-NC_000943.1     GAGGGCCCGGA--AAAGACCGGGCGUCAAUAUGCUGAAACGCGGCAGUCC--CCG-UG-   100
MBFV-NC_006551.1     GAGGGCCCGGA--AAAACGGGCUCAUCAAUAUGCUGAAACGCGGCCAUCC--CC-----   100
MBFV-NC_001563.2     GAGGGCCCGGA--AAAGACCGGGCGUCAAUAUGCUGAAACGCGGUAUGCC--CG-CU-   100
MBFV-NC_009942.1     GAGGGCCCGC--AAGAGCCGGGCGUCAAUAUGCUGAAACGCGGUGAAUGCC--CG-CU-   100
MBFV-NC_009026.2     GAAGACCCGGA--AAGAACGGGGCGUCAAUAUGCUGUAGCGUACGGCGAC--CAA-CG-   100
MBFV-NC_034151.1     GAGACCCCGGA--AAGAACGGGGCGUCAAUAUGCUGAAGCGCACAUCCCG--CGC-UA-   100
MBFV-NC_002031.1     AAGCUCAGGGA--AAGACCCUGCGGUGCUAUAUGGUGCGCUGGAAAGAGC--GACUGAUUU-   100
ISFV-NC_016997.1     AGCUUAGGAGGCGCCGUUAACGGCCGUCUAUAUUAAAAUAAAGCCUGUAGGCUGU---   100
MBFV-NC_027999.1     GGAUCUCGGGU--AAAACCGAGGUGUCAAUAUGUUGAUUGAGCCGGCCAAGGUA--------   100
MBFV-NC_008719.1     CAAGUCUCGGGCUCACGACCCGGGGUCAAUAUGGUGAGCGGCCAAGAUAA---------   100
MBFV-NC_012735.1     UAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUGAGCUGCUAGAGUGA---------   100
NKV-NC_005039.1      AAACACCCG------AAUUUCG---GUUGUCAAUAUGCUAAAAAUUCCAAAAAAACAAA   99
NKV-NC_004119.1      AAAAACGCGAAUAAAAAAGAAAGCAAAAAAUCUCUUAAGACCAAUUCCUUGGGUCGAA   100
NKV-NC_003635.1      AAAAAGCCGUGAAAGGGGUUUCAAAGAAAGAAAGAUGUCGUUGAAGGAAUG--------   100
                     20.......130.......140.......150.......160.......170.......1
```

(secondary-structure track: `)))))))))))`)

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ISFV-NC_034204.1     ----NNNNNNNACG---ACCACCUCUCGUGAAAUGUUAAUAAAACUGUGCGG-AAAAUUCCA   152
ISFV-NC_034017.1     ----NNNNNNNGCG---ACCACCUCUCGUGAAAUGUUAAUAAAACUGUGCGG-AAAAUUCCA   152
ISFV-NC_024299.2     ----NNNNNNNAGU---A-CACCCCCCGGUUGUGUUAUAUAGCCCAUGUU-GAGUCUGUA   151
ISFV-NC_027817.1     ----NNNNNNNAGU---AGCCGUCACCCCGGUUGUGAAAACCGAUUGGUAGCUUACACAGUA   152
ISFV-NC_027819.1     ----NNNNNNNUUA-GCCACACCCCCCGAAGUGAUUUAU--AUGAUGAC-AGAA-AUCA   152
ISFV-NC_008604.2     ----NNNNNNNGUG-ACGCCACCCCCCGAAAUGUGUAUGU-AACAGAC-AUGC-CUGC   152
ISFV-NC_021069.1     ----NNNNNNNUG-ACGCCACCCCCCGAAAUGUGUAUUGU-GACAG-GC-UUGC-UCGC   151
ISFV-NC_012671.1     ----NNNNNNNGUG-ACGCCACCCCCCGAAAUGUGUAUUGU-GACAG-GC-UUGC-UCGC   151
ISFV-NC_001564.2     ----NNNNNNN-AGCCGAACCCUCUCCGUGAAAUGAAUGGAAAGGGGAAAU-AAGAACCCA   153
ISFV-NC_005064.1     ----NNNNNNNCGU----ACCCCCCUCG-UAAUGUGAAAAAAAUAUGGGGGAA-AAGAACCCA   152
ISFV-NC_012932.1     ----NNNNNNNUUGGCGAACCCCCCCG-AAAUUGUAAAAAUAUUAUGGGGA---AAGUACCCA   154
TBFV-NC_001809.1     ----NNNNNNNG-UUUUUCCUCCUCCCUAUCAAAUCAUAU---CCCUCGACAGAGGGGGG   152
TBFV-NC_027709.1     ----NNNNNNNGAUUUUUCCUCCUCCUAUACCAAAU-UCC---CCCUCGACAGAGGGGGG   152
TBFV-NC_003690.1     ----NNNNNNNGAUUUUUCCUCCUCCUAUACAAAU-UCCC---CCCUCGUCAGAGGGGGG   152
TBFV-NC_001672.1     ----NNNNNNNUAUACAAAAUCUUCCUCGGU-AGAGGG--GGGCGGUUCUUGUUCUUC   152
TBFV-NC_005062.1     ----NNNNNNNUAUCAAAUCUUCCUCUCUAUCAC-UGAGGG---GGGCGGUUCUUGUUGUUC   152
TBFV-NC_003687.1     ----NNNNNNN-AUUUUCCUCCCUCUAUAACUAAAUGG----UCCAGGAAACUGGGGGU   152
