SMRT-Analysis
RRID:SCR_002942
Type: Tool

Proper Citation

SMRT-Analysis (RRID:SCR_002942)

Resource Information

URL: https://github.com/PacificBiosciences/SMRT-Analysis/

Description: Open-source bioinformatics software suite for analyzing single molecule, real-time DNA sequencing data. Users can choose from a variety of analysis protocols that utilize PacBio and third-party tools. Analysis protocols include de novo genome assembly, cDNA mapping, DNA base-modification detection, and long-amplicon analysis to determine phased consensus sequences.

Resource Name: SMRT-Analysis
Proper Citation: SMRT-Analysis (RRID:SCR_002942)
Resource Type: Resource, software resource
Keywords: software suite
Resource ID: SCR_002942
Availability: Open unspecified license
Website Status: Last checked up
Alternate IDs: OMICS_05142
Alternate URLs: http://www.pacb.com/devnet/
Mentions Count: 47

Ratings and Alerts
No rating or validation information has been found for SMRT-Analysis.
No alerts have been found for SMRT-Analysis.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 47 mentions in open access literature.

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


