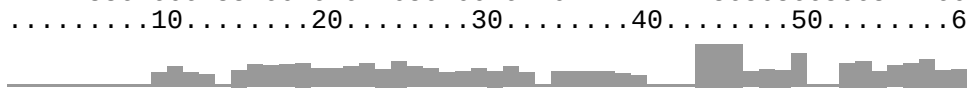
```
                   ............................................................
NKV-NC_003635.1    ---------GUGCGCGUCACGCCACUUUUUGUG-GCGGGA---AUGUGCG--AAAAUAAC 45
NKV-NC_004119.1    ---------CCGGCUGGAAACCACGUCGCUCUU-CGUCAA---AUGACGA--AAGGCAAA 45
ISFV-NC_027817.1   ---------CAAACUCUGGUGUCCAACGUUUGU-AUUAAA---AUGAGUG--GA-UUAGG 44
ISFV-NC_034204.1   --------GCGACAGAAGGAACCCAAAAAGGGA-ACCAG----AUGGAGCAAAACGGA-A 46
ISFV-NC_034017.1   --------UAAAAAUUUCUUUGGCGACGUACGG-AAAGU----AUGGAGCAAUCGAGG-A 46
ISFV-NC_001564.2   ---------GAGAGACCGUGAGGUUGAACCCGG-CAAGGA---AUGAAGAGGAAGGAU-U 46
ISFV-NC_005064.1   ---------AGUUGUUCGGAAAAGUGACCUUGG-UUCGUU---AUGAAUAGAGGAAGC-U 46
ISFV-NC_012932.1   ---------AGUUGUUUGGAAAAGAGAUAUCUG-GUUCGU---AUGAAUAGGAUGCUC-U 46
ISFV-NC_008604.2   ---------UUGUUUAUAAUAGUCGUUUUUCUC-GCAGAA---AUGGGAAAGGACGACGG 47
ISFV-NC_021069.1   ---------CCUUAUUACCUCUGGUACUAAAGU-GGCACU---AUGAAGAAGAAGGA--G 45
ISFV-NC_012671.1   ---------CUUUAUUACCUCUGGUACUAACGU-GGCGCU---AUGAAGAAGAAGGA--G 45
ISFV-NC_024299.2   ---------UUGAAAAACUUUUGCGACGUUGUU-GAGAAG---AUGGAGCGAAAGAAAGG 47
ISFV-NC_027819.1   ---------UCACGGAAUAGCUUGAGAAGAAGC-UUACUU---AUGUCAAGUAAGGAAGC 47
TBFV-NC_003687.1   --------AGCUUAGGAGAACAAGAGCUGGGAG--UGGUU---AUG------AUGACCAC 41
TBFV-NC_005062.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGGG---AUGGCCGGGAAGGCCAU 47
TBFV-NC_001672.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGGG---AUGGUCAAGAAGGCCAU 47
TBFV-NC_001809.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGGG---AUGGGCAGGAAGACCAU 47
TBFV-NC_027709.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGGG---AUGGGCAGGAAGUCCAU 47
TBFV-NC_003690.1   --------AAAGACAGCUUA-GGAGAACAAGAG-CUGGGA---AUGGCCGGGAAGGCCGU 47
MBFV-NC_001437.1   -------UUAAAC-AGUUUUUUAGAACGGAAGA--UAACC---AUGACUA---AAAAACC 44
ISFV-NC_040610.1   ----AAAGAAGUC---AAAACUAGAAAUUUA-AUU-U-CA---AUGGUUA---GACAACU 44
ISFV-NC_024017.1   ----UGAU-AGAG---UUGACUAGAUAACGA-AUUAU-CG---AUGGUGA---CUAAAUC 44
MBFV-NC_034151.1   --------UGAAU-AGUUUUUUG-AACAGAGGUGAAAACC---AUGUCUA---AAAAACC 44
MBFV-NC_009028.2   ------UCUGAAU-AGUUUUUU--AGCAAGGGA-UUUCCC---AUGUCUA---AAAAACC 44
