```
MBFV-NC_012735.1  .-UUUCUACUGAAAGGGGUAGAAGAA--------AGGAGAUUCAUUCCCAUGGCAACAAAG  52
MBFV-NC_008719.1  AUUUCUACUGAAAG-GUAGAGAAA-------AGGAGUUUUGCUUCUCAUGGCAACAAGA  52
MBFV-NC_002031.1  UAAUCGUUCGUUGAGCGAUUAGCAG---------AGAACUGAC-CAGAACAUGUC------  45
MBFV-NC_012532.1  AACAGGUUUU-------AAUUUGGAUUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAACCCC  52
MBFV-NC_035889.1  ACAGGUUUU----------AUUUUGGAUUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAAA--CCC  51
MBFV-NC_012533.1  AACAGGUUUU-------CGUUGUUGAGAAACGAGAG-UCUUCGGAAAGAAAA---CCC  50
MBFV-NC_002640.1  -------UCUA-ACAGUUUGUUUGAAUAGAGAGCAGAUCUCUGGAAAAAAUG----AA--CC  47
MBFV-NC_001474.2  -CGUAGUUCUA-ACAGUUU-AUUAUUAGAGAGCAGAUCUCUG-----AUGAAUAA--CC  50
MBFV-NC_001477.1  ACGUAGUUCUA-ACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG-----AUGAAUAA--CC  50
MBFV-NC_001475.2  ACGUAGUGCUG-ACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG-----AUGAACAA--CC  50
MBFV-NC_009028.2  U--U---UAACACAGUCUGAAUAGUUUUU--AGCAAGGGA-UUUCCCAUGUCU---AAA  49
MBFV-NC_040776.1  U--U---UAACACAGUCUGAAUAGUUUUUU--AGCAGAGGU-ACUACCAAGUCU---AAA  49
MBFV-NC_034151.1  -------UAACACAGUCUGAAUAGUUUUUUGAACAGAGGUGAAAACCAUGUCU---AAA  49
MBFV-NC_001563.2  A--U---UAACACAGUGCGAGCUGUUUCUUGGCAC-GAAGAACUC-GAUGUCU---AAG  49
MBFV-NC_009942.1  A--U---UAACACAGUCGGACAGUUUUUUUGGCAGC-GAAGAUCUC-GAUGUCU---AAG  49
MBFV-NC_000943.1  -------UAACGCAGUUGAACAGUUUUUUUGGAGC-GUUUUGAAUC-AAGUCU---AAA  49
MBFV-NC_006551.1  ---U---UAACACAGUUCCGGCAGUUUUUUGGAGC-GUUGAUUUUCAAUGUCU---AAA  49
MBFV-NC_001437.1  ---U---UAGUGCAGUUUAAACAGUUUUUUGGAGC-GUUGAUUUUCAAUGUCU---AAA  49
MBFV-NC_007580.2  ---U---AACUUGACUGCGAACAGUGUUUUUAGCAGGGAAU-UACCCAUGUCU---AAA  49
MBFV-NC_012534.1  CCGU-----AAACACAGUUUGAAC-GUUUUUUUGGAGGAGAAUUUAU---AUGUUUAACAAA  52
MBFV-NC_015843.1  ACGU-----AAACACAGUUUGAAC-GUUUUUUUGGAGAUAGAGACAACU---UGUCUAACAAA  52
MBFV-NC_009029.2  UGAUUACAUAAACACAGUGUGAACAGUUUUUUGAACAG-----C-----AUGAACC---AAA  49
MBFV-NC_032088.1  UC----UAUAACUCAGUGUGAACAAGUUUUUGUAUCAGUGGAU-----AUGACU---AAA  49
MBFV-NC_009026.2  AG---------AAAAGUUUUCGAGAAAGACAGAGAAGCGAAUUCUCUG-AAUGGCU---AGA  49
TBFV-NC_027709.1  --GUGUUGAGAAAAAGACAGCUUUAGGAGAACAAGAGCUGGGG-----AUGGGUCAAGAAG  52
TBFV-NC_001672.1  --GUGUUGAGAAAAAGACAGCUUUAGGAGAACAAGAGCUGGGG-----AUGGGCAAGAAG  52
TBFV-NC_001809.1  --GUGUUGAGAAAAAGACAGCUUUAGGAGAACAAGAGCUGGGG-----AUGGGCGGGAAG  52
