

# Resource Summary Report

Generated by T1D on Aug 1, 2026

## TRIAGE

RRID:SCR\_016609

Type: Tool

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### Proper Citation

TRIAGE (RRID:SCR\_016609)

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### Resource Information

**URL:** <https://triage.niaid.nih.gov>

**Proper Citation:** TRIAGE (RRID:SCR\_016609)

**Description:** Platform to facilitate prediction, analysis, and hypothesis generation from genome wide perturbation studies like those designed with RNAi and CRISPR technologies.

**Abbreviations:** TRIAGE

**Synonyms:** Throughput Ranking by Iterative Analysis of Genomic Enrichment

**Resource Type:** data analysis service, web application, service resource, production service resource, software resource, analysis service resource

**Keywords:** prediction, analysis, hypothesis, generation, genome, perturbation, study, RNAi, CRISP

**Funding:** NIH

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

**Resource Name:** TRIAGE

**Resource ID:** SCR\_016609

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260801T042800+0000

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### Ratings and Alerts

No rating or validation information has been found for TRIAGE.

No alerts have been found for TRIAGE.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

Zhao Q, et al. (2025) TRIAGE: an R package for regulatory gene analysis. Briefings in bioinformatics, 26(1).