

Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

Sequencing Analysis Software

RRID:SCR_000718

Type: Tool

Proper Citation

Sequencing Analysis Software (RRID:SCR_000718)

Resource Information

URL: <https://www.thermofisher.com/order/catalog/product/4474950>

Proper Citation: Sequencing Analysis Software (RRID:SCR_000718)

Description: A software that gives the user the ability to basecall, trim, display, edit, and print data for the entire line of capillary DNA sequencing instruments for data analysis and quality control. This software benefits from being able to obtain longer read lengths, greater accuracy on the 5' end, and the ability to filter out low-quality sequence ends.

Abbreviations: Sequencing Analysis Software

Resource Type: software resource

Keywords: basecall, capillary dna, quality control, low-quality sequence ends

Availability: Restricted

Resource Name: Sequencing Analysis Software

Resource ID: SCR_000718

Alternate IDs: OMICS_01814

License: Commercial license

Record Creation Time: 20220129T080203+0000

Record Last Update: 20260912T195514+0000

Ratings and Alerts

No rating or validation information has been found for Sequencing Analysis Software.

No alerts have been found for Sequencing Analysis Software.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Kobunai T, et al. (2010) The frequency of KRAS mutation detection in human colon carcinoma is influenced by the sensitivity of assay methodology: a comparison between direct sequencing and real-time PCR. Biochemical and biophysical research communications, 395(1), 158.