TBFV-NC_005062.1  --GUGUUGAGAAAAAGACAGCUUUAGGAGAACAAGAGCUGGGG-----AUGGGCGGGAAG  52
TBFV-NC_003690.1  --GUGUUGAGAAAAAGACAGCUUUAGGAGAACAAGAGCUGGGA-----AUGGGCGGGAAG  52
TBFV-NC_003687.1  ---------AGAAAAAGACAGCUUAGGAGAACAAGAGCUGGGAGUGGGUUAUG------AUG  46
MBFV-NC_018705.3  CCGU-----AAACACAGUUUGAAC-GUUUUUUUGGAGAGACUACU----UGUCUAACAAA  52
                  ........10........20........30........40........50......6
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MBFV-NC_012735.1  G-GGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUGAUAUAUGGUAG--CGGCUAGA---  106
MBFV-NC_008719.1  G-GAGCAAGCAAGUCUCGGGUCACGUACCCGGGCGUGAUAUAUGGUGA--CGGCCAAG---  106
MBFV-NC_002031.1  ---UGGUCGUAAGUCUCAGGGAAAACCCUGGGCGUGAUAUAUGGU-A--CGACGAGGAGU  99
MBFV-NC_012532.1  A--AGAAGAAAU--CC--GGAGGA-UCCGGAUUGUGAUAUAUGUCUAAAACGCGGGAGUAGC  105
MBFV-NC_035889.1  A--AAAAAGAAAU--CC--GGAGGAUUCCGGAUUGUGAUAUAUGUCUAAAACGCGGGAGUAGC  105
MBFV-NC_012533.1  A--AAAAAGG---------GCCGCAUCGGUCCUCGAUAUAUGUCUAAAACGCUUUGGAUC  99
MBFV-NC_002640.1  A--ACGAAAAAAG--GU---G---GUUAGGACACCUUUGCGUGAUAUAUGUCUGAAACGCGAGAGAAAA  99
MBFV-NC_001474.2  A--ACGGAAAAAG--GC---G---AAAAACACGCCUUUGCGUGAUAUAUGUCUGAAACGCGCGAGAAA  102
MBFV-NC_001477.1  A--ACGGAAAAAG--AC---G---GUCGACCGUCUCAUGCGUGAUAUAUGUCUGAAACGCGUGAGAAA  102
MBFV-NC_001475.2  A--ACGGAAGAAG--AC---G---GGAAAACGCUCUAUGCGUGAUAUAUGUCUGAAACGCGUGAGAAA  102
MBFV-NC_009028.2  A--AACCAGGAAAGUCC--GCUGCAAAGCGGACGUGUGCAUAUAUGUCUAAAAGCCGUGGCUAG  105
MBFV-NC_040776.1  A--AACCAGGAAAAA---GCCGGAAAGCUGUGUGCAAUAUAUGUCUAAAAGCCCAGCCAG  105
MBFV-NC_034151.1  A--AACCAGGAGACCC--GGAAAGAACCGGGUCGUGUUAUAUGUCUAAAAGCCGCACACCCG  105
MBFV-NC_001563.2  A--AACCAGGAGGGCCC--GGUAAAAACGCGGGUCGUGUCAUAUAUGUCUAAAAGCGGGAAUGCC  105
MBFV-NC_009942.1  A--AACCAGGAGGGCCC--GGCAAGAGCCGGGCUGUGUCAUAUAUGUCUAAAAGCGGGAAUGCC  105
MBFV-NC_000943.1  A--AACCAGGAGGGCCC--GGGAAGCCCGGGUCGUGUCAUAUAUGUCUAAAAGCGGGCAUACC  105
MBFV-NC_006551.1  A--AACCAGGAGGGCCC--GGAAGAAACGGGUCAUCUAUAUGUCUAAAAGCGGGCAUACC  105
MBFV-NC_001437.1  A--AACCAGGAGGGCCC--GGUAAAAAGCCGGGUCGUGUCAUAUAUGUCUAAAAGCGGGCCUAAC  105
MBFV-NC_007580.2  A--AACCAGGAAAACCC--GGUAGAAAGCCGGGUCGUGUCAUAUAUGUCUAAAAGCGGGGCCUAUC  105
MBFV-NC_012534.1  A--AACCAGGAAAACCC--GGGGGCAACCCGGGUGUCAAUAUAUGUCUAAAAGCGCGGUUGCAUC  108
MBFV-NC_015843.1  A--AACCAGGAAGACCC--GGCUCAGGCCGGGUGUCAAUAUAUGUCUAAAAGCGCGGAACAGC  108
