

Resource Summary Report

Generated by RRID on Aug 13, 2026

Numbat

RRID:SCR_019207

Type: Tool

Proper Citation

Numbat (RRID:SCR_019207)

Resource Information

URL: <https://github.com/bgcarlisle/Numbat>

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Description: Software tool for managing extraction of large volumes of data from primary sources among multiple users, and then reconciling differences between them. Designed for use in systematic review projects in academic context.

Synonyms: Numbat Systematic Review Manager

Resource Type: data management software, software resource, software application

Keywords: Manage large references databases, large data, manage different extraction levels, multiple extraction forms, codebooks, reference network generation, publication database, systematic review, academic context

Availability: Free, Available for download, Freely available

Resource Name: Numbat

Resource ID: SCR_019207

Alternate URLs: <https://github.com/bgcarlisle/Numbat>, <https://numbat.bgcarlisle.com>

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260812T044404+0000

Ratings and Alerts

No rating or validation information has been found for Numbat.

No alerts have been found for Numbat.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Feng C, et al. (2026) Spatial tumor evolution panorama of ovarian cancer. Cell reports. Medicine, 7(3), 102666.

Kim Y, et al. (2026) Single-nucleus multi-omics delineates distinct epigenetic programs associated with tumor progression in lung adenocarcinoma. Clinical epigenetics, 18(1).

Shi H, et al. (2026) Single-cell profiling reveals distinct populations of tumor-associated macrophages and metastatic tumor cells in breast cancer brain metastasis. Cell death & disease, 17(1).

Collot R, et al. (2026) IGSF11-VISTA is a critical and targetable immune checkpoint axis in diffuse midline glioma. Cancer cell, 44(3), 641.

Ruchiy Y, et al. (2024) Genomic tumor evolution dictates human medulloblastoma progression. Neuro-oncology advances, 6(1), vdae172.

Iarkaeva A, et al. (2024) Workflow for detecting biomedical articles with underlying open and restricted-access datasets. PloS one, 19(5), e0302787.

Iskander R, et al. (2024) The Benefits and Risks of Receiving Investigational Solid Tumor Drugs in Randomized Trials : A Systematic Review and Meta-analysis. Annals of internal medicine, 177(6), 759.

Hutchinson N, et al. (2024) Probability of Regulatory Approval Over Time: A Cohort Study of Cancer Therapies. JCO oncology practice, 20(2), 247.

Akay EMZ, et al. (2023) Artificial Intelligence for Clinical Decision Support in Acute Ischemic Stroke: A Systematic Review. Stroke, 54(6), 1505.

Bicer S, et al. (2023) Timing for First-in-Minor Clinical Trials of New Cancer Drugs. The Journal of pediatrics, 263, 113705.

Hutchinson N, et al. (2022) How informative were early SARS-CoV-2 treatment and prevention trials? a longitudinal cohort analysis of trials registered on ClinicalTrials.gov. PloS one, 17(1), e0262114.