MBFV-NC_040776.1   ------UCUGAAU-AGUUUUUU--AGCAGAGGU-ACUACC---AUGUCUA---AAAAACC 44
MBFV-NC_001563.2   ------UGCGAGC-UGUUUCUUGGCAC-GAAGA-UCU-CG---AUGUCUA---AGAAACC 44
MBFV-NC_009942.1   ------UGCGAGC-UGUUUCUUAGCAC-GAAGA-UCU-CG---AUGUCUA---AGAAACC 44
MBFV-NC_006551.1   ------GCCG-GC-AGUUUCUUUGAGC-GUUGAUUUU-CA---AUGUCUA---AGAAACC 44
MBFV-NC_000943.1   ------U--GAAC-AGUUUUUUGGAGCUUUUGAUUUC-AA---AUGUCUA---AAAAACC 44
MBFV-NC_009029.2   ACACAGUGUGAACAAGUUUUUUG-AACAG---------CA---AUGACCA---AAAAACC 44
MBFV-NC_032088.1   ----AGUGUGAACAAGUUUUUUG-AUCAGUGGA-----UA---AUGACUA---AAAAACC 44
MBFV-NC_007580.2   -------GCGAAC-AGUUUUUU--AGCAGGGAAUUACCCA---AUGUCUA---AAAAACC 44
MBFV-NC_012534.1   ----AGUUUGAAC--GUUUUUUGGAGAGAGA----UUAUU---AUGUUUAACAAAAAACC 47
MBFV-NC_015843.2   ----AGUUUGAAC--GUUUUUUGGAUAGAGA----CAACU---AUGUCUAACAAAAAACC 47
MBFV-NC_009026.2   -------UCGAGG-AAAGACGAGAAGCGAAUU--CUCUGA---AUGGCUA---GAAAACC 44
MBFV-NC_012532.1   -------A---UU-UGGAUUUGGAAACGAGAGUUUCUGGUC--AUGAAAA--ACCCCAAA 45
MBFV-NC_035889.1   -------U---UU-UGGAUUUGGAAACGAGAGUUUCUGGUC--AUGAAAA--A-CCCAAA 44
MBFV-NC_012533.1   -------G---UG-UUGAGAAAGAAA--ACAGAUUCUUCGGAAAUGAAAA----CCCAAA 43
MBFV-NC_001474.2   -------ACAGUU-UUUUAAUUAGAGAGCAGAUCUCUG-----AUGAAUA--A-CCAACG 44
MBFV-NC_001477.1   -------AACAGU-UUUUUAUUAGAGAGCAGAUCUCUG-----AUGAACA--A-CCAACG 44
MBFV-NC_001475.2   -------GACAGU-UUUUUAUUAGAGAGCAGAUCUCUG-----AUGAACA--A-CCAACG 44
MBFV-NC_002640.1   -------UGUUUG-AAUAGAGAGCAGAUCUCUGGAAAA-----AUG---A--A-CCAACG 41
ISFV-NC_017086.1   -------UGAAUA-GGUUUAUUGUGAAUCGAAAUACCG-----AUGGCGA--A-UAAACC 44
MBFV-NC_008719.1   ------------AAAGGGUAGAGAAAAGGAGUUUUGCUUCU-CAUGGCAACAAGAGGAGC 47
MBFV-NC_012735.1   ------------AAGGAGUAGAAGAAAGGAGAUUCAUUCCC-AAUGGCAACAAAGGGGAU 47
ISFV-NC_016997.1   ------------CAUUAGAAUAGGUUUGAAAGAGUUGAAUC-AAUGAGAUCAAUGGUCUU 47
ISFV-NC_027999.1   ------------UUGUUUGAGAGAACAGCGAGAAGUACUAC-UAUGGUUGAGAAAACUAG 47
NKV-NC_005039.1    -----CGGAAGAACAACUGUCCGGAAUCAAAGACG--------AUGAAAAAUCGUA---- 43
MBFV-NC_002031.1   -----UUGAGCGAUUAGCAGAGAACUGACCAGAAC--------AUGUCUGGUCGUAAAGC 47
                   .........10........20........30........40........50........6
```



```
                   ...............................(((((((((.....................
