

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

U3-Nomogram

RRID:SCR_027696

Type: Tool

Proper Citation

U3-Nomogram (RRID:SCR_027696)

Resource Information

URL: <https://github.com/MANO-B/U3-Nomogram>

Proper Citation: U3-Nomogram (RRID:SCR_027696)

Description: Software application for C-CAT CALICO database. Used for Germline-focUsed analysis of tUmoUr-detected variants with Nomogram for C-CAT CALICO database.

Resource Type: source code, software application, software resource

Keywords: C-CAT CALICO database, Germline-focUsed analysis, tUmoUr-detected variants

Availability: Free, Available for download, Freely available

Resource Name: U3-Nomogram

Resource ID: SCR_027696

Record Creation Time: 20251122T055350+0000

Record Last Update: 20260815T183309+0000

Ratings and Alerts

No rating or validation information has been found for U3-Nomogram.

No alerts have been found for U3-Nomogram.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Ikegami M, et al. (2026) Germline Pathogenic Variant Prediction Model for Tumor-Only Sequencing Based on Japanese Clinicogenomic Database. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(4), 770.