

Resource Summary Report

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

XenofilteR

RRID:SCR_026196

Type: Tool

Proper Citation

XenofilteR (RRID:SCR_026196)

Resource Information

URL: <https://github.com/NKI-GCF/XenofilteR>

Proper Citation: XenofilteR (RRID:SCR_026196)

Description: Software tool for computational deconvolution of mouse and human reads in tumor xenograft sequence data.

Resource Type: software application, software resource, source code

Defining Citation: [PMID:30286710](#)

Keywords: computational deconvolution, mouse, human, reads, tumor xenograft sequence data,

Funding: European Research Council under the European Union's Seventh Framework Programme ;
Dutch Cancer Society ;
Cancer Research UK ;
The Wellcome Trust

Availability: Free, Available for download, Freely available

Resource Name: XenofilteR

Resource ID: SCR_026196

License: GNU GPL v3

Record Creation Time: 20241218T053316+0000

Record Last Update: 20260805T044556+0000

Ratings and Alerts

No rating or validation information has been found for XenofilterR.

No alerts have been found for XenofilterR.

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](#) .

He S, et al. (2025) Comprehensive assessment of computational methods for cancer immunoediting. Cell reports methods, 5(3), 101006.

Mal A, et al. (2025) HIF-2-Dependent Regulation of PTHrP and Paraneoplastic Hypercalcemia in Aggressive Clear-Cell Renal Cell Carcinoma. Cancer discovery, 15(12), 2485.

Montaudon E, et al. (2020) PLK1 inhibition exhibits strong anti-tumoral activity in CCND1-driven breast cancer metastases with acquired palbociclib resistance. Nature communications, 11(1), 4053.