NKV-NC_003635.1    AGGAAAAAGAACCCUGGAAAGGGUUCAAAGAAGAAAAAGAAUGCUGGAAGG-AAUG---- 100
NKV-NC_004119.1    AUGAAAAACGCGAAUAAAAAGAAAGCAAAAAACUCUAAGACCAAUCCUGGU-CGAA---- 100
ISFV-NC_027817.1   AGGCCUUCUUCCCCUCCGAGGGAAGAAAAAGAAAGCCCCUGUGAUACAAUCACAGG---- 100
ISFV-NC_034204.1   AGUUUU--CCCGACCGGGAGGUGGUCGGGCCCGGCCCAGAGAGGGGCUGG-GGAAGA--- 100
ISFV-NC_034017.1   AGUUUU--CCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGAGGGGCUGG-CGAAGA--- 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_008604.2   UAAGAA--GAAGAAGGGGCCGGGCUC-CUCCGGAUGGUUGUUACCACCGGGG-CGCG--- 100
ISFV-NC_021069.1   UCGGGA--GUAGUGGAGAGAGGACGCGGUACGCGCCCUCCGACCAACCGUGGAGGAA--- 100
ISFV-NC_012671.1   UCUGGA--GUGGUGGAGAGAGGACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAG--- 100
ISFV-NC_024299.2   AGUAAA--UUAUGGGAGAAGGGAAGC--UAUCCCGCCCCCUCCCAUUAAGGAGAAAA--- 100
ISFV-NC_027819.1   ACUGAA--GAAGAAGGGAAUUUCCCUGGGAGGAGGAGGAAAACGGGGCUUUGAG--A--- 100
TBFV-NC_003687.1   UUCUAAAGGAAAGGGGGGCGGUCCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAAGU- 100
TBFV-NC_005062.1   UCUGAAAGGAAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA------- 100
TBFV-NC_001672.1   CCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA------- 100
TBFV-NC_001809.1   CCUGAAAGGCAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCGA------- 100
TBFV-NC_027709.1   CCUGAAAGGUAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA------- 100
TBFV-NC_003690.1   UCUGAAAGGAAAGGGGGGCGGUCCCCCUCGACGAGCGUCGAAAGUGGCCCAA------- 100
MBFV-NC_001437.1   AGGAGG--GCCCGGUAAAAACCGGGCUA-UCAAUAUGCUGAAACGCGGCCUACCCCGCG- 100
ISFV-NC_040610.1   UGGAAA--GACCGUUGCAAGACGGUCUG-UCAAUAUGCGAAAGCGGAUUACCACCCGCU- 100
ISFV-NC_024017.1   AAGAAA--GCCCGCCGCAAAGCGGGCUG-UUAAUAUGCUAACACGCAUCACCAAGCGUG- 100
MBFV-NC_034151.1   AGGAGG--ACCCGGAAAGAACCGGGUCG-UUAAUAUGCUGAAGCGCACAUCCCGCGCUA- 100
MBFV-NC_009028.2   AGGAAA--GUCCGCUGCAAAGCGGACUG-UCAAUAUGCUAAAACGCCUGGCUAGCGUGU- 100
MBFV-NC_040776.1   AGGAGG--ACCCGCCGGAAGGCGGGUUG-UCAAUAUGCUGAAACGCCCAGCCAGCGUUU- 100
MBFV-NC_001563.2   AGGAGG--GCCCGGUAAAAACCGGGCUG-UCAAUAUGCUAAAACGCGGUAUGCCCCGCG- 100
