FreeBayes
RRID:SCR_010761
Type: Tool

Proper Citation
FreeBayes (RRID:SCR_010761)

Resource Information

URL: https://github.com/ekg/freebayes

Description: A Bayesian genetic variant detector designed to find small polymorphisms, specifically SNPs, indels, MNPs, and complex events smaller than the length of a short-read sequencing alignment.

Resource Name: FreeBayes

Proper Citation: FreeBayes (RRID:SCR_010761)

Resource Type: Resource, software resource

Keywords: single-nucleotide polymorphism, indel, insertion, deletion, multi-nucleotide polymorphism, complex event, composite insertion, substitution event

Resource ID: SCR_010761

Availability: Acknowledgement requested

Website Status: Last checked up

Alternate IDs: OMICS_00059

Abbreviations: FreeBayes

Mentions Count: 575
No rating or validation information has been found for FreeBayes.

No alerts have been found for FreeBayes.

**Data and Source Information**

**Source:** SciCrunch Registry

**Usage and Citation Metrics**

We found 575 mentions in open access literature.

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


Wang TT, et al. (2020) Whole Genome Sequencing of Spontaneously Occurring Rat Natural Killer Large Granular Lymphocyte Leukemia Identifies JAK1 Somatic Activating Mutation. Cancers, 12(1).


