

NC_132943.1:11258037..11258168 Ciconia boyciana chromosome 10, ASM3463844v1, whole genome shotgun sequence

11,258,040 | 11,258,050 | 11,258,060 | 11,258,070 | 11,258,080 | 11,258,090 | 11,258,100 | 11,258,110 | 11,258,120 | 11,258,130 | 11,258,140 | 11,258,150 | 11,258,160

AGTCAATCACAGGCTGTTCCTCATCCCTCGCGCACCAAGGGTACCCAGGCTGTCTGTATGCTCTCCAGTGCTGGCTCTTAGCTCCCTTCACTTCATCCCTTCCCTCGCTTTTCCCTGGCT

TCAGTTAGTGTCCGACAAAGGAAGTAGGGAGCGCGTGGTCTCCAGTGGTCCGACAGACATACGAGAGGTCACACCGAGAATCGAGGAAAGTGAAGTAGGGAAAGGGAGGAGCGAAAAAAAAGGAAGGACCGA

(U) Alignment for group templates, BLAST Results for: XM_072874453.1 PREDICTED: Ciconia boyciana...

Query_8336811

(U) BLAST Results for: XM_072874453.1 PREDICTED: Ciconia boyciana...

Query_8336811

Genes, NCBI RefSeq Annotation GCF_034638445.1-RS_2025_04

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< X < < < < < < < < < < XP_072730554.1 < < < < < < < < < < < < < < < < < < <

TCA GTT AGT GTC CGA CAA AGG AAG TAG GGA GCG CGT GGT CTC CAG TGG TCC GAC AGA TAC GAG AGG TCA CAC CGA GAA TCG AGG AAA GTG AAG TAG GGA AAG GGA GGA GCG AAA KAK AAA KAK GGA AGG ACC GAK

L X L S N G E D R A C W L D G P Q R H E G T H S K A G K V E D R E R A K K R G P K

Genes, Ensembl Rapid release 2024-10

RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF_034638445.1-RS_2025_04 - log 2 scaled

2 2 2 2 2

RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF_034638445.1-RS_2025_04 - log 2 scaled

0 0 0 0 0

RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF_034638445.1-RS_2025_04