```
                                          1
lcl|NC_009127.1_cds_YP_001096113.1_77   MAPHLRSRSL TARHLTESRT EGLSFHGQRP DGTILAVDRG

                                          41
lcl|NC_009127.1_cds_YP_001096113.1_77   RLLCTHADID ELERTLLGSS APVSHKDLVP IYCDRCLPYT

                                          81
lcl|NC_009127.1_cds_YP_001096113.1_77   LPWSQAPVSH DRDGLLPAHG YTVVGTAGVF SRDLVETKGM

                                          121
lcl|NC_009127.1_cds_YP_001096113.1_77   EKTSQHYHLP LRDLELFTHT QKSNRVLLVH CDIPIGSVIC

                                          161
lcl|NC_009127.1_cds_YP_001096113.1_77   YWHVHDRENA HCAVNALIVI TDQAFHTIFS RCKGLSWTTT

                                          201
lcl|NC_009127.1_cds_YP_001096113.1_77   SVVASELSLV GIPGRASSLI HSISRLSDLT QGTLPVSRKP

                                          241
lcl|NC_009127.1_cds_YP_001096113.1_77   DLRSLGYGRN SPRHADVVMA ALSKGTENPE ETASDQRREW

                                          281
lcl|NC_009127.1_cds_YP_001096113.1_77   KQMVEMLMGQ LDLLYEATIA KGPDDDGQAS KRYLSEGRDL

                                          321
lcl|NC_009127.1_cds_YP_001096113.1_77   FKKLHELKAH LEKLESASTP AAVSAAAPAP STSVPAPQQT

                                          361
lcl|NC_009127.1_cds_YP_001096113.1_77   TETPKEDSRA GTPEASVTSQ QAQSAPVPSQ PVTSVPSQPV

                                          401
lcl|NC_009127.1_cds_YP_001096113.1_77   TSVPSQPVTS VPSQPVTSVP SQPVTSVPSQ PVTSVQQPNK

                                          441
lcl|NC_009127.1_cds_YP_001096113.1_77   MASSDDIRDA VAKALADVIK PSIQQYTPPP QPNHGGGATG

                                          481
lcl|NC_009127.1_cds_YP_001096113.1_77   FRDFVEMFKL MQQMHAPAQA PAPVAAPAPV VAQVPQSLSH

                                          521
lcl|NC_009127.1_cds_YP_001096113.1_77   HGLVDDEHPS TSAHQRGGKR KLELSDDDIK LFKKLREQDE

                                          561
lcl|NC_009127.1_cds_YP_001096113.1_77   LSRREKERLA LKEELKKEMM AEFSSSVPQA AAPVVVPSTS

                                          601
lcl|NC_009127.1_cds_YP_001096113.1_77   APAPVTDVKQ LVSEAIKELL AVQQQQQQQA VVPAPQGVPG

                                          641
lcl|NC_009127.1_cds_YP_001096113.1_77   PATTISLPVA TLAAATGRSP AAAAALQSAI NNVHESASQL

                                          681
lcl|NC_009127.1_cds_YP_001096113.1_77   VPGGLNLGVA GVPVNASTVV STPQPAVPST STQVVSVSAG

                                          721
lcl|NC_009127.1_cds_YP_001096113.1_77   LEVQGGGNKK SLNDDARKMM MLMMDH*
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