

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 28, 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: data analysis software, data processing software, software application, software resource

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: 20260821T194116+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 7 mentions in open access literature.

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

Culliford R, et al. (2026) Contrasting Features of Papillary and Chromophobe Renal Cell Carcinoma Revealed by Whole-Genome Sequencing. *Molecular cancer research : MCR*, 24(5), 327.

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.