

# 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](#)

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## 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.

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.

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

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