DNA Sequence Prediction Report

Formatted DNA Sequence	ттессетете
Predicted Mutation Type	Deletion

ID: 10

Sequence: TTGCCGTGTGCTGGTGACGCAATTACAATCAAGAAAGCGCTACGAGAGGAGCGGTGTGTCAGGTAACTATAGTTAAGTAACGTCGCAGTGTGTCG

AAAGTCGTCT

Length: 110

GC_Content: 47.27

Mutation_Type: nan

Gene: Gene_110

Ancestral_Region: Africa

Likelihood_of_Mutation: 0.59

Genetic_Condition: Huntington's Disease

Protein_Synthesis_Effect: No

Genealogy_Markers: ECJMTIL7

Inheritance_Type: Autosomal Recessive

Phenotypic_Expression: Normal

Pathogenic_Score: 0.57

Ethnic Association: Africa

Evolutionary_Clade: Reptile

Epigenetic_Modifications: nan

Health is not valued till sickness comes.