

# Resource Summary Report

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## 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 application, software resource

**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:** 20260912T200240+0000

### Ratings and Alerts

No rating or validation information has been found for CCRAVAT.

No alerts have been found for CCRAVAT.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.