```
                                     ..................(((((((((((((((((..(--.......................((.(((((
dISFV-NC_016997.1   C-AUUAGAAUAGGUUUGAAAGAGUUGAAU--CA-------AUGAGAUCAAUGGUCUUUCG 50
dISFV-NC_040610.1   AAAGAAGUCAAAACUAGAAAAUUUAA-UUU--CA-------AUG---------GUUAGACA 41
dISFV-NC_024017.1   UGAU-AGAGUUGACUAGAUAACGAAUUAU--CG-------AUG---------GUGACUAA 41
dISFV-NC_017086.1   UGAAUAGGUUUAUUGUGAAUCGAAA-UAC--CG-------AUG---------GCGAAUAA 41
dISFV-NC_027999.1   UUGUUUGAGAGAAC---AGCGAGAAGUACUACU-------AUG---GUUGAGAAAACUAG 47
TBFV-NC_003690.1    AAAG----ACAGCUUAGGAGAACAAGAGCUGGGA------AUGGCCGGGAAGGCCGUUCU 50
TBFV-NC_001809.1    AAAG----ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGACCAUCCU 50
TBFV-NC_027709.1    AAAG----ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGGCAGGAAGUCCAUCCU 50
TBFV-NC_005062.1    AAAG----ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGCCGGGAAGGCCAUUCU 50
TBFV-NC_001672.1    AAAG----ACAGCUUAGGAGAACAAGAGCUGGGG------AUGGUCAAGAAGGCCAUCCU 50
TBFV-NC_003687.1    ---------AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG------AUGACCACUUC 44
                    .........10........20........30........40........50.......6

                                     .........((((((.............................................
dISFV-NC_016997.1   AGCUAGGAGGCCCCGUU---AAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC------- 100
dISFV-NC_040610.1   ACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGC--GGAUUACCACCCG- 98
dISFV-NC_024017.1   AUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACAC--GCAUCACCAAGCG- 98
dISFV-NC_017086.1   ACCCAAGAAGCCCGG---AAGACGGGCUAUCGAUAGAUGGAGACGUGCGUUGCCUCGCGU 98
dISFV-NC_027999.1   AGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUGAGGCGUAC---------UGG 98
TBFV-NC_003690.1    AAAAGGAAAGGGGGGGGGGUCCCC--CUCGACGAGCGUCGAAAGUGGCCC-----CAA--- 100
TBFV-NC_001809.1    GAAAGGCAAGGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGACCG----CGA---- 100
TBFV-NC_027709.1    GAAAGGUAAGGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGACCG----CAA---- 100
TBFV-NC_005062.1    GAAAGGAAAGGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGACCG----CUA---- 100
TBFV-NC_001672.1    GAAAGGUAAGGGGGGGCGGUCCCC--CUCGACGAGUGUCGAAAGAGACCG----CAA---- 100
TBFV-NC_003687.1    UAAAGGAAAGGGGGGGCGGUCCCC--CUAGGCGCAAGCUUAAAGUGACCG----CAAAUAA 98
                    0........70........80........90........100.......110.......1

                                     ...............................................))))))).......))))).)))))))
dISFV-NC_016997.1   --NNNNNNNCAAAAAU-GU-AUAGGCUAUUUAUAUCGACACCUAGG--AAAGACUGGGAGAU 154
dISFV-NC_040610.1   CUNNNNNNNNCUCAUUA-UUUUAUAUAACAGCAUAUUGACA-CCAGGG-AUAGACCGGAGAA 155
dISFV-NC_024017.1   UGNNNNNNNNCU-GCUG-AGAGCAAAAUAGCAUAUUGACA-CUUGGGAAGAGACCAGAGAU 155
dISFV-NC_017086.1   CUNNNNNNNNCCCAAAA-AUGA-AGGGCACUAUAUCGACACUUGGG--AAAGACCAGAGGU 154
dISFV-NC_027999.1   AANNNNNNNNC-CAUUG-GAAAUUUAGCUUCAUAUUGACACC-AGGGAAAAAGACCGGAGAA 155
TBFV-NC_003690.1    --NNNNNNNGAUAUUUCCUCCUCCUAUACCAAA-UUCCCCUCGUCAGAGGGGGGCGGU 157
TBFV-NC_001809.1    --NNNNNNNG-UUUUCCUCCUCCUAUACCAAAUUCCCCUCGACAGAGGGGGGCGGU 157
TBFV-NC_027709.1    --NNNNNNNGAUUUUCCUCCUCCUAUACCAAA-UUCCCCUCGACAGAGGGGGGCGGU 157
TBFV-NC_005062.1    --NNNNNNN--------------UAUACCAAA-UUCCCCUCAUCUGAGGGGGGCGGU 142
TBFV-NC_001672.1    --NNNNNNN--------------UAUACAAAA-UUCCCCUCGGUAGAGGGGGGCGGU 142
TBFV-NC_003687.1    GUNNNNNNN-AUUUUUCCGCCUCCUAUACUAAAAUUUCCCCAGGAAACUGGGGGGGCGGU 157
                    20.......130.......140.......150.......160.......170.......1

                    )))))).))))))..............).
dISFV-NC_016997.1   ACCUUUGGUUC-------------UU 167
dISFV-NC_040610.1   AUACACAAUUU-------------C- 167
dISFV-NC_024017.1   AUG-AUGUUGU-------------UU 167
dISFV-NC_017086.1   ACUCGCUGAUU-------------CA 167
dISFV-NC_027999.1   ACUU-CUUGCU-------------UA 167
TBFV-NC_003690.1    UCUUG-UUCUC-------------- 167
TBFV-NC_001809.1    UCUUG-UUCUC-------------- 167
TBFV-NC_027709.1    UCUUG-UUCUC-------------- 167
TBFV-NC_005062.1    UCUUG-UUCUCCCUGAGCCACCAUCA 167
TBFV-NC_001672.1    UCUUG-UUCUCCCUGAGCCACCAUCA 167
TBFV-NC_003687.1    UCUUG-UUCUC-------------- 167
                    80.......190.......200....
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