10   20   30   40   50   60   70   80   90   100   110   120
GGG GGA TAG AT TA ACG C GA GCACAT T T GCCAAG AAT GT CT T CAT T AAT CAAGAAC G AAAGTCAG AG G T T CGAAG GCG AT TAG AT ACC GC CAT AGT TCT G ACCTT AAACAAT ACCT GCT G GCG AA
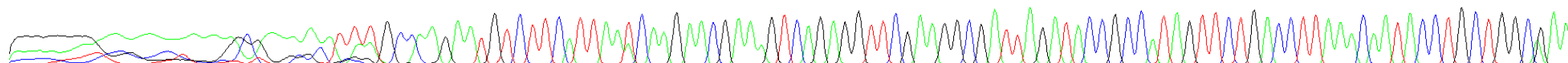
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CC G AT G GT G GCG AAAGCT GCCT T T T C GGGG G AGCACCCC GG AAAC G AAAGC AT AT C GGAT CC GGGGG T AGTAT G GT T GCAAAGCT GAAACT T AAA GGAAT T GACG GAAG AGCACCACC AGG AGT G G
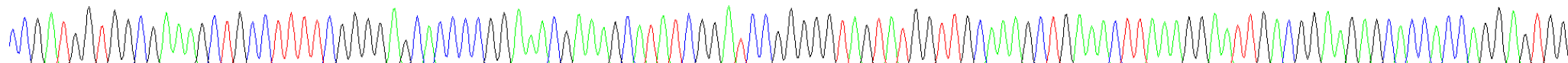
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AGCCT GCGGCT T AAT T T G ACT CAACACGGG AAACCT CACCC GGCCC GGACACC GT AAG GAT T GT CAGACT G AT T GCT CT T AT AT GAT T CGGT G GT T GGT G GT GCAT GGCC GT T CT T AGT T G GT GAA
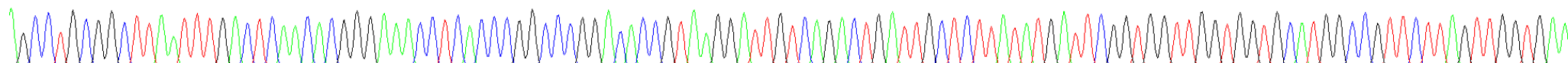
380   390   400   410   420   430   440   450   460   470   480   490   500
GC GAT T T GT CA GGT T T AT T CC GGT AAC GAGC GAGACT CT AGGCT CT AAT T GAAT AC GGG AAT CT T T GAAT T T CGGT ACT T T CT T AAAGGAAT T GGGGG T AAAAAAC GGCAT G CAAAAAC CAAT AA
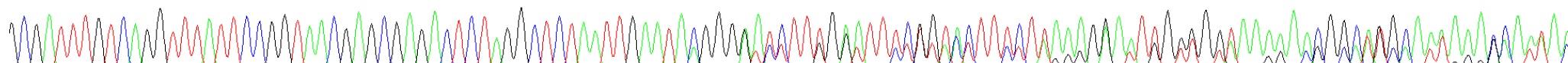
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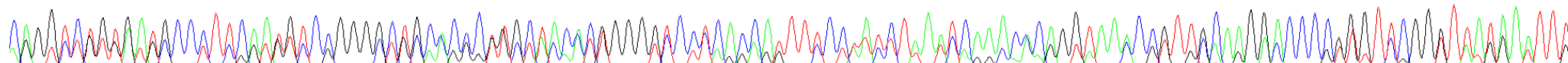
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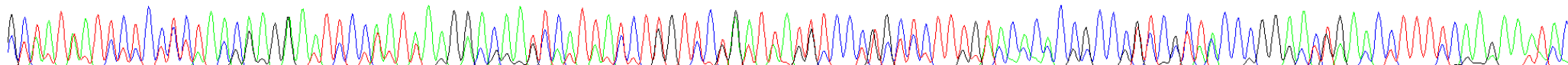
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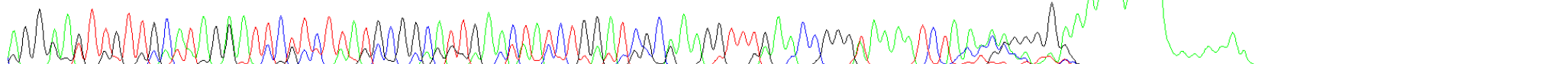
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