MBFV-NC_009029.2  A--AACCAGGAAGACCC--GGUAGAAACGGGCUGUCAUAUAUGUCUAAAAGCGCGGUUGCGUC  105
MBFV-NC_032088.1  A--AACCAGGAAAGUCC--GGCAAUCGUCGGACGUGUCAUAUAUGUCUAAAAGCGAGCGGUC  105
MBFV-NC_009026.2  A--AACCAGGAAGACCC--GGAGGAAACGGGUCGUGCAUAUAUGUCUAAAAGCGUACGGCGGC  105
TBFV-NC_027709.1  UCCAUCCUGAAAGGUAAGGGGGCGGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCAA--  110
TBFV-NC_001672.1  GCCAUCCUGAAAGGUAAGGGGGGCGGUCUCCCUCGACGAGUGUCGAAAGAGACCGCAA--  110
TBFV-NC_001809.1  GCCAUUCUGAAAGGCAAGGGGGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA--  110
TBFV-NC_005062.1  GCCAUUCUGAAAGGGAAGGGGGGUCCCCCUCGACGAGUGUCGAAAGAGACCGCUA--  110
TBFV-NC_003690.1  GCCGUUCUGAAAGGGAAGGGGGGUCCCCCUCGACGAGUGUCGAAAGUGGCCCCAA--  110
TBFV-NC_003687.1  ACCACUUCUGAAAGGGAAAGGGGGUGGCCCCUAGGCGCAAGCUUAAAAGUGACCGCAAAU  106
MBFV-NC_018705.3  A--AACCAGGAAGACCC---GGGGCAACCCGGGUGUGCAAUAUGUCUAAAAGCGCGGCGCAUC  108
                  0........70........80........90........100.......110.......1
```

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MBFV-NC_012735.1  -------GUGANNNNNNN-----------C-CCAUUCCCUCCCGAGCACACAUAGC-GGAC  147
MBFV-NC_008719.1  -------GCCANNNNNNN----------GAGGAGACCCUCCCGAGCAC-UAUAGC-GGAC  147
MBFV-NC_002031.1  UCGCUCCUUGUNNNNNNN----------GAGACCCUCCCAGGGAACAAAUAGUGGGAC  146
MBFV-NC_012532.1  CCGUG-------NNNNNNN------GGUUAGAGGAGACCCCCCGGAAAACGCAAAACAG  151
MBFV-NC_035889.1  CCGUG-------NNNNNNN------GGUUAGAGGAGACCCCCCGGAAAACGCAAAACAG  150
MBFV-NC_012533.1  UAGCGCUGGCCNNNNNNN------GAGACCCCCCGCGCCCAUAACCAACAUAAAACAG  151
MBFV-NC_002640.1  CCGCGUAUCAANNNNNNN----ACUAGAGGUUAGAGGAGACCCCCCCAACA--CAAAAACAG  155
MBFV-NC_001474.2  CCGCGUGU----NNNNNNN---CUAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAAACAG  155
MBFV-NC_001477.1  CCGCGUGU----NNNNNNN---CUAGAGGUUAGAGGAGACCCCCC-GCCAACAACAAAACAG  155
MBFV-NC_001475.2  CCGUGUGU----NNNNNNN---ACUAGAGGUUAGAGGAGACCCCCC-G---CAAAUAAAAACAG  155
MBFV-NC_009028.2  CGUGU-----NNNNNNN----GUGGAGACCCUCAGCAC-CAAGCGCGAA--ACAAACAG  153
MBFV-NC_040776.1  CGUUU-----NNNNNNN----A-GGAGACCCUUGGCACACAAGCGCAAA-GACAAACAG  154
MBFV-NC_034151.1  CGCUA-----NNNNNNN-------AGACCCUUGGGCAGUGAA-CGCAAA-UAAAAACAG  150
MBFV-NC_001563.2  CCGCG-----NNNNNNN----GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAAACAG  153
MBFV-NC_009942.1  CCGCG-----NNNNNNN----GGAGACCCCGUGCCACACAAGCACCACA-ACAAAACAG  153
MBFV-NC_000943.1  CCGCG-----NNNNNNN----GACCCCACUCUCAAAAGCAUC-AA-AU--AACAG  147
