Resource Summary Report

Generated by RRID on Aug 13, 2026

CancerMine

RRID:SCR_027748
Type: Tool

Proper Citation

CancerMine (RRID:SCR_027748)

Resource Information

URL: http://bionlp.bcgsc.ca/cancermine/

Proper Citation: CancerMine (RRID:SCR_027748)

Description: Text-mined and routinely updated database of drivers, oncogenes and tumor suppressors in different types of cancer.

Resource Type: data or information resource, database

Defining Citation: PMID:31110280

Keywords: Cancer genes, literature-mined database, drivers, oncogenes, tumor suppressors, cancer

Funding: BC Cancer Foundation and Genome British Columbia ;
Vanier Canada Graduate Scholarship

Availability: Free, Freely available

Resource Name: CancerMine

Resource ID: SCR_027748

Record Creation Time: 20251209T055453+0000

Record Last Update: 20260811T043828+0000

Ratings and Alerts

No rating or validation information has been found for CancerMine.

No alerts have been found for CancerMine.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Li H, et al. (2026) CASER: A semi-supervised model with multi-omics data integration prioritizes cancer-associated epigenetic regulator genes. PLoS computational biology, 22(4), e1014253.

Shin YJ, et al. (2022) Portrait of Molecular Signaling and Putative Therapeutic Targets in Prostate Cancer with ETV4 Fusion. Biomedicines, 10(10).

Martinez-Fundichely A, et al. (2022) Modeling tissue-specific breakpoint proximity of structural variations from whole-genomes to identify cancer drivers. Nature communications, 13(1), 5640.

Grimaldi AM, et al. (2020) The New Paradigm of Network Medicine to Analyze Breast Cancer Phenotypes. International journal of molecular sciences, 21(18).

Terkelsen T, et al. (2020) Secreted breast tumor interstitial fluid microRNAs and their target genes are associated with triple-negative breast cancer, tumor grade, and immune infiltration. Breast cancer research : BCR, 22(1), 73.

Shademan M, et al. (2019) Expression profile analysis of two antisense lncRNAs to improve prognosis prediction of colorectal adenocarcinoma. Cancer cell international, 19, 278.