```
                    ..................................(((((.....((((..((((.............(((((.......((((.....))))).(((((((((((..
dISFV-NC_024017.1   UC-----GCGCAUUUUGAUAGAGUU---GACUAGAUAACGAAUUAUCGAUGGUG------ACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAAC 93
dISFV-NC_040610.1   -------GAACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAAUU-UCAAUGGUU------AGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAA 93
dISFV-NC_027999.1   GGAUUAAUUGUUGUUUGAGAGAAC----AGCGAGAAGUACUACU----AUGGUUGAGAAAACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAG 99
dISFV-NC_017086.1   ----AUAACACGUUUGAAUAGGUUU----AUUGUGAAUCGAAAUACCGAUG---------GCGAAUAAACCCAAGAAGCCCGGAAGACGGGCUAUCGAUAUAGUGAG 90
dISFV-NC_016997.1   AAACAUAACGCAUUAGAAUAGGUUU--------GAAAGAGUUGAAUCAAUGAGAUCAAUGGUCUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAA 99
                    .........10........20........30........40........50........60........70........80........90........100.....

                    .(((.........................))).......(((...........))))))))))))))))))))).....))))..))))........)))))....
dISFV-NC_024017.1   ACGCAUCACCAAGCGUG---NNNNNNNCCCC---ACUGCUGAGAG------CAAAAUAGCAUAUUGACACUUGGGAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAU 188
dISFV-NC_040610.1   GCGGAUUACCACCCGCU---NNNNNNNACCC---CCUCAUUAUUUA-----UAUAACAGCAUAUUGACACCAGGG-AUAGACCGGAGAAAU-ACACAAUUUCUGGGC 187
dISFV-NC_027999.1   GCGUACUG------GAA---NNNNNNNUGGCGAACCCCAUUGGAAA-----UUUAGCUUCAUAUUGACACCAGGGAAAAGACCGGAGAAACUUCUUGCUUACUCGCA 192
dISFV-NC_017086.1   ACGUGCGUUGCCUCGCGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGACACUUGGGAA-AGACCAGAGGUACUCGCUGAUUCACCGC- 193
dISFV-NC_016997.1   ACGUAAGCUGC---------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAGGAA-AGACUGGAGAUAC-CUUUGGUUCUUCGA- 192
                    ..110.......120.......130.......140.......150.......160.......170.......180.......190.......200.......210..

                    .....(((((((..(((.......)))).......)))))))).............................
dISFV-NC_024017.1   GCUGACAACAGCCAUGGCACAGAGCGCCUGAAGAGAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU--- 258
dISFV-NC_040610.1   ACUAAACAACUACA-AGCACAGAGCGCUUCGAGAAGUAGUUGUUUCUUGACGAGAGAAUAU-GUAUAUCUUGG 258
dISFV-NC_027999.1   GC-AACCACCGAAA-GGCACAGUAUGCCUGAAACAUUCGGUGGUUGAUGUAGUAAACAACCAGUUUCU----- 258
dISFV-NC_017086.1   -CACCAG-ACUACACGGCACAGCGCGCCGGAAAAGGUAGU-UUGGAGGUUGUAAAACAACAAGUAUCU----- 258
dISFV-NC_016997.1   -CACUAACUCCUAAUGGCACAGAGCGCCAUAAGAAACGGA-GUUAGUAUGGAAAAAACAUAAGUAUCU----- 258
                    .....220.......230.......240.......250.......260.......270.......280.....
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