

# Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 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:** analysis service resource, data analysis service, production service resource, service resource

**Related Condition:** Cancer

**Resource Name:** CanPredict

**Resource ID:** SCR\_008216

**Alternate IDs:** OMICS\_00142

**Record Creation Time:** 20220129T080246+0000

**Record Last Update:** 20260919T195716+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 [nidm-terms](#) .

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.