10   20   30   40   50   60   70   80   90   100   110   120
GCCCAGG C AT G A G GC T GG T CCCT G AT G G T CATTGCC TCTTGTAGGGAGCTCCA GCCTGACTCGCGGCCTTG GATGA GGCATG AACTGAGGGGAAAACC CAAATTGCT GCCCCCCAATTTTT G

130   140   150   160   170   180   190   200   210   220   230   240
GGGC GGAGGAT CG CCCTC GGG TTT GTGTACCT GTAT CCA GGG ATCTCT GGCACATATACT G AACCGCCACT AAGGGGAACTTTACTT GAAGAC GGCTGCAATCCAACAGGT GATTGCT GGGGAC T

250   260   270
TT GCACTT GTCCATCCCCCACC CCTCTGTAAA