```
                    ..............................(((((......((((..((.........((..
dISFV-NC_016997.1   AAACAUAACGCAUUAGAAUAGGUUU----GAAAGAGUUGAAUCAAUGAGAUCAAUGGUCU 56
dISFV-NC_017086.1   ----AUAACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG---------GCGA 47
dISFV-NC_040610.1   --GAACAAUUUUAAAGAAGUCAAAAC-UAGAAAUUU-AAUUUCAAUG---------GUUA 47
dISFV-NC_024017.1   UCGCGCAUUUUGAUAGAGUU---GAC-UAGAUAACGAAUUAUCGAUG---------GUGA 47
dISFV-NC_027999.1   GGAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUA----CUAUG---GUUGAGAAAA 53
                    .........10........20........30........40........50........6

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

                    ..........................................))))))))))))))))))
dISFV-NC_016997.1   ------NNNNNNN--AGGAAACCCCCCCAAAAAUGUAUAGGCUAUUAUAUCGACACCUAG 162
dISFV-NC_017086.1   GCGUCUNNNNNNNAGAGGAGACCCCCCCAAAAAUG--AAGGGCACUAUAUCGACACUUGG 162
dISFV-NC_040610.1   CCGCU-NNNNNNN--------ACCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGG 156
dISFV-NC_024017.1   GCGUG-NNNNNNN--------CCCCACU-GCUGAGAGCAAAAUAGCAUAUUGACACUUGG 155
dISFV-NC_027999.1   CUGGAANNNNNNNUGGCGAACCCCAUUGGAAAUU-----UAGCUUCAUAUUGACACCAGG 159
                    20.......130.......140.......150.......160.......170.......1

                    ).....))))..))))........)))))).............................
dISFV-NC_016997.1   GAA-AGACUGGAGAUAC-CUUUGGUUCUUCGACACUAACUCCUA--AUGGCACAGAGCGC 218
dISFV-NC_017086.1   GAA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUAC--ACGGCACAGCGCGC 218
dISFV-NC_040610.1   G-AUAGACCGGAGAAAU-ACACAAUUUCUGGGCACUAAACAACUACA-AGCACAGAGCGC 213
dISFV-NC_024017.1   GAAGAGACCAGAGAUAUGAUGUUGUUUGUCAAUGCUGACAACAGCCAUGGCACAGAGCGC 215
dISFV-NC_027999.1   GAAAAGACCGGAGAAACUUCUUGCUUACUCG-CAGCAACCACCGAAA-GGCACAGUAUGC 217
                    80.......190.......200.......210.......220.......230.......2

                    ...............................................
dISFV-NC_016997.1   CAUAAGAA--ACGGAGUUAGUAUGGAAAAAACAUAA-GUAUCU--- 258
dISFV-NC_017086.1   CGGAAAAG--GUAGUUUGGAGGUUGUAAAACAACAA-GUAUCU--- 258
dISFV-NC_040610.1   UUCGAGAAGUAGUUGUUUCUUGACGAGAGAAUAU-GUAUAUCUUGG 258
dISFV-NC_024017.1   CUGAAGAGAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAUCU--- 258
dISFV-NC_027999.1   CUGAAACAUUCGGUGGUU--GAUGUAGUAAACAACCAGUUUCU--- 258
                    40.......250.......260.......270.......280....
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