# **Resource Summary Report**

Generated by <u>dkNET</u> on Apr 29, 2025

# **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

Funding:

Availability: Commercial license

Resource Name: Sequencing Analysis Software

Resource ID: SCR\_000718

Alternate IDs: OMICS\_01814

Record Creation Time: 20220129T080203+0000

Record Last Update: 20250420T014003+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: <u>SciCrunch Registry</u>

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

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.