

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

CCRAVAT

RRID:SCR_009141

Type: Tool

Proper Citation

CCRAVAT (RRID:SCR_009141)

Resource Information

URL: <http://www.sanger.ac.uk/resources/software/rarevariant/>

Proper Citation: CCRAVAT (RRID:SCR_009141)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16, 2026. Software application for enabling the analysis of rare variants in large-scale case control and quantitative trait association studies. CCRaVAT (Case-Control Rare Variant Analysis Tool) and QuTie (Quantitative Trait) are software packages that enable efficient large-scale analysis of rare variants across specific regions or genome-wide. These programs implement a rare variant super-locus or collapsing method that investigates the accumulation of rare variant alleles in either a case-control or quantitative trait study design. (entry from Genetic Analysis Software)

Abbreviations: CCRAVAT

Synonyms: Case-Control RAre Variant Analysis Tool QUTIE

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: CCRAVAT

Resource ID: SCR_009141

Alternate IDs: nlx_154260

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260731T042808+0000

Ratings and Alerts

No rating or validation information has been found for CCRAVAT.

No alerts have been found for CCRAVAT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.