

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## DT

RRID:SCR\_024270

Type: Tool

### Proper Citation

DT (RRID:SCR\_024270)

### Resource Information

**URL:** <https://cran.r-project.org/web/packages/DT/index.html>

**Proper Citation:** DT (RRID:SCR\_024270)

**Description:** Software R package for data objects in R to render as HTML tables using JavaScript library DataTables. DataTables library has been included in this R package.

**Synonyms:** DT: A Wrapper of the JavaScript Library 'DataTables', r-cran-dt, DataTables

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

**Keywords:** data objects in R, HTML tables, JavaScript library DataTables,

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

**Resource Name:** DT

**Resource ID:** SCR\_024270

**Alternate URLs:** <https://sources.debian.org/src/r-cran-dt/>

**License:** GPL-3

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260905T133106+0000

### Ratings and Alerts

No rating or validation information has been found for DT.

No alerts have been found for DT.

### Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Branco MA, et al. (2026) Modeling embryonic heart vascular plexus development and sympathetic innervation on a human heart organoid. iScience, 29(3), 115071.

Cacheiro P, et al. (2024) Computational identification of disease models through cross-species phenotype comparison. Disease models & mechanisms, 17(6).

Zou J, et al. (2024) iProPhos: A Web-Based Interactive Platform for Integrated Proteome and Phosphoproteome Analysis. Molecular & cellular proteomics : MCP, 23(1), 100693.

Hacking JPR, et al. (2023) Deriving a continuum score for group 3 and 4 medulloblastoma tumor samples analyzed via RNA-sequencing or DNA methylation microarray. STAR protocols, 4(3), 102509.