HISAT2
RRID:SCR_015530
Type: Tool

Proper Citation

HISAT2 (RRID:SCR_015530)

Resource Information

URL: http://ccb.jhu.edu/software/hisat2/index.shtml

Description: Graph-based alignment of next generation sequencing reads to a population of genomes.

Resource Name: HISAT2

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Resource Type: Resource, source code, data analysis software, data processing software, software application, sequence analysis software, software resource

Keywords: alignment program, mapping reads, population genomics, human genome

Resource ID: SCR_015530

Parent Organization: Johns Hopkins University; Maryland; USA

Funding Agency: NIGMS, NLM, NSF

Related resources: TopHat

References: PMID:25751142

Availability: Available for download

Website Status: Last checked up

Alternate URLs: https://github.com/infphilo/hisat2

Mentions Count: 1331
Ratings and Alerts

No rating or validation information has been found for HISAT2.

No alerts have been found for HISAT2.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1331 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](#).


Qin G, et al. (2020) Diversity of metabolite accumulation patterns in inner and outer seed coats of pomegranate: exploring their relationship with genetic mechanisms of seed coat


