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

                    ..................(((......((((.(((((((((((.................
dISFV-NC_024017.1   CUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAACACGCA--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-UGNNNNNNN--------CCCCACUGCUGAGAG-CAAAAUAGCAUAUUGACACUUGG 155
dISFV-NC_040610.1   CCG-CUNNNNNNN--------ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGG 156
dISFV-NC_017086.1   GCGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGACACUUGG 162
dISFV-NC_016997.1   ------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAG 162
dISFV-NC_027999.1   CUGGAANNNNNNNNUGGCGA-ACCCCAUUGGAAAUUU----AGCUUCAUAUUGACACCAGG 159
                    20.......130.......140.......150.......160.......170.......1

                    .....)))))..))))).......)))))).))..............................
dISFV-NC_024017.1   GAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACAGAGCGC 215
dISFV-NC_040610.1   G-AUAGACCGGAGAAAUACACAAUUUCUGGGCACUAAACAACUACAA--GCACAGAGCGC 213
dISFV-NC_017086.1   GAA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA--CACGGCACAGCGCGC 218
dISFV-NC_016997.1   GAA-AGACUGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCU--AAUGGCACAGAGCGC 218
dISFV-NC_027999.1   GAAAAGACCGGAGAAACUUCUUGCUUACUCGCAGCAACCACCG--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-AGGUAGUUUGGAGGUUGUAAAACAACAAGUA-----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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