

Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

Primate Orthologous Exon Database

RRID:SCR_002065

Type: Tool

Proper Citation

Primate Orthologous Exon Database (RRID:SCR_002065)

Resource Information

URL: <http://giladlab.uchicago.edu/orthoExon/>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022. Database of orthologous exon regions in the genomes of human, chimpanzee, and rhesus macaque. It can be used in analysis of multi-species RNA-seq expression data, allowing for comparisons of exon-level expression across primates, as well as comparative examination of alternative splicing and transcript isoforms.

Abbreviations: Primate Orthologous Exon Database

Resource Type: database, data or information resource

Keywords: alternative splicing, transcript isoform, ortholog, exon, gene, rna-seq, primate, genome

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Primate Orthologous Exon Database

Resource ID: SCR_002065

Alternate IDs: OMICS_01895

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260725T191122+0000

Ratings and Alerts

No rating or validation information has been found for Primate Orthologous Exon Database.

No alerts have been found for Primate Orthologous Exon Database.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Zhu Y, et al. (2014) XSAnno: a framework for building ortholog models in cross-species transcriptome comparisons. BMC genomics, 15(1), 343.