Picard
RRID:SCR_006525
Type: Tool

Proper Citation

Picard (RRID:SCR_006525)

Resource Information

URL: http://broadinstitute.github.io/picard/

Proper Citation: Picard (RRID:SCR_006525)

Description: Java toolset for working with next generation sequencing data in the BAM format.

Resource Type: Resource, software resource, software toolkit, source code

Keywords: next generation sequencing, java, bam

Parent Organization: Broad Institute, SourceForge

Availability: Available for download, Free

Website Status: Last checked up

Resource Name: Picard

Resource ID: SCR_006525

Alternate IDs: OMICS_01066


Ratings and Alerts

No rating or validation information has been found for Picard.
No alerts have been found for Picard.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5550 mentions in open access literature.

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


Li L, et al. (2021) Tumor-derived mutations in postoperative plasma of colorectal cancer with microsatellite instability. Translational oncology, 14(1), 100945.

Jia R, et al. (2021) Post-radiation circulating tumor DNA as a prognostic factor in locally
advanced esophageal squamous cell carcinoma. Oncology letters, 21(1), 68.


Sun KA, et al. (2020) Endogenous itaconate is not required for particulate matter-induced NRF2 expression or inflammatory response. eLife, 9.