MBFV-NC_001437.1     ----NNNNNNNCAUAUUGACACCUUGGGAUAGAGCUGGGAGAUCUUCUG------CUCUAUC   151
MBFV-NC_002640.1     UCAANNNNNNNCCCCCCAUCA--CAAAAACAGCAUAUUGACG--CUGGGA--AAGACCA-G   150
MBFV-NC_001477.1     U--NNNNNNNCCGCACAC---AACAAACAGCAUAUUGACG-CUGGGA---GAGACCA-G   150
MBFV-NC_001475.2     U--NNNNNNNCCCCCGCAAA--UAAAAACAGCAUAUUGACG--CUGGGA---GAGACCA-G   150
MBFV-NC_001474.2     U--NNNNNNNCCCCAC---AA--AAAAAACAGCAUAUUGACG--CUGGGA---AAGACCA-G   150
MBFV-NC_012532.1     ----NNNNNNNCCCGGAAAACGCAAAACAGCAUAUUGAC--G---UGGGA--AAGACCGA-G   150
MBFV-NC_035889.1     ----NNNNNNN-CCCGGAAAACGCAAAACAGCAUAUUGAC--G--CUGGGA--AAGACCG-G   150
MBFV-NC_012533.1     GGCNNNNNNNAUAACCAACA-UAAAAACAGCAUAUUGACACCUUGGGA---AAGACCG-G   150
ISFV-NC_017086.1     ----NNNNNNNCAAAAAUG-A-AGGGCACUAAUAGUGACACUUUGGGG--AAAGACCA-G   150
ISFV-NC_040610.1     ----NNNNNNNCUCAUUAUU--AUAUAACAGCAUAUUGACACCAGGG--AUAGACCG-G   151
ISFV-NC_024017.1     ----NNNNNNNC-GCUGUGA--GCAAAAUGACAUAUUGACA--CUGAGG--GGUGAGACCA-A   150
MBFV-NC_009029.2     ----NNNNNNN-AGCGCAAA---ACAAAACAGCAUAUUGACA--CCUGGGA--UGACUA-UG   150
MBFV-NC_032088.1     ----NNNNNNN-UACCGCAA---ACAAAACAGCAUAUUGACA--CCUGGGA---UGACUA-UG   150
MBFV-NC_012534.1     ----NNNNNNNAACACCAAA-UAAAAACAGCAUAUUGACA--CCUGGGA--UAGACUA-UG   150
MBFV-NC_015843.1     ----NNNNNNNACACCACA-A-GAAACAGCAUAUUGACA--CCUGGGA---UAGACUA-UG   150
MBFV-NC_007580.2     ----NNNNNNNCCGUAAACGC-AAACAACAGCAUAUUGACG--CCU-GGA---AAGACAG-G   150
MBFV-NC_009028.2     ----NNNNNNNCAAGCGCGAA--ACAACAGCAUAUUGACG--CCUGGGA---AAGACUA-A   150
MBFV-NC_040776.1     ----NNNNNNNCAAGCAA-UGA--ACAACAGCAUAUUGACA--CCUGGGA---UAGACUA-UG   150
MBFV-NC_000943.1     ----NNNNNNNAAAGCAUCAA---CAAAACAGCAUAUUGACA--CCUGGGAA--UAGACUA-UG   152
MBFV-NC_006551.1     ----NNNNNNNAAACCAUCAA--CAAAACAGCAUAUUGACA--CCUGGGA---UAGACUA-A   150
MBFV-NC_001563.2     ----NNNNNNNAAACCAUCAA--CAAAACAGCAUAUUGACA--CCUGGGA---UAGACUA-UG   150
MBFV-NC_009942.1     ----NNNNNNNAAACCAUCAA--CAAAACAGCAUAUUGACA--CCUGGGA---UAGACUA-UG   150
MBFV-NC_009026.2     ----NNNNNNNCAAACCA-AGAAAACAGCAUAUUGACGACUGGGA---UAGACCGGA   152
MBFV-NC_034151.1     ----NNNNNNNAAACCUCUG-CUAUUU-CAGCAUAUUGACG--CCUGGGA---UAGACUCGGA   151
MBFV-NC_002031.1     ----NNNNNNNC---AAAAAUUGUAACGGCACAUAUUGAC--UGUA---AGGA---AAGACCG-G   150
ISFV-NC_016997.1     ----NNNNNNNC--AAAUGUUAUAGCU-UCAUAUUGACA--CCUAGGA---AAAGACC-C   150
MBFV-NC_027999.1     ----NNNNNNNCCAUUGGAAAUUUAGCU-UCAUAUUGACA--CCAGGGA---AAAGACC-C   150
MBFV-NC_008719.1     ----NNNNNNNCC-GAGCACACAUAUGGUGGACACAUUGACG--CCUGGGA---AAGACC-G   151
MBFV-NC_012735.1     ----NNNNNNNC-GAGCACACAUAUGGCAACAUAUUGACG--CCAGGGA---AAGACC-G   150
NKV-NC_005039.1      A----NNNNNNNNUUGCCUCAUCGUGAAUGGGACGGGCAGAUU------UGGAU-AUAUUACUCCAU   150
NKV-NC_004119.1      ----NNNNNNNUGUGCCUAAUUCC-CGUUACAGGGGGU----GGGGUUGUAUUAUAGCCAUGG   150
NKV-NC_003635.1      ----NNNNNNNUUUCCCAAUUGU-CCGUGUUAUACAGU--AAGGA---AAGACCU-C   150
                     80......190.....
