

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

[wgbstools](#)

RRID:SCR_028257

Type: Tool

Proper Citation

wgbstools (RRID:SCR_028257)

Resource Information

URL: https://github.com/nloyfer/wgbs_tools

Proper Citation: wgbstools (RRID:SCR_028257)

Description: Software tools for working with Bisulfite Sequencing data while preserving reads intrinsic dependencies.

Resource Type: software toolkit, source code, software resource

Keywords: Bisulfite Sequencing data, preserving reads intrinsic dependencies,

Availability: Free, Available for download, Freely available

Resource Name: wgbstools

Resource ID: SCR_028257

License URLs: https://github.com/nloyfer/wgbs_tools?tab=License-1-ov-file#readme

Record Creation Time: 20260407T061137+0000

Record Last Update: 20260815T183310+0000

Ratings and Alerts

No rating or validation information has been found for wgbstools.

No alerts have been found for wgbstools.

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 [RRID](#) .

Murphy L, et al. (2025) Platelets sequester extracellular DNA, capturing tumor-derived and free fetal DNA. Science (New York, N.Y.), 389(6761), eadp3971.

Kueng N, et al. (2023) Investigation of Different Library Preparation and Tissue of Origin Deconvolution Methods for Urine and Plasma cfDNA Methylome Analysis. Diagnostics (Basel, Switzerland), 13(15).

Jaber M, et al. (2022) Comparative parallel multi-omics analysis during the induction of pluripotent and trophectoderm states. Nature communications, 13(1), 3475.