

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

[cgpBattenberg](#)

RRID:SCR_017092

Type: Tool

Proper Citation

cgpBattenberg (RRID:SCR_017092)

Resource Information

URL: <https://github.com/cancerit/cgpBattenberg>

Proper Citation: cgpBattenberg (RRID:SCR_017092)

Description: Software tool as installation helper, perl wrapper and R program Battenberg which detects subclonality and copy number in matched NGS data.

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

Keywords: installation, helper, perl, wrapper, detect, subclonality, copy, number, NGS, next, generation, sequencing, data

Availability: Free, Available for download, Freely available

Resource Name: cgpBattenberg

Resource ID: SCR_017092

License: GNU AGPL v3

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260805T044405+0000

Ratings and Alerts

No rating or validation information has been found for cgpBattenberg.

No alerts have been found for cgpBattenberg.

Data and Source Information

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Domenico D, et al. (2025) Enabling whole genome sequencing analysis from FFPE specimens in clinical oncology. Nature communications, 16(1), 10649.

Pacyna CN, et al. (2024) Multifocal, multiphenotypic tumours arising from an MTOR mutation acquired in early embryogenesis. Oncogene, 43(44), 3268.

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

Pilet J, et al. (2023) Preneoplastic liver colonization by 11p15.5 altered mosaic cells in young children with hepatoblastoma. Nature communications, 14(1), 7122.

Andersen LVB, et al. (2023) Non-BRCA1/BRCA2 high-risk familial breast cancers are not associated with a high prevalence of BRCAness. Breast cancer research : BCR, 25(1), 69.

Ketola K, et al. (2021) Subclone Eradication Analysis Identifies Targets for Enhanced Cancer Therapy and Reveals L1 Retrotransposition as a Dynamic Source of Cancer Heterogeneity. Cancer research, 81(19), 4901.

Woodcock DJ, et al. (2020) Prostate cancer evolution from multilineage primary to single lineage metastases with implications for liquid biopsy. Nature communications, 11(1), 5070.

Glodzik D, et al. (2020) Comprehensive molecular comparison of BRCA1 hypermethylated and BRCA1 mutated triple negative breast cancers. Nature communications, 11(1), 3747.

Maciejowski J, et al. (2020) APOBEC3-dependent kataegis and TREX1-driven chromothripsis during telomere crisis. Nature genetics, 52(9), 884.

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

Steele CD, et al. (2019) Undifferentiated Sarcomas Develop through Distinct Evolutionary Pathways. Cancer cell, 35(3), 441.