## **Resource Summary Report**

Generated by NIF on Apr 28, 2025

# targetscan.Hs.eg.db

RRID:SCR\_001068

Type: Tool

### **Proper Citation**

targetscan.Hs.eg.db (RRID:SCR\_001068)

### Resource Information

#### **URL:**

http://www.bioconductor.org/packages/release/data/annotation/html/targetscan.Hs.eg.db.html

**Proper Citation:** targetscan.Hs.eg.db (RRID:SCR\_001068)

**Description:** R software that predicts biological targets of miRNAs by searching for the presence of conserved 8mer and 7mer sites that match the seed region of each miRNA.

**Resource Type:** data processing software, sequence analysis software, software resource, data analysis software, software application

Keywords: software, biological targets, mirna, prediction, r, sequence analysis software

**Funding:** 

Availability: Open source

Resource Name: targetscan.Hs.eg.db

Resource ID: SCR 001068

Alternate IDs: OMICS\_00790

**Record Creation Time:** 20220129T080205+0000

Record Last Update: 20250428T052848+0000

### **Ratings and Alerts**

No rating or validation information has been found for targetscan. Hs. eg.db.

No alerts have been found for targetscan. Hs. eg.db.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We have not found any literature mentions for this resource.