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GGGCCCAACT GCATTGAGCA CAACGAGGCC CTCTTGGAGG AGGCGAAGAT 197
GATGAACAGA CTGAGACACA GCCGGGTGGT GAAGCTCCTG GGCGTCATCA 247
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CTGATGCACG TGCTGAAAGC CGAGATGAGT ACTCCGCTTT CTGTAAAAGG 347
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CAAACTGAAT AATGAAGAGC ACAATGAGCT GAGGGAAGTG GACGGCACCG 547  G181S
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GACGTCAACG CAAAGCCCAC AGAGAAGTCG GATGTGTACA GCTTTGCTGT 647  Y212*
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