DINDEL
RRID:SCR_001827
Type: Tool

Proper Citation
DINDEL (RRID:SCR_001827)

Resource Information

URL: http://www.sanger.ac.uk/science/tools/dindel

Description: Software program for calling small indels from short-read sequence data ("next generation sequence data"). It is currently designed to handle only Illumina data. Dindel takes BAM files with mapped Illumina read data and enables researchers to detect small indels and produce a VCF file of all the variant calls. It has been written in C++ and can be used on Linux-based and Mac computers (it has not been tested on Windows operating systems).

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Resource Type: Resource, software resource, software application

Keywords: indel, short-read, next generation sequence, illumina, gene, genetic, genomic, c++, linux, macos

Resource ID: SCR_001827

Parent Organization: Wellcome Trust Sanger Institute; Hinxton; United Kingdom

References: PMID:20980555

Website Status: Last checked up

Alternate IDs: nlx_154283, OMICS_00096

Old URLs: http://www.sanger.ac.uk/resources/software/dindel/
Abbreviations: Dindel

Mentions Count: 41

Ratings and Alerts

No rating or validation information has been found for DINDEL.

No alerts have been found for DINDEL.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 41 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch Infrastructure.


