

Resource Summary Report

Generated by [RRID](#) on Aug 14, 2026

OutLyzer

RRID:SCR_025120

Type: Tool

Proper Citation

OutLyzer (RRID:SCR_025120)

Resource Information

URL: <https://github.com/EtieM/outLyzer>

Proper Citation: OutLyzer (RRID:SCR_025120)

Description: Software tool for extracting low-allele-frequency tumor mutations from sequencing background noise in clinical practice. Detects variations, specifically low allele frequency variation, in next generation sequencing data.

Resource Type: software resource, source code

Defining Citation: [PMID:27825131](#)

Keywords: Detects variations, variant caller, low allele frequency variation, extracting low-allele-frequency tumor mutations, next generation sequencing data

Funding: Institut National du Cancer

Availability: Free, Available for download, Freely available

Resource Name: OutLyzer

Resource ID: SCR_025120

Alternate URLs: <https://github.com/EtieM/outLyzer>

Record Creation Time: 20240320T053304+0000

Record Last Update: 20260813T055319+0000

Ratings and Alerts

No rating or validation information has been found for OutLyzer.

No alerts have been found for OutLyzer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

C??saire M, et al. (2019) Sensitization of chondrosarcoma cells with PARP inhibitor and high-LET radiation. Journal of bone oncology, 17, 100246.