Resource Summary Report

Generated by NIF on Aug 9, 2026

MoDIL

RRID:SCR_010764
Type: Tool

Proper Citation

MoDIL (RRID:SCR_010764)

Resource Information

URL: http://compbio.cs.toronto.edu/modil/

Proper Citation: MoDIL (RRID:SCR_010764)

Description: Software for a novel method for finding medium sized indels from high throughput sequencing datasets.

Abbreviations: MoDIL

Synonyms: MoDIL: Detecting INDEL Variation with Clone-end Sequencing

Resource Type: software resource

Keywords: bio.tools

Resource Name: MoDIL

Resource ID: SCR_010764

Alternate IDs: OMICS_00066, biotools:modil

Alternate URLs: https://bio.tools/modil

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260808T185935+0000

Ratings and Alerts

No rating or validation information has been found for MoDIL.

No alerts have been found for MoDIL.

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 NIF .

Russell CW, et al. (2019) Comprehensive Identification of Fim-Mediated Inversions in Uropathogenic Escherichia coli with Structural Variation Detection Using Relative Entropy. mSphere, 4(2).

Xia LC, et al. (2016) A genome-wide approach for detecting novel insertion-deletion variants of mid-range size. Nucleic acids research, 44(15), e126.

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.

Keane TM, et al. (2014) Identification of structural variation in mouse genomes. Frontiers in genetics, 5, 192.