

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

CanPredict

RRID:SCR_008216

Type: Tool

Proper Citation

CanPredict (RRID:SCR_008216)

Resource Information

URL: <http://research-public.gene.com/Research/genentech/canpredict/index.html>

Proper Citation: CanPredict (RRID:SCR_008216)

Description: Web application that uses a combination of computational methods to identify those changes most likely to be cancer-associated.

Abbreviations: CanPredict

Synonyms: CanPredict: A computational tool for predicting Cancer-associated mutations

Resource Type: service resource, production service resource, analysis service resource, data analysis service

Related Condition: Cancer

Resource Name: CanPredict

Resource ID: SCR_008216

Alternate IDs: OMICS_00142

Record Creation Time: 20220129T080246+0000

Record Last Update: 20260731T042752+0000

Ratings and Alerts

No rating or validation information has been found for CanPredict.

No alerts have been found for CanPredict.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Yang S, et al. (2025) Photoreceptor metabolic window unveils eye-body interactions. Nature communications, 16(1), 697.

Tian R, et al. (2015) Computational methods and resources for the interpretation of genomic variants in cancer. BMC genomics, 16 Suppl 8(Suppl 8), S7.

Doss CG, et al. (2014) Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. BioMed research international, 2014, 895831.