macrogen

10   20   30   40   50   60   70   80   90   100   110   120
C CGG AAG CAGA A AAA A GCT GT G ATAT AA GCAAT AGT T GC TAAAGTT AAAAG AGGG AACAT T AGG GATA CCAAAG CACC CAAT GT AGAG ACG ATT T GTT GTG G GATA GCACG AAT CCT ATG ACAT

130   140   150   160   170   180   190   200   210   220   230   240
TAC CAAG TAAG TT GAGT TGAT AGTAAG AT GT GTT CTT CATT G GTAGTAAT CAAT CAC CTG CAAAAG ATT GT GAT GAT AATTTT GGCCAG ATT CTT CT GTT GGG AGG AGGG CACATG TGCAG

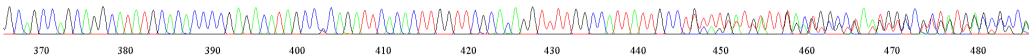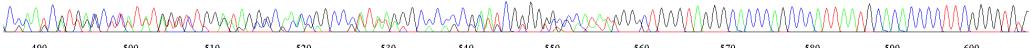

250   260   270   280   290   300   310   320   330   340   350   360
GCAGC CAGC AGG CATAT GCACACC CCG AG ACAT AGC CGC AG ATT CACACT TGGTT CGCACAGG TTTT GGCCATT ATT GCCT GTTT GAT GATT GACTG GCC TAAT GGGCG CTATG AGCGCCG

370   380   390   400   410   420   430   440   450   460   470   480
CC AAT CGT GAGC CTTT ATA GGCT GGA TAGGGC AACCG GCC CATAAGG GCCCCC G AAT CGGGGGG G CATAT GGGGCT GGGT AGGG CACCCAGC CCATT AAT GCACACCCCTT CGGGGTGT G

490   500   510   520   530   540   550   560   570   580   590   600
C ATAT GGG TGC C GC TT GCGCG CA CCTC CT A AT CG TGTCCCT CT CTCTCTGCGCTT C ACCCA GGGG G CG GG TCAT G GCG CCT ACA AAT CGT TACC A GTTT CTAAT GCA CGACGCCC AAGGG

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         610          620          630          640          650          660          670          680          690          700          710          720          730
GG GGTGT ATATC GGGTT TTT ATC GTGGAT GT CTT TATTTT CT CTT T CTTTT TTA G TGG ACTT TGT TTCT TT ACT AT TAAT TCT T TCGTG TAAAACT T TTA TGT TATTTT ATC TGT TT GTAAT TG T

         740          750          760          770          780          790          800          810          820          830          840          850          860
TT TT TTTT AT TATCTAAAAT TATGT AGTG A TTGAATTA C GTA CA AACAT C TGGT GGGA TGGTG GTTA TA TGGA TTCCGCGTT ACTTTA TC CT AATTA GTGGTAAT GCT CT CTTGTTT GGT TGTAA GTATG

         870          880          890          900          910          920          930          940          950          960          970          980          990
GA GTT AA GTA CTTAGC CTCCAAATTTAC CTGAC GTTA CCGCGT AAGCTCATCTTT GTTTTTTTCC TGTT GGCTGAGATATA AAGTT GGGAGCTGAAGTTTTGAAGGAAA AACAC TAACTTCTCCC TATA

        1000         1010         1020         1030         1040         1050         1060         1070         1080         1090         1100         1110         1120
T ATATTTTT GATGGAATTTGGTT TTGTGGGGGT CTCCTTCAAATATAAGGGCACC AATAA AT AC TCTCCTCCCT TTTGTGGGG ATGGAAAA GGGGT GT GTGGTT CTAAATA AAAAAAAA GGGGAG AGG GT GGGC

        1130         1140         1150         1160         1170         1180         1190         1200         1210         1220         1230         1240         1250         1260
CCT TC CGT TA GT TGT GC CCCC CTTT TCCCCC G GGAGCA TTAAAAAATCAACCAATAA TGTGT G AAACT GATTTTTCCC TCTTGGTCGCAAAGTAACCAGGATGCCC CCCCT CATTT GTGGTTT CTAA TCCCCC CAC ACA
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