

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

Nonhuman Primate Reference Transcriptome Resource

RRID:SCR_017534

Type: Tool

Proper Citation

Nonhuman Primate Reference Transcriptome Resource (RRID:SCR_017534)

Resource Information

URL: <http://nhprtr.org/>

Proper Citation: Nonhuman Primate Reference Transcriptome Resource (RRID:SCR_017534)

Description: Nonhuman Primate reference transcriptome resource consisting of deep sequencing complete transcriptomes (RNA-seq) from multiple NHP species.

Abbreviations: NHPRTR

Synonyms: Nonhuman Primate Reference Transcriptome Resource

Resource Type: data or information resource, portal, project portal

Keywords: Nonhuman, primate, reference, transcriptome, deep, sequencing, RNAseq, data, species

Funding: NIH

Availability: Free, Freely available

Resource Name: Nonhuman Primate Reference Transcriptome Resource

Resource ID: SCR_017534

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260731T043011+0000

Ratings and Alerts

No rating or validation information has been found for Nonhuman Primate Reference Transcriptome Resource.

No alerts have been found for Nonhuman Primate Reference Transcriptome Resource.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Merchant W, et al. (2025) The molecular components of the anti-inflammatory cholinergic pathway are extrasplenic. PloS one, 20(9), e0331707.

Yee SW, et al. (2024) Illuminating the function of the orphan transporter, SLC22A10, in humans and other primates. Nature communications, 15(1), 4380.

Yee SW, et al. (2023) Illuminating the Function of the Orphan Transporter, SLC22A10 in Humans and Other Primates. Research square.

Yee SW, et al. (2023) Illuminating the Function of the Orphan Transporter, SLC22A10 in Humans and Other Primates. bioRxiv : the preprint server for biology.

Rogers J, et al. (2022) Genomic resources for rhesus macaques (*Macaca mulatta*). Mammalian genome : official journal of the International Mammalian Genome Society, 33(1), 91.

Ramos EI, et al. (2021) Characterization of the Testis-specific LINC01016 Gene Reveals Isoform-specific Roles in Controlling Biological Processes. Journal of the Endocrine Society, 5(11), bvab153.

Bhattacharya A, et al. (2021) Multiple Alu Exonization in 3'UTR of a Primate-Specific Isoform of CYP20A1 Creates a Potential miRNA Sponge. Genome biology and evolution, 13(1).

Santos-Rodriguez G, et al. (2021) Evolutionary dynamics of circular RNAs in primates. eLife, 10.

Chondrou V, et al. (2020) LRF/ZBTB7A conservation accentuates its potential as a therapeutic target for the hematopoietic disorders. Gene, 760, 145020.

Peng X, et al. (2014) Assessment and improvement of Indian-origin rhesus macaque and Mauritian-origin cynomolgus macaque genome annotations using deep transcriptome sequencing data. Journal of medical primatology, 43(5), 317.