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ISFV-NC_034204.1     AAGACUGUCC-CCACC-  167
ISFV-NC_034017.1     AAGAAUGUCC-CCACC-  167
ISFV-NC_024299.2     CAACAGAGUU-CCACCG  167
ISFV-NC_027817.1     AAGUCGAGAG-UCUCC-  167
ISFV-NC_027819.1     GAUGGAAGAU-CUCCA-  167
ISFV-NC_008604.2     AGCAGCAGGU-CUCCA-  167
ISFV-NC_021069.1     AGCAGCAAAU-CUCCAC  167
ISFV-NC_012671.1     AGCAGCAAAU-CUCCAC  167
ISFV-NC_001564.2     UCUUA--ACC-UCCUAC  167
ISFV-NC_005064.1     UCUAAAUCUC-CCACCG  167
ISFV-NC_012932.1     UCUAA---AGC-CUCCCA  167
TBFV-NC_001809.1     GCGGUUCUUU-GUUUC-  167
TBFV-NC_027709.1     GCGGUUUUU--GUUCUC-  167
TBFV-NC_003690.1     GCGGUUUUU--GUUCUC-  167
TBFV-NC_001672.1     CCUGAGCCA--CCAUCA-  167
TBFV-NC_005062.1     CCUGAGCCA--CCAUCA-  167
TBFV-NC_003687.1     GCGGUUUCU--GUUCUC-  167
MBFV-NC_001437.1     UCAACAUCA-GCUACUA  167
MBFV-NC_002640.1     AGAUCCUGCUGCUCUUC  167
MBFV-NC_001477.1     AGAUCUGUCUGUCUCUA  167
MBFV-NC_001475.2     AGAUCCGUCUGUCUCUA  167
MBFV-NC_001474.2     AGAUCUGUCUGUCUCUA  167
MBFV-NC_012532.1     GACUCCAUGGAGUUUCC  167
MBFV-NC_035889.1     GACUCCAUGAGUUUCCC  167
MBFV-NC_012533.1     AGACUCUGUUGGCUUUC  167
ISFV-NC_017086.1     AGGUAUCGCUGGAAUUC  167
ISFV-NC_040610.1     AGAAAUACACAAUUUUC-  167
ISFV-NC_024017.1     AGAUAUG-AUGUGUUUG-  167
MBFV-NC_009029.2     GGGAUUGUCUGUCCUGG  167
MBFV-NC_032088.1     GAGACCGUCUGUCCUGG  167
MBFV-NC_012534.1     GAGACCCUCUGUCCUAA  167
MBFV-NC_015843.1     GAGACCCUCUGUCCUAA  167
MBFV-NC_007580.2     GAGACCCUCUGUCCUUC  167
MBFV-NC_009028.2     GAGACCCUCUGUCCUUC  167
MBFV-NC_040776.1     GAGACCCUCUGUCCUUC  167
MBFV-NC_000943.1     GAGAUCUUCUGUCUCUG  167
MBFV-NC_006551.1     GAGAUCUUCUGUCCUUC-  167
MBFV-NC_001563.2     GAGAUCUUCUGUCUUG-  167
MBFV-NC_009942.1     GAGAUCUUCUGUCUGC-  167
MBFV-NC_009026.2     GAGAUCCUCUGUCCGGA  167
MBFV-NC_034151.1     AGACCCUCUG-CUUU-  167
MBFV-NC_002031.1     GUG-GUUCUC-UGCCUUU  167
ISFV-NC_016997.1     AGAUACUUUGAGUUUU-  167
MBFV-NC_027999.1     GAGAAACUUCUUGCUUA  167
MBFV-NC_008719.1     GAGACACUAUAGGCGCC-  167
MBFV-NC_012735.1     AGACA-CUCCUUGAUUC-  167
NKV-NC_005039.1      GUGUUUGUCGUGAUUU-  167
NKV-NC_004119.1      CCAGAAAAGACCUCAGAU  167
NKV-NC_003635.1      UCGCACAGAUCAUGCU-  167
                     80......190.....
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