

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[VARiD](#)

RRID:SCR_000241

Type: Tool

Proper Citation

VARiD (RRID:SCR_000241)

Resource Information

URL: <http://compbio.cs.utoronto.ca/varid/>

Proper Citation: VARiD (RRID:SCR_000241)

Description: Software using a Hidden Markov Model for SNP (single nucleotide polymorphism) and indel identification with AB-SOLiD color-space as well as regular letter-space reads.

Abbreviations: VARiD

Resource Type: software resource

Defining Citation: [PMID:20529926](#)

Keywords: c, single nucleotide polymorphism, indel, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: VARiD

Resource ID: SCR_000241

Alternate IDs: OMICS_02163, biotools:varid

Alternate URLs: <https://bio.tools/varid>

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260801T190112+0000

Ratings and Alerts

No rating or validation information has been found for VARiD.

No alerts have been found for VARiD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.