Resource Summary Report

Generated by PRECISE-TBI on Sep 9, 2026

cgpPindel

RRID:SCR_017090
Type: Tool

Proper Citation

cgpPindel (RRID:SCR_017090)

Resource Information

URL: https://github.com/cancerit/cgpPindel

Proper Citation: cgpPindel (RRID:SCR_017090)

Description: Software tool as cancer genome project insertion or deletion detection workflow for Pindel.

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

Availability: Free, Available for download, Freely available

Resource Name: cgpPindel

Resource ID: SCR_017090

Alternate URLs: http://cancerit.github.io/cgpPindel/

License: GNU AGPL v3

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260905T133008+0000

Ratings and Alerts

No rating or validation information has been found for cgpPindel.

No alerts have been found for cgpPindel.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 26 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Davies HR, et al. (2026) Homologous recombination deficiency in primary ER-positive and HER2-negative breast cancer. Communications medicine, 6(1), 118.

Leongamornlert D, et al. (2026) Genomic evolution and natural history of myeloproliferative neoplasms on therapy. Cancer discovery.

Wong K, et al. (2025) Wnt/?-catenin activation by mutually exclusive FBXW11 and CTNNB1 hotspot mutations drives salivary basal cell adenoma. Nature communications, 16(1), 4657.

McLoughlin MA, et al. (2025) Telomere attrition becomes an instrument for clonal selection in aging hematopoiesis and leukemogenesis. Nature genetics, 57(9), 2215.

Brzozowska N, et al. (2025) Selection for somatic escape variants in SERPINA1 in the liver of patients with alpha-1 antitrypsin deficiency. Nature genetics, 57(4), 875.

Kapadia CD, et al. (2025) Clonal dynamics and somatic evolution of haematopoiesis in mouse. Nature, 641(8063), 681.

Domenico D, et al. (2025) Enabling whole genome sequencing analysis from FFPE specimens in clinical oncology. Nature communications, 16(1), 10649.

Minello A, et al. (2025) A recurrent pathogenic BRCA2 truncating variant reveals a role for BRCA2-PCAF complex in modulating NF-?B-driven transcription. Nature communications, 16(1), 10953.

Yamaguchi TN, et al. (2025) The Germline and Somatic Origins of Prostate Cancer Heterogeneity. Cancer discovery, 15(5), 988.

Ferreira I, et al. (2025) The molecular cartography of malignant and benign sebaceous tumours. Nature communications, 17(1), 14.

Kapadia CD, et al. (2024) Clonal dynamics and somatic evolution of haematopoiesis in mouse. bioRxiv : the preprint server for biology.

Keahi DL, et al. (2024) G-quadruplexes are a source of vulnerability in BRCA2 deficient granule cell progenitors and medulloblastoma. bioRxiv : the preprint server for biology.

Pacyna CN, et al. (2024) Multifocal, multiphenotypic tumours arising from an MTOR mutation acquired in early embryogenesis. Oncogene, 43(44), 3268.

Basyuni S, et al. (2024) Large-scale analysis of whole genome sequencing data from formalin-fixed paraffin-embedded cancer specimens demonstrates preservation of clinical utility. Nature communications, 15(1), 7731.

Olafsson S, et al. (2023) Effects of psoriasis and psoralen exposure on the somatic mutation landscape of the skin. Nature genetics, 55(11), 1892.

Hutten SJ, et al. (2023) A living biobank of patient-derived ductal carcinoma in situ mouse-intraductal xenografts identifies risk factors for invasive progression. Cancer cell, 41(5), 986.

Li R, et al. (2022) Mapping single-cell transcriptomes in the intra-tumoral and associated territories of kidney cancer. Cancer cell, 40(12), 1583.

Mitchell E, et al. (2022) Clonal dynamics of haematopoiesis across the human lifespan. Nature, 606(7913), 343.

Petljak M, et al. (2022) Mechanisms of APOBEC3 mutagenesis in human cancer cells. Nature, 607(7920), 799.

Rose Li Y, et al. (2020) Mutational signatures in tumours induced by high and low energy radiation in Trp53 deficient mice. Nature communications, 11(1), 394.