MBFV-NC_009942.1   AGGAGG--GCCCGGCAAGAGCCGGGCUG-UCAAUAUGCUAAAACGCGGAAUGCCCCGCG- 100
MBFV-NC_006551.1   AGGAGG--GCCCGGAAGAAACCGGGCCA-UCAAUAUGCUGAAACGCGGCAUACCCCGCG- 100
MBFV-NC_000943.1   AGGAGG--ACCCGGGAAGCCCCGGGUCG-UCAAUAUGCUAAAACGCGGCAUACCCCGCG- 100
MBFV-NC_009029.2   AGGAAG--GCCCGGUAGAAAUCGGGCCG-UCAAUAUGCUAAAGCGCGGUGCGUCCCCGCG- 100
MBFV-NC_032088.1   AGGAAA--GUCCGGCAAUCGUCGGACUG-UCAAUAUGCUAAAGCGCGGAGCAUCCCGCG- 100
MBFV-NC_007580.2   AGGAAA--ACCCGGUAGAAACCGGGUUG-UCAAUAUGCUAAAACGCGGCGUAUCCCGCG- 100
MBFV-NC_012534.1   AGGAAA--ACCCGGGGCAACCCGGGUUG-UCAAUAUGCUAAAGCGCGGUGCAUCCC---- 100
MBFV-NC_015843.2   AGGAAG--ACCCGGCUCAGGCCGGGUUG-UCAAUAUGCUAAAGCGCGGAACGUCCC---- 100
MBFV-NC_009026.2   AGGAAG--ACCCGGAGGAAACCGGGUCG-UCAAUAUGCUAAAGCGUACGGCGGCCAACG- 100
MBFV-NC_012532.1   GAAGAA--AUCCGGAGGA-UCCGGAUUG-UCAAUAUGCUAAAACGC---GGAG-UAGCCC 97
MBFV-NC_035889.1   AAAGAA--AUCCGGAGGAUUCCGGAUUG-UCAAUAUGCUAAAACGC---GGAG-UAGCCC 97
MBFV-NC_012533.1   AAAGG-------GCCGCAUCGCGGCCUA-UCAAUAUGCUGAAGCGCUUUGGAUCUAGCGC 95
MBFV-NC_001474.2   GAAAAA--GGCGAAAAAC---ACGCCUU-UCAAUAUGCUGAAACGCGAGAGAAACCGCGU 98
MBFV-NC_001477.1   GAAAAA--GACGGGUCGA---CCGUCUU-UCAAUAUGCUGAAACGCGCGAGAAACCGCGU 98
MBFV-NC_001475.2   GAAGAA--GACGGGAAAA---CCGUCUA-UCAAUAUGCUGAAACGCGUGAGAAACCGUGU 98
MBFV-NC_002640.1   AAAAAA--GGUGGUUAGA---CCACCUU-UCAAUAUGCUGAAACGCGAGAGAAACCGCGU 95
ISFV-NC_017086.1   CAAGAA--GCCCGGAAGA---CGGGCUA-UCGAUAUAGUGAGACGUGCGUUGCCUCGCGU 98
MBFV-NC_008719.1   AAGCAAGUCUCGGGUCACGACCCGGGGC-GUCAAUAUGGUGACGGC------CAAGGCCA 100
MBFV-NC_012735.1   GAAUAAGUCUCGGGCUCGAUCCCGAGGC-GUCAAUAUGGUAGCGGC------UAGAGUGA 100
ISFV-NC_016997.1   UCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACG-------UAAGCUGC 100
ISFV-NC_027999.1   AGUAGGAUCUCGGGUUAAAACCCGAGGU-GUCAAUAUGAUUAGGCG------UACUGGAA 100
NKV-NC_005039.1    GACAAAACACCCGAAUU---UCGGGU----GUCAAUAUAAAAAAGCAAAAU-CCAAAAAA 95
MBFV-NC_002031.1   UCAGGGAAAAACC--CU---GGGCGU----CAAUAUGGUACGACGAGGAGU-UC--GCUC 95
                   0........70........80........90........100.......110.......1
```

```
                   ...............................))))))))))...................
