

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

battenberg

RRID:SCR_017098

Type: Tool

Proper Citation

battenberg (RRID:SCR_017098)

Resource Information

URL: <https://github.com/Wedge-Oxford/battenberg>

Proper Citation: battenberg (RRID:SCR_017098)

Description: Software R package for subclonal copy number estimation.

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

Keywords: subclonal, copy, number, estimation

Availability: Free, Available for download, Freely available

Resource Name: battenberg

Resource ID: SCR_017098

License: GNU AGPL v3

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260729T044431+0000

Ratings and Alerts

No rating or validation information has been found for battenberg.

No alerts have been found for battenberg.

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 [nidm-terms](#) .

Woolley CE, et al. (2025) Coevolution of Atypical BRAF and KRAS Mutations in Colorectal Tumorigenesis. Molecular cancer research : MCR, 23(4), 300.

Treger TD, et al. (2025) Predisposition Footprints in the Somatic Genome of Wilms Tumors. Cancer discovery, 15(2), 286.

M Naeini M, et al. (2023) Multi-omic features of oesophageal adenocarcinoma in patients treated with preoperative neoadjuvant therapy. Nature communications, 14(1), 3155.

Buhigas C, et al. (2022) The architecture of clonal expansions in morphologically normal tissue from cancerous and non-cancerous prostates. Molecular cancer, 21(1), 183.

Liu LY, et al. (2020) Quantifying the influence of mutation detection on tumour subclonal reconstruction. Nature communications, 11(1), 6247.

Lee JJ, et al. (2019) Tracing Oncogene Rearrangements in the Mutational History of Lung Adenocarcinoma. Cell, 177(7), 1842.