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

                     ...............................(((.........((((.(((((((((((((((................
dISFV-NC_024017.1    CUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAAUAUGCUAAACACGCA--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--------CCCACUGCUGAGAG-CAAAAUAGCAUAUUGACACUUGGG 155
dISFV-NC_040610.1    CCG-CUNNNNNNN--------CCCCCUCAUUAUUUAUAUAACAGCAUAUUGACACCAGGG 156
dISFV-NC_017086.1    GCGUCUNNNNNNNNGAGGGAGACCCCCCCAAAAAUG--AAGGGCACUAUUGACGACUUGGG 162
dISFV-NC_016997.1    ------NNNNNNN--GGAAACCCCCCCCAAAAAUGUAUAGGCUAUUAUAUCACCUAGG 162
dISFV-NC_027999.1    CUGGAANNNNNNNNGGCGA-ACCCCAUUGGAAAUUU----AGCUUCAUAUUGACACCAGGG 159
                     20.......130.......140.......150.......160.......170.......1

                     .....)))))..))))).......)))))...............................
dISFV-NC_024017.1    AAGAGACCAGAGAUAUGAUGUGUGUUUGUCAAUGCUGACAACAGCCAUGGCACA 208
dISFV-NC_040610.1    -AUAGACCGGAGAAAUACACAAUUUCUGGGCACUAAACAACUACAAGCACAGA 208
dISFV-NC_017086.1    AA-AGACCAGAGGUACUCGCUGAUUCACCGCCACCAG-ACUA-----CACGGC 208
dISFV-NC_016997.1    AA-AGACUGGAGAUAC-CUUUUGGUUCUUCGACACUAACUCCU-----AAUGGC 208
dISFV-NC_027999.1    AAAAGACCGGAGCAAACUUCUUGCUUACUCGCAGCAACCACCGA----AAGGCA 208
                     80.......190.......200.......210.......220.......230.
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