NKV-NC_003635.1    -----NNNNNNNUUUUCCAAAUUCCCGU---UACCGGGGGUGGGGUGA---U-UAGCCAU 148
NKV-NC_004119.1    -----NNNNNNNUUGCCUCAUCUUGAAU---GGGGGGCGGCGUGGAUA---UAUACUCCA 149
ISFV-NC_027817.1   -----NNNNNNN-AGUAGCGUCACCCCC---CGGUUGUGAAAACGAUU---GCGACUAGA 148
ISFV-NC_034204.1   -----NNNNNNN----ACGACCACCCUC---UGUAAUGU-U-AAUAAAACUGUGCGGAAA 146
ISFV-NC_034017.1   -----NNNNNNN----GCGACCACCCUC---UGUAAUGU-U-UAAUAAGUUGUGCGGAAA 146
ISFV-NC_001564.2   -----NNNNNNN---AGCGAACCUCCUC---CGUUAAUG-UGAAAAAGUAUGGGG-AAAG 147
ISFV-NC_005064.1   -----NNNNNNN-----CGUACCCCCUC---CG-UAAUG-UGAAAAAGUAUGGGGAAAAG 145
ISFV-NC_012932.1   -----NNNNNNN--UUGCGAACCCCCUC---CG-AAAUG-UGAAAAAUUAUGGGGAAAAG 148
ISFV-NC_008604.2   -----NNNNNNN--CUGACGCCACCCUC---CGAAAUGU-GUAUAGUAACAGAGCAUGCC 149
ISFV-NC_021069.1   -----NNNNNNN--GUGACGCCACCCCC---CGAAAUGU-GUAUUGUGACAG-GCUUGCU 148
ISFV-NC_012671.1   -----NNNNNNN--GUGACGCCACCCCC---CGAAAUGU-GUAUUGUGACAG-GCUUGCU 148
ISFV-NC_024299.2   -----NNNNNNN-----ACGACACCCCC---CGAAAUGU-GUAUAUAGCCCAUGUUGAGU 146
ISFV-NC_027819.1   -----NNNNNNN--UUAGCCACACCCCC---CGAAACGU-GAUUUAUAUGAUGACAAGAA 149
TBFV-NC_003687.1   -----NNNNNNN-AUUUUUCCGCCUCCUAUACUAAAUUU-CCCCCAGGAAACUGGGGGGG 153
TBFV-NC_005062.1   -----NNNNNNN--------------UAUACCAAA-UU-CCCCCUCAUCUGAGGGGGGG 138
TBFV-NC_001672.1   -----NNNNNNN---------------UAUACAAAA-UU-CCCCCUCGGUAGAGGGGGGG 138
TBFV-NC_001809.1   -----NNNNNNNG-UUUUUCCUCCUCCUAUACCAAAUUU-CCCCCUCGACAGAGGGGGGG 153
TBFV-NC_027709.1   -----NNNNNNNGAUUUUUCCUCCUCCUAUACCAAA-UU-CCCCCUCGACAGAGGGGGGG 153
TBFV-NC_003690.1   -----NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UU-CCCCCUCGUCAGAGGGGGGG 153
MBFV-NC_001437.1   -----NNNNNNN-CAUAUUGACAC-CUGGGAAUAGACUGGGAGAUCUUC--UGCUCUAUC 151
ISFV-NC_040610.1   -----NNNNNNNCUCAUUAUUUAU-AUAACAGCAUAUUGACACCAGGG-AUAGACCGGAG 153
ISFV-NC_024017.1   -----NNNNNNNCUGCUGAGAG-C-AAAAUAGCAUAUUGACACUUGGGAAGAGACCAGAG 153
MBFV-NC_034151.1   -----NNNNNNNGAA-CGCAAAUA-AAAACAGCAUAUUGACACCUGGGA--UAGACUGGG 151
MBFV-NC_009028.2   -----NNNNNNNCAAGCGCGAA-A-CAAACAGCAUAUUGACGCCUGGGA--AAGACCGGG 151
MBFV-NC_040776.1   -----NNNNNNN-AAGCGCAAAGA-CAAACAGCAUAUUGACACCUGGGA--AAGACCGGG 151
