Resource Summary Report

Generated by RRID on Aug 15, 2026

MutPanningV2

RRID:SCR_027729
Type: Tool

Proper Citation

MutPanningV2 (RRID:SCR_027729)

Resource Information

URL: https://github.com/vanallenlab/MutPanningV2

Proper Citation: MutPanningV2 (RRID:SCR_027729)

Description: Software tool designed to detect rare cancer driver genes from aggregated whole-exome sequencing data.

Synonyms: MutPanning

Resource Type: source code, software application, software resource

Defining Citation: PMID:32015527

Keywords: Cancer Genes, SNV, indel, detect rare cancer driver genes, aggregated whole-exome sequencing data,

Availability: Free, Available for download, Freely available

Resource Name: MutPanningV2

Resource ID: SCR_027729

License: BSD-3-Clause license

Record Creation Time: 20251209T055452+0000

Record Last Update: 20260815T183309+0000

Ratings and Alerts

No rating or validation information has been found for MutPanningV2.

No alerts have been found for MutPanningV2.

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 .

Newell F, et al. (2022) Comparative Genomics Provides Etiologic and Biological Insight into Melanoma Subtypes. Cancer discovery, 12(12), 2856.