

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## Progenetix

RRID:SCR\_019220

Type: Tool

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### Proper Citation

Progenetix (RRID:SCR\_019220)

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### Resource Information

**URL:** <https://progenetix.org/>

**Proper Citation:** Progenetix (RRID:SCR\_019220)

**Description:** Collection of mutation data in cancer, with focus on copy number abnormalities for all types of human malignancies.

**Resource Type:** data or information resource, database

**Keywords:** Cancer genome data, genome profiles, cancer mutation data, copy number abnormalities, human malignancies types

**Availability:** Free, Freely available

**Resource Name:** Progenetix

**Resource ID:** SCR\_019220

**Alternate IDs:** r3d100012820

**Alternate URLs:** <https://progenetix.net>, <https://doi.org/10.17616/R3KV46>

**Record Creation Time:** 20220129T080344+0000

**Record Last Update:** 20260905T133217+0000

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### Ratings and Alerts

No rating or validation information has been found for Progenetix.

No alerts have been found for Progenetix.

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### Data and Source Information

## Usage and Citation Metrics

We found 11 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Yang Z, et al. (2026) CNAttention: an attention-based deep multiple-instance method for uncovering copy number aberration signatures across cancers. *Briefings in bioinformatics*, 27(1).

Zhao H, et al. (2025) CNAdjust: enhancing CNA calling accuracy through systematic baseline adjustment. *Frontiers in genetics*, 16, 1674138.

Liu SC, et al. (2023) A 3-gene signature comprising CDH4, STAT4 and EBV-encoded LMP1 for early diagnosis and predicting disease progression of nasopharyngeal carcinoma. *Discover. Oncology*, 14(1), 119.

Huth T, et al. (2023) Chromosome 8p engineering reveals increased metastatic potential targetable by patient-specific synthetic lethality in liver cancer. *Science advances*, 9(51), eadh1442.

Huang Q, et al. (2022) Candidate targets of copy number deletion events across 17 cancer types. *Frontiers in genetics*, 13, 1017657.

Acheson E, et al. (2021) Extracting and modeling geographic information from scientific articles. *PloS one*, 16(1), e0244918.

Huang Q, et al. (2021) The Progenetix oncogenomic resource in 2021. *Database : the journal of biological databases and curation*, 2021.

Funato K, et al. (2021) Dissecting the impact of regional identity and the oncogenic role of human-specific NOTCH2NL in an hESC model of H3.3G34R-mutant glioma. *Cell stem cell*, 28(5), 894.

Gao B, et al. (2021) Signatures of Discriminative Copy Number Aberrations in 31 Cancer Subtypes. *Frontiers in genetics*, 12, 654887.

, et al. (2021) Voices of GA4GH members: Collaborating in technology and policy development. *Cell genomics*, 1(2), 100036.

Yang Y, et al. (2020) LncRNA PPP1R14B-AS1 Promotes Tumor Cell Proliferation and Migration via the Enhancement of Mitochondrial Respiration. *Frontiers in genetics*, 11, 557614.