MBFV-NC_001563.2   -----NNNNNNN-AAACACCAAAA-GAAACAGCAUAUUGACACCUGGGA--UAGACUAGG 151
MBFV-NC_009942.1   -----NNNNNNN-AACACCACAAC-AAAACAGCAUAUUGACACCUGGGA--UAGACUAGG 151
MBFV-NC_006551.1   -----NNNNNNN-UAAGCAUCAAA-AAAACAGCAUAUUGACACCUGGGAAUUAGACUAGG 153
MBFV-NC_000943.1   -----NNNNNNN-AAAAGCAUCAA-ACAACAGCAUAUUGACACCUGGGAA-AAGACUAGG 152
MBFV-NC_009029.2   -----NNNNNNN-AGCGCAAACAC-AAAACAGCAUAUUGACACCUGGGA--AAGACUAGG 151
MBFV-NC_032088.1   -----NNNNNNN-UACCGCAAAAC-AAAACAGCAUAUUGACACCUGGGA--UAGACUAGG 151
MBFV-NC_007580.2   -----NNNNNNNCCGUUAACGCAA-ACAACAGCAUAUUGACACCUGGAA---AGAC-AGG 150
MBFV-NC_012534.1   -----NNNNNNN-AACACCAAAAC-AAAACAGCAUAUUGACACCUGGGA--GAGACUAGG 151
MBFV-NC_015843.2   -----NNNNNNN-GAACACCACAA-GAAACAGCAUAUUGACACCUGGGA--UAGACUAGG 151
MBFV-NC_009026.2   -----NNNNNNN-ACGCACCAAAA-GAAACAGCAUAUUGACACCUGGGA--UAGACCGGA 151
MBFV-NC_012532.1   --GUGNNNNNNNCCCCGGAAAACGCAAAACAGCAUAUUGACG-UGGG-A--AAGACC-AG 150
MBFV-NC_035889.1   --GUGNNNNNNN-CCCGGAAAACGCAAAACAGCAUAUUGACGCUGGG-A--AAGACC-AG 150
MBFV-NC_012533.1   UGGCCNNNNNNNAUAACCAACAUA-AAACAGCAUAUUGACACCUGGG-A--AAGACC-GG 150
MBFV-NC_001474.2   GU---NNNNNNNCCCCCGAAACAA-AAAACAGCAUAUUGACGC-UGGGA--AAGACC-AG 150
MBFV-NC_001477.1   GU---NNNNNNNCCCGCACAACAA-CAAACAGCAUAUUGACGC-UGGGA--GAGACC-AG 150
MBFV-NC_001475.2   GU---NNNNNNNCCCCCGCAAAUA-AAAACAGCAUAUUGACGC-UGGGA--GAGACC-AG 150
MBFV-NC_002640.1   AUCAANNNNNNNCCCCCCAACACA-AAAACAGCAUAUUGACGC-UGGGA--AAGACC-AG 150
ISFV-NC_017086.1   CU---NNNNNNNCCCA-AAAAUGA-AGGGCACUAUAUCGACACUUGGGA--AAGACC-AG 150
MBFV-NC_008719.1   -----NNNNNNNCCGAGCAC-UAUAGCGGACCAUAUUGACGCCUGGG-A--AAGAC-CGG 150
MBFV-NC_012735.1   -----NNNNNNNCCGAGCACACAUAGCGGACCAUAUUGACGCCAGGG-A--AAGAC-CGG 151
ISFV-NC_016997.1   -----NNNNNNN-CAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGG-A--AAGAC-UGG 150
ISFV-NC_027999.1   -----NNNNNNNCC-AUUGGAAAUUUAGCUUCAUAUUGACACCAGGGAA--AAGAC-CGG 151
NKV-NC_005039.1    CAAAANNNNNNN-UACGCAUGGAU-UUUGCAAUAUGUUGACAUCAGGGA--AAGACCGGG 151
MBFV-NC_002031.1   CUUGUNNNNNNN-AGGGAACAAAUAGUGGGACCAUAUUGACGCCAGGGA--AAGACCGGA 152
                   20.......130.......140.......150.......160.......170.......1
```

```
                   ..............................
