Resource Summary Report

Generated by dkNET on Aug 11, 2026

TNBCtype

RRID:SCR_026238
Type: Tool

Proper Citation

TNBCtype (RRID:SCR_026238)

Resource Information

URL: https://cbc.app.vumc.org/tnbc/

Proper Citation: TNBCtype (RRID:SCR_026238)

Description: Website for predicting the subtype of triple negative breast cancer sample based on its gene expression profile.

Resource Type: web application, software resource

Defining Citation: PMID:22872785

Keywords: predicting subtype, triple negative breast cancer sample, gene expression profile,

Funding: NCI CA068485;
NCI CA95131;
NCI CA148375;
NCI CA105436;
NCI CA070856;
NCI CA009385;
American Cancer Society ;
Komen Foundation

Availability: Free, Freely available

Resource Name: TNBCtype

Resource ID: SCR_026238

Record Creation Time: 20250108T053322+0000

Record Last Update: 20260808T190736+0000

Ratings and Alerts

No rating or validation information has been found for TNBCtype.

No alerts have been found for TNBCtype.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Torossian N, et al. (2025) Reference-free RNA profiling predicts triple negative breast cancer chemoresistance to neoadjuvant treatment. NAR cancer, 7(4), zcaf036.

Torres ER, et al. (2025) Sequencing analysis reveals evidence of immune activation in advanced HER2 negative breast cancer responders treated with entinostat + nivolumab + ipilimumab. Research square.

Chen W, et al. (2022) High PANX1 Expression Leads to Neutrophil Recruitment and the Formation of a High Adenosine Immunosuppressive Tumor Microenvironment in Basal-like Breast Cancer. Cancers, 14(14).

Font-Clos F, et al. (2022) Classification of triple negative breast cancer by epithelial mesenchymal transition and the tumor immune microenvironment. Scientific reports, 12(1), 9651.