

NC_087265.1:12645345..12645411 Pogoniulus pusillus isolate bPogPus1 chromosome 2, bPogPus1.pri, whole genome shotgun sequence

12,645,350 12,645,360 12,645,370 12,645,380 12,645,390 12,645,400

C T G T T T G T A T G C T C T C C A G C G T G G T T C T T A G C T T C A T C C C T T C C C T C C T T G C T T T T T C T T C C C

G A C A A A C A T A C G A G G T C G C A C C A A G A A T C G A A G T A G G G A A A G G G A G G A A C G A A A A A A A G A A G G G

(U) Alignment for group templates, BLAST Results for: XM_064166482.1:1-1533 PREDICTED: Pogoniulus...

Query_6960787

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Query_6960787

Genes, NCBI RefSeq Annotation GCF_015220805.1-RS_2024_03

SCTR

XM_064166482.1

XP_064022552.1

GAC Q AAA K TAC H GAG E AGG G TCG A CAC H CAA N GAA K TCG A AAG E TAG D GGA R AAG E GGA R GGA R ACG A AAA K AAA K AGA R AGG G

RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF_037610285.1-RS_2024_03 - log 2 scaled



RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF_037610285.1-RS_2024_03 - log 2 scaled



RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF_037610285.1-RS_2024_03