

Resource Summary Report

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

Xenome

RRID:SCR_026266

Type: Tool

Proper Citation

Xenome (RRID:SCR_026266)

Resource Information

URL: <https://github.com/data61/gossamer/blob/master/docs/xenome.md>

Proper Citation: Xenome (RRID:SCR_026266)

Description: Software tool to distinguish host and graft reads from next generation sequencing assays of xenograft samples. Performs classification of xenograft-derived sequence read data.

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

Defining Citation: [PMID:22689758](#)

Keywords: distinguish host and graft reads, next generation sequencing assays, xenograft samples, classification of xenograft-derived sequence read data, xenograft-derived sequence read data,

Funding: Prostate Cancer Foundation of Australia ;
Victorian Government Operational Infrastructure Support

Availability: Free, Freely available,

Resource Name: Xenome

Resource ID: SCR_026266

Record Creation Time: 20250110T053324+0000

Record Last Update: 20260815T042913+0000

Ratings and Alerts

No rating or validation information has been found for Xenome.

No alerts have been found for Xenome.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Avolio M, et al. (2026) The Molecular and Functional Landscape of Resistance to FOLFIRI Chemotherapy in Metastatic Colorectal Cancer. Cancer discovery, 16(2), 270.

Neill NJ, et al. (2025) Integrative Proteogenomics and Forward Genetics Reveal a Novel Mitotic Vulnerability in Triple-Negative Breast Cancer. Cancer discovery, 15(11), 2326.