

NC_134317.1:11020924..11021056 Calonectris borealis chromosome 6, bCalBor7.hap1.2, whole genome shotgun sequence

11,020,930 | 11,020,940 | 11,020,950 | 11,020,960 | 11,020,970 | 11,020,980 | 11,020,990 | 11,021 K | 11,021,010 | 11,021,020 | 11,021,030 | 11,021,040 | 11,021,050
AGTCAACCACAGGCTGTTCCATCCCTGCACCAAGGCCACTGGGTGTCGTGCTCCAGCGTGGCTTAGCTCCTTCACTCACCCCTTCCCTCACTTTTCCCTCCCAGCTT
TCAGTTGGTGTCCGACAAAGGAAGTAGGAAACGCGTGGTCTCCGGTACCCGACAGACACGAGAGGTGCAACGAGAATCGAGGAAAGTGAAGTGGGGAAAGGGAGGAGTGAAAAAAGGAAAGGGTCGAA
(U) Alignment for group templates, BLAST Results for: XM_075153922.1 PREDICTED: Calonectris borealis...

Query_4085757
(U) BLAST Results for: XM_075153922.1 PREDICTED: Calonectris borealis...

Genes, NCBI RefSeq Annotation GCF_964195595.1-RS_2025_07
SCTR
- XM_075153922.1
- XP_075010023.1
TCA GTT GGT GTC CGA CAA AGG AAG TAG GGA ACG CGT GGT CTC CGG TGA CCC GAC AGA CAC
T L W L S N G E D R A C W L G S P Q R H
RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF_964195595.1-RS_2025_07 - log 2 scaled

0 0 0 0
RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF_964195595.1-RS_2025_07 - log 2 scaled

10 10 10 10
4 4 4 4
RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF_964195595.1-RS_2025_07