

# Resource Summary Report

Generated by [RRID](#) on Aug 11, 2026

## rGREAT

RRID:SCR\_028008

Type: Tool

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### Proper Citation

rGREAT (RRID:SCR\_028008)

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### Resource Information

**URL:** <https://jokergoo.github.io/rGREAT/>

**Proper Citation:** rGREAT (RRID:SCR\_028008)

**Description:** Software R package for functional enrichment on genomic regions.  
Functional enrichment analysis directly performed on genomic regions.

**Synonyms:** Genomic Regions Enrichment of Annotations Tool

**Resource Type:** software toolkit, software resource

**Defining Citation:** [PMID:36394265](#)

**Keywords:** Functional enrichment analysis, functional enrichment on genomic regions,  
genomic regions,

**Availability:** Free, Available for download, Freely available

**Resource Name:** rGREAT

**Resource ID:** SCR\_028008

**License:** MIT license

**Record Creation Time:** 20260224T055257+0000

**Record Last Update:** 20260810T040950+0000

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### Ratings and Alerts

No rating or validation information has been found for rGREAT.

No alerts have been found for rGREAT.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Comerford M, et al. (2026) Mapping the gene regulatory landscape of archaic hominin introgression in modern Papuans. PLoS genetics, 22(3), e1012067.

Hussain S, et al. (2023) Short tandem repeats are important contributors to silencer elements in T cells. Nucleic acids research, 51(10), 4845.

Cieslak A, et al. (2020) Blueprint of human thymopoiesis reveals molecular mechanisms of stage-specific TCR enhancer activation. The Journal of experimental medicine, 217(9).