## **Resource Summary Report**

Generated by NIF on May 4, 2025

# **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: production service resource, analysis service resource, service resource,

data analysis service

Related Condition: Cancer

**Funding:** 

Resource Name: CanPredict

Resource ID: SCR\_008216

Alternate IDs: OMICS\_00142

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

Record Last Update: 20250503T060006+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 NIF.

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.