

# Resource Summary Report

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

## tsne

RRID:SCR\_024305

Type: Tool

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### Proper Citation

tsne (RRID:SCR\_024305)

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### Resource Information

**URL:** <https://cran.r-project.org/package=tsne>

**Proper Citation:** tsne (RRID:SCR\_024305)

**Description:** Software R package as implementation of the t-SNE algorithm.

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

**Keywords:** t-SNE algorithm implementation, R

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

**Resource Name:** tsne

**Resource ID:** SCR\_024305

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

**License:** GPL-2 | GPL-3 [expanded from: GPL]

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

**Record Last Update:** 20260905T133107+0000

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### Ratings and Alerts

No rating or validation information has been found for tsne.

No alerts have been found for tsne.

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### Data and Source Information

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

## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Renatino Canevarolo R, et al. (2025) Epigenetic Plasticity Drives Carcinogenesis and Multi-Therapy Resistance in Multiple Myeloma. Research square.

Tasis A, et al. (2024) Single-Cell Analysis of Bone Marrow CD8+ T Cells in Myeloid Neoplasms Reveals Pathways Associated with Disease Progression and Response to Treatment with Azacitidine. Cancer research communications, 4(12), 3067.

Yang L, et al. (2024) Dietary Folate and Cofactors Accelerate Age-dependent p16 Epimutation to Promote Intestinal Tumorigenesis. Cancer research communications, 4(1), 164.

Wu ZX, et al. (2024) CD69+CD103+CD8+ tissue-resident memory T cells possess stronger anti-tumor activity and predict better prognosis in colorectal cancer. Cell communication and signaling : CCS, 22(1), 608.

Fang S, et al. (2024) The role of the S100A8/S100A9 in gastric tumor progression. Scientific reports, 14(1), 23574.

Sudalagunta PR, et al. (2024) The Functional Transcriptomic Landscape Informs Therapeutic Strategies in Multiple Myeloma. Cancer research.