

NC_079295.1:33770293..33770355 Falco biarmicus isolate bFalBia1 chromosome 8, bFalBia1.pri, whole genome shotgun sequence

33 770 300

|33 770 310

33 770 320

33 770 330

b3 770 340

33 770 35

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

(U) Alignment for group templates, BLAST Results for: XM_056349686.1:2215-3849 PREDICTED: Falco...
Query_310767

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Genes, NCBI RefSeq Annotation GCF_023638135.1-RS_2023_05

SCTR
XM_056349686.1
XP_056205661.1

GAA E AGG R GAC D GAA E GTG V AAA K GGA C GCT A AGG R AAC N CAA Q ACT T GGA C GAG E CAT H AGA P CAG Q ACC T AGT S GAC D

RNA-seq exon coverage, aggregate (filtered), NCBI RefSeq Annotation GCF_030062545.1-RS_2023_05 - log 2 scaled

RNA-seq intron-spanning reads, aggregate (filtered), NCBI RefSeq Annotation GCF_030062545.1-RS_2023_05 - log 2 scaled

RNA-seq intron features, aggregate (filtered), NCBI RefSeq Annotation GCF_030062545.1-RS_2023_05