MBFV-NC_006551.1  CCGCG-----NNNNNNN----ACCCCGCA-UCAUAAAGCAUCAAA-AA--AACAG  146
MBFV-NC_001437.1  CCGCG-----NNNNNNN------UAGAGGAGACCCCGUCAUUUGCAUC-A---AACAG  147
MBFV-NC_007580.2  CCGCG-----NNNNNNNUUAGAGGAGACCCCUUGCGUUAAC-----GC-AAACAACAG  153
MBFV-NC_012534.1  CC--------NNNNNNN----GGAGACCCUUGCGAGGUUAACACCAAA-ACAAAACAG  153
MBFV-NC_015843.1  CC--------NNNNNNN------AGGAGACCCCUUGCGAGUGAGACACCACA-A-GAAACAG  153
MBFV-NC_009029.2  CCGCG-----NNNNNNNAGAGG---AGACCCUCCCACAAAGAAGCGCAAACACAAAACAG  157
MBFV-NC_032088.1  CCGCG-----NNNNNNN----GGAGGUUAGAGGAGACCCCCCACAUACCGCAAA-ACAAAACAG  158
MBFV-NC_009026.2  CAACG-----NNNNNNNCCCCGUG------GCAAAACAGUUAACGCACCAAAAGAAAACAG  154
TBFV-NC_027709.1  ----------NNNNNNNCUCUCUUCAGGAUUUUUUCUCCUCCU-----------AUAC  147
TBFV-NC_001672.1  ----------NNNNNNNCUCUCUUCAGGAUUUUUUCUCCUCCU-----------CAUCCCU  147
TBFV-NC_001809.1  ----------NNNNNNN-UCUCUUCAGGGGUUUUUCUCCUCCU-----------AUAC  146
TBFV-NC_005062.1  ----------NNNNNNNCUCUCUUCAGGAAAUUUUCUCCUCCU-----------AUAC  147
TBFV-NC_003690.1  ----------NNNNNNNCUCUCUUCAGGAUAUUUCUCCUCCU-----------AUAC  147
TBFV-NC_003687.1  AAGU------NNNNNNNCU-UCCUCGGGAUUUUUCUCCUCCU-----------AUAC  146
MBFV-NC_018705.3  CC--------NNNNNNNCCCCGCGGGAAAUCAAAGCAAGGCCCAACCUG--GCAAAGCUG  157
                  20.......130.......140.......150.......160.......170.......1
```

```
MBFV-NC_012735.1  CAUAUUGACGCCAGGGAA--AGACCGG-AGACAUCCUUGAUU-CUCACCUUUCUCGCCC  203
MBFV-NC_008719.1  CAUAUUGACGCCUGGGAA--AGACCGG-AGACAUCCUUGAUU-CUCACCUUUCUCACCC  203
MBFV-NC_002031.1  CAUAUUGACGCCAGGGAA--AGACCGG-AGUGGUUCUCUCAAGAAGGUCUG  202
MBFV-NC_012532.1  CAUAUUGACG--UGGGAA--AGACCAGAGACU-CCAUGAGUUU--CCACCACGCUGGCCGC  205
MBFV-NC_035889.1  CAUAUUGACG--CUGGGAA--AGACCAGAGACU-CCAUGAGUUU--CCACCACGCUGGCCGC  205
MBFV-NC_012533.1  CAUAUUGACACCUGGGAA--AGACCGGAGACUCUGUUGUGCUUU-CC--CCACACCAGCCGC  207
MBFV-NC_002640.1  CAUAUUGACG--CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---UGCAACAUCAAUC  207
MBFV-NC_001474.2  CAUAUUGACG--CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---UUCAACAUCAUUC  207
MBFV-NC_001477.1  CAUAUUGACG--CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---UACAGCAUCAUUC  207
MBFV-NC_001475.2  CAUAUUGACG--CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---UUCAACAUCAUUC  207
MBFV-NC_009028.2  CAUAUUGACGCCUGGGAA--AGACCGGGAGAUCCU--CCAUCACCAGCCA  206
