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

Ratings and Alerts
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).


