```
              .........(( .. (((((.................(((( .. ((................
dISFV-NC_024017.1  UCGCGCAUUUUGAUAGAGU-U---GACUAGAUAACGAAUUAUCGAUG---------GUGA  47
dISFV-NC_040610.1  --GAACAAUUUUAAAGAAG-UCAAAACUAGAAAUUUAAUU-UCAAUG---------GUUA  47
dISFV-NC_017086.1  ----AUAACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG---------GCGA  47
dISFV-NC_016997.1  AAACAUAACGCAUUUGAAUAGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGUCU  56
dISFV-NC_027999.1  GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA----CUAUG---GUUGAGAAAA  53
              .........10........20........30........40........50.......6

              ..................(((.....((((.((((((((((((((((................
dISFV-NC_024017.1  CUAAAUCAAGAAAGCCCACCGCAAAGCGGGCUGUUAAUAUGCUAAACACGCA--UCACCAA  105
dISFV-NC_040610.1  GACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGA--UUACCAC  105
dISFV-NC_017086.1  AUAAACCCAAGAAGCCCGG---AAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUC  104
dISFV-NC_016997.1  UUCGAGCUAGGAGGCCCGU---UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC---  110
dISFV-NC_027999.1  CUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUA---------  104
              0........70........80........90........100.......110........1

              ..............................)))))))))))))))))))))).
dISFV-NC_024017.1  GCG-UGNNNNNNNN--------CCCCACUGCUGAGAG-CAAAAUAGCAUAUUGACACUUGG  155
dISFV-NC_040610.1  CCG-CUNNNNNNNN--------ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGG  156
dISFV-NC_017086.1  GCGUCUNNNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCUACAUAUCGACACUUGG  162
dISFV-NC_016997.1  ------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAG  162
dISFV-NC_027999.1  CUGGAANNNNNNNUGGCGA-ACCCCAUUGGAAAUUU----AGCUUCAUAUUGACACCAGG  159
              20.......130.......140.......150.......160.......170.......1

              .....))))))..)))))........)))))).))
dISFV-NC_024017.1  GAAGAGACCCAGAGAUAUGAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACAGAGCGC  215
dISFV-NC_040610.1  G-AUAGACCGGAGAAAUACACAAUUUCUGGGCACUAAACAACUACAA--GCACAGAGCGC  213
dISFV-NC_017086.1  GAA-AGACCCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA--CACGGCACAGCGCGC  218
dISFV-NC_016997.1  GAA-AGACUGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCU--AAUGGCACAGAGCGC  218
dISFV-NC_027999.1  GAAAAGACGGAGAAACUUCUUGCUUACUCGCAGCAACCACCG--AAAGGCACAGUAUGC  217
              80.......190.......200.......210.......220.......230.......2

              ....................................................
dISFV-NC_024017.1  CUGAAG-A--GAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU  258
dISFV-NC_040610.1  UUCGAG-AAGUAGUUGUUUCUUGACGAGAGAAUAUGUAUAUCUUGG  258
dISFV-NC_017086.1  CGGAAA-AGGUAGUUUUGGAGGUUGUAAAACAACAAGUA-----UCU  258
dISFV-NC_016997.1  CAUAAG-AAACGGAGUUAGUAUGGAAAAAACAUAAGUA---UCU  258
dISFV-NC_027999.1  CUGAAACAUUCGGUGGUUGAUGUAGUAAACAACCAGU-----UUCU  258
              40.......250.......260.......270.......280....
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