

Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

Knime4Bio

RRID:SCR_005376

Type: Tool

Proper Citation

Knime4Bio (RRID:SCR_005376)

Resource Information

URL: <https://code.google.com/p/knime4bio/>

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Description: A set of custom nodes for the KNIME (The Konstanz Information Miner) graphical workbench, for analysing next-generation sequencing (NGS) data without the requirement of programming skills.

Abbreviations: Knime4Bio

Synonyms: Knime4Bio: custom nodes for the interpretation of Next Generation Sequencing data with KNIME

Resource Type: software resource

Defining Citation: [PMID:21984761](#)

Keywords: node, next-generation sequencing, knime, bioinformatics, workflow, sequencing, flow, data, bam, wig, bed, bio.tools

Availability: GNU General Public License, v3

Resource Name: Knime4Bio

Resource ID: SCR_005376

Alternate IDs: biotools:knime4bio, OMICS_01143

Alternate URLs: <https://bio.tools/knime4bio>

Record Creation Time: 20220129T080229+0000

Record Last Update: 20260725T190603+0000

Ratings and Alerts

No rating or validation information has been found for Knime4Bio.

No alerts have been found for Knime4Bio.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Rimbert A, et al. (2016) Identification of novel APOB mutations by targeted next-generation sequencing for the molecular diagnosis of familial hypobetalipoproteinemia. *Atherosclerosis*, 250, 52.

Veerman CC, et al. (2016) hiPSC-derived cardiomyocytes from Brugada Syndrome patients without identified mutations do not exhibit clear cellular electrophysiological abnormalities. *Scientific reports*, 6, 30967.