NKV-NC_003635.1    GGUCGCACAGAUCAAG----------CUC 167
NKV-NC_004119.1    GCCAGAAAAGACUCAG----------AU- 167
ISFV-NC_027817.1   ACUAAAGUCGAGAGUC----------UCC 167
ISFV-NC_034204.1   AAUCCAAAGACUGUCC-------CCACC- 167
ISFV-NC_034017.1   AAUCCAAAGAAUGUCC-------CCACC- 167
ISFV-NC_001564.2   AACUCAUCUUAACCUC-------CCAC-- 167
ISFV-NC_005064.1   AACCCAGCUAAAUCUC-------CCACCG 167
ISFV-NC_012932.1   UACCCAUCUAAGCCUC-------CCA--- 167
ISFV-NC_008604.2   UGCAGCAGCAGGUCU--------CCA--- 167
ISFV-NC_021069.1   CGCAGCAGCAAAUCU--------CCAC-- 167
ISFV-NC_012671.1   CGCAGCAGCAAAUCU--------CCAC-- 167
ISFV-NC_024299.2   CUAUACAACAGGCUU--------CCACCG 167
ISFV-NC_027819.1   UCAGAUGGAAGAUCC--------CCA--- 167
TBFV-NC_003687.1   CGGUUCUUGUUCUC--------------- 167
TBFV-NC_005062.1   CGGUUCUUGUUCUCCCUGAGCCACCAUCA 167
TBFV-NC_001672.1   CGGUUCUUGUUCUCCCUGAGCCACCAUCA 167
TBFV-NC_001809.1   CGGUUCUUGUUCUC--------------- 167
TBFV-NC_027709.1   CGGUUCUUGUUCUC--------------- 167
TBFV-NC_003690.1   CGGUUCUUGUUCUC--------------- 167
MBFV-NC_001437.1   UCAACAUCAGCUACUA------------- 167
ISFV-NC_040610.1   AAAUACACAAUUUC--------------- 167
ISFV-NC_024017.1   AUAUGAUGUUGUUU--------------- 167
MBFV-NC_034151.1   AGACCCUCUGC-UUAU------------U 167
MBFV-NC_009028.2   AGAUCCUCUGC-UUUC------------C 167
MBFV-NC_040776.1   AGACCCUCUGC-UCUG------------C 167
MBFV-NC_001563.2   GGAUCUUCUGC-UCUG------------C 167
MBFV-NC_009942.1   AGAUCUUCUGC-UCUG------------C 167
MBFV-NC_006551.1   AGAUCUUCUGC-UCU-------------- 167
MBFV-NC_000943.1   AGAUCUUCUGC-UCUA------------- 167
MBFV-NC_009029.2   GGAUUUGCUGC-UCUG------------G 167
MBFV-NC_032088.1   AGAUCCGCUGC-UCUG------------G 167
MBFV-NC_007580.2   AGAUCCCCUGCUUUUU------------C 167
MBFV-NC_012534.1   AGA-CCCUCUGUCCUA------------A 167
MBFV-NC_015843.2   AGA-CCCUCUGUCCUA------------A 167
MBFV-NC_009026.2   GAAUUUGCUGCCUC-G------------C 167
MBFV-NC_012532.1   AGACUCCAUGAGUUUC------------C 167
MBFV-NC_035889.1   AGACUCCAUGAGUUUC------------C 167
MBFV-NC_012533.1   AGACUCUGUUGGCUUU------------C 167
MBFV-NC_001474.2   AGAUCCUGCUGUCUCC------------U 167
MBFV-NC_001477.1   AGAUCCUGCUGUCUCU------------A 167
MBFV-NC_001475.2   AGAUCCUGCUGUCUCC------------U 167
MBFV-NC_002640.1   AGAUCCUGCUGUCUCU------------G 167
ISFV-NC_017086.1   AGGUACUCGCUGAUUC------------A 167
MBFV-NC_008719.1   AGACACUCCUUGAUUC------------U 167
MBFV-NC_012735.1   AGACACUCCUUGAUUC------------- 167
ISFV-NC_016997.1   AGAUACCUUUGGUUCU------------U 167
ISFV-NC_027999.1   AGAAACUUCUUGCUUA------------- 167
NKV-NC_005039.1    UGUUUGUCGGUUCCGG------------- 167
MBFV-NC_002031.1   -GUGGUUCUCUGCUUU------------- 167
                   80.......190.......200.......
```