```
                  ...............(((((.....................((((..((.........((..
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 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 GCGUCUNNNNNNAGAGGAGACCCCCCCAAAAAUG--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 GAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACA 209
dISFV-NC_040610.1 G-AUAGACCGGAGAAAUACACAAUUUCUGGGCACUAAACAACUACAAGCACAGA 209
dISFV-NC_017086.1 GAA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA-----CACGGC 209
dISFV-NC_016997.1 GAA-AGACUGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCU-----AAUGGC 209
dISFV-NC_027999.1 GAAAAGACCGGAGAAACUUCUUGCUUACUCGCAGCAACCACCGA----AAGGCA 209
                  80.......190.......200.......210.......220.......230..
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