MBFV-NC_040776.1  CAUAUUGACACCUGGGAA--AGACCGGGAGACCU-CUGCUCU----UGCAACACCAGCCA  207
MBFV-NC_034151.1  CAUAUUGACACCUGGGAU--AGACUGGGAGACCU-CUGCUUA--UUUCAACACCAGCCA  205
MBFV-NC_001563.2  CAUAUUGACACCUGGGAA--UAGACUAGGGAGAUCU--CUGCU----GCAACAACAGCCA  206
MBFV-NC_009942.1  CAUAUUGACACCUGGGA---UAGACUAGGAGAUCU--CUGCUC----GCAACAACAGCCA  204
MBFV-NC_000943.1  CAUAUUGACACCUGGGAA--AGACUAGGAGAUCU--CUGCUCU--AUUCCAACAUCAGUCA  204
MBFV-NC_006551.1  CAUAUUGACACCUGGGAA---UGACUAGGAGAUCU--CUGCUCU--AUUCCAACAUCAACCA  204
MBFV-NC_001437.1  CAUAUUGACACCUGGGAAU--AGACUGGGAGAUCU--CUGCUCU--AUCUCAACAUCAGUCA  205
MBFV-NC_007580.2  CAUAUUGACACCUGGAAGA----CAGGAGACCCU-CUGCUUU----UUCAACACCAGCCA  203
MBFV-NC_012534.1  CAUAUUGACACCUGGGAA---GAGACUAGGAGUCU--CUGCUCU-AA--CAACACCAGCCA  207
MBFV-NC_015843.1  CAUAUUGACACCUGGGAA---UAGACUAGGAGAUCU--CUGCUCU----CAACACCAGCCA  207
MBFV-NC_009029.2  CAUAUUGACACCUGGGAA---UAGACUAGGAGAUUUG-CUGCUCU----UGGACUUCCGGCU  209
MBFV-NC_032088.1  CAUAUUGACACCUGGGA---UAGACUAGGAGACCG-CUGCUCU----UGGAUCCCAGCC  210
MBFV-NC_009026.2  CAUAUUGACACCUGGGA---UAGACUGG-AGAAUU----UGCUUGC-CUCGCAACCACUCCCA  207
TBFV-NC_027709.1  CAAA--UUCCCCCUUCGAC----AGAGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCAUCAC  203
TBFV-NC_001672.1  AAAA--UUCCCCCUUCGGU----UAGAGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCAUCAC  203
TBFV-NC_001809.1  CAAA--UUCCCCCUUCGAC----AGAGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCAUCAC  203
TBFV-NC_005062.1  CAAA--UUCCCCCUUCAUC----UGAGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCAUCAC  203
TBFV-NC_003690.1  CAAA--UUCCCCCUUCGUC----UGAGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCAUCAC  203
TBFV-NC_003687.1  UAAAU-UUCCCCCAGGAA----ACUGGGGGGGCGGUUCUUGUUCUCCCGGAGCCACCACCAU  203
MBFV-NC_018705.3  CAAA--CCAGGGGAACUAGAGGUUAGAGGAGACCCCU---------UGCGAGUGAACA  207
                  80........190.......200.......210.......220.......230.......2
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MBFV-NC_012735.1  UAAAGCACAGAUUGCUU--GAAUGCAGGGCGG-GGAAGUUGAGAACCAACUAGUGUCU  258
MBFV-NC_008719.1  UUAAGCACAGAUUGCUU--GAAUGCAGUGGUGG-GGAAGUUGGGAACCAACUAGUGUCU  258
MBFV-NC_002031.1  U-GAGCACAGUUUGCUUCAAGAAUAAGCAGACC--UUUUGGAUGACAAACACAAAACCACU  258
MBFV-NC_012532.1  CA-GGCACAGAUCGCCGA--ACUUCGGCGGCC-GGUGUGGG-GAAAUCCAUGGUUUCU  258
MBFV-NC_035889.1  CA-GGCACAGAUCGCCGA--ACUUCGGCGGCC-GGUGUGGG-GAAAUCCAUGGUUUCU  258
