

Resource Summary Report

Generated by [NIF](#) on Sep 4, 2026

FlowSorted.Blood.450k R package

RRID:SCR_018003

Type: Tool

Proper Citation

FlowSorted.Blood.450k R package (RRID:SCR_018003)

Resource Information

URL: <https://bioconductor.org/packages/FlowSorted.Blood.450k/>

Proper Citation: FlowSorted.Blood.450k R package (RRID:SCR_018003)

Description: Illumina HumanMethylation data on sorted blood cell populations.

Resource Type: data or information resource, data set

Defining Citation: [PMID:22848472](#)

Keywords: Illumina, human methylation, DNA, data, sorted blood cell,

Availability: Free, Available to download, Freely available

Resource Name: FlowSorted.Blood.450k R package

Resource ID: SCR_018003

Record Creation Time: 20220129T080338+0000

Record Last Update: 20260904T000824+0000

Ratings and Alerts

No rating or validation information has been found for FlowSorted.Blood.450k R package.

No alerts have been found for FlowSorted.Blood.450k R package.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Malta TM, et al. (2024) The Epigenetic Evolution of Glioma Is Determined by the IDH1 Mutation Status and Treatment Regimen. *Cancer research*, 84(5), 741.

Rovira P, et al. (2020) Epigenome-wide association study of attention-deficit/hyperactivity disorder in adults. *Translational psychiatry*, 10(1), 199.

Chakravarthy A, et al. (2018) Pan-cancer deconvolution of tumour composition using DNA methylation. *Nature communications*, 9(1), 3220.