MBFV-NC_012533.1  CU-GGCACAGAUCGCCGA----AGAUCGGUGGU-GUGGUGGG-GCAACCCAACAGGACU  258
MBFV-NC_002640.1  CA-GGCACAGAACGCCGC--AAGAUGGAUUUG-UGUCUGU----UGAUCCAACAGGUUCU  258
MBFV-NC_001474.2  CA-GGCACAGAACGCCAG--AAAAUGGAUUGG-UGCUGU---UGAUCCAACAGGUUCU  258
MBFV-NC_001477.1  CA-GGCACAGAACGCCAA--AAAAUGGAUGG-UGCUGU---UGAUCCAACAGGUUCU  258
MBFV-NC_001475.2  CA-GGCACAGAACGCCAG--AAAAUGGAUUGG-UGCUGU---UGAUCAACAGGUUCU  258
MBFV-NC_009028.2  CUAGGCACAGAUCGCCGC--AAGUAGGUGGCUG-GUGGUGAA---AACACAUGGAUCU  258
MBFV-NC_040776.1  CUAGGCACAGAUCGCCGA--AAUCUGGUGCUG-GUGCUGCA---AACACAGGAUCU  258
MBFV-NC_034151.1  CUCGGCACAGACGCCGU--AAGU-GUGGCUG-GUGUGUGAAAAUAAACAGAGAUCU  258
MBFV-NC_001563.2  CACGGCACAGUGCGCCGA--CAUAGGUGGCUG-GUGGUGC--UAGAACACA-GGAUCU  258
MBFV-NC_009942.1  CACGGCACAGUGCGCCGA--CAAUGGUGGCUG-GUGGUGC--GAGAACACA-GGAUCU  258
MBFV-NC_000943.1  CAAGGCACAGACGCCGA--ACACUGUGACUG-AUGGGGGAAGAACAACACA-GGAUCU  258
MBFV-NC_006551.1  CAAGGCACAGACGCCGC--AAAUUGUGAGCUG-GUGGGGGAAGAACAACACA-GGAUCU  258
MBFV-NC_001437.1  CUAGGCACAGACGCCGA--AGUAUGUAGCUG-GUGGUGAGAAAAAACACA-GGAUCU  258
MBFV-NC_007580.2  CAGGGCACAGACGCCGU--AAG-UGUGGCUG-GUGGUG--GGACACA-GUGUCU  258
MBFV-NC_012534.1  UCCGGCACAGACGCCGU-UAAG-UGUGGCUG-GUGGUG--GUAGAACACA-GUGUCU  258
MBFV-NC_015843.1  CUCGGCACAGAC-GCCGA--AAG-UGUGGCUG-GUGGUG--GUAGAACACA-GUGUCU  258
MBFV-NC_009029.2  CUCGGCACAGAACGCCGU--UGA--GGAGCCGG-AGGCCCAA----AACAC-CAGAUCU  258
MBFV-NC_032088.1  CUAGGCACAGAACGCCGA--CA-UGACGGG-AGGCCCAA----AACAC-CGAUCU  258
MBFV-NC_009026.2  CCGGCACAGAAGACGCCGA--CA--UGUGGGGAG-GGGCGUGAGACACCA---GAUUCU  258
TBFV-NC_027709.1  CCAGAACAGAUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGU  258
TBFV-NC_001672.1  CCAGAACAGAUGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGU  258
TBFV-NC_001809.1  CCAGAACAGUAGUCUGACAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGU  258
TBFV-NC_005062.1  CCAGAACAGCAGUCUAACAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGU  258
TBFV-NC_003690.1  CUAGAACAGUAGUCUGAAAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGCU  258
TBFV-NC_003687.1  CUAGAACAGCAGUCUGAAAAGGAGGUGAUGCGUGACUCGGAAAACAG---CCGU  258
MBFV-NC_018705.3  CCACAAAAAACAGCAUAUUGACACCUGGGAUGGGGAAAGACUAGGAGACCCUC------UGU  258
                  40.......250.......260.......270.......280.......290.......
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