GenABEL (RRID:SCR_001842)

**Type:** Tool

**Proper Citation:**

GenABEL (RRID:SCR_001842)

**Resource Information**

**URL:** [http://www.genabel.org/packages/GenABEL](http://www.genabel.org/packages/GenABEL)

**Description:** An R software library for genome-wide association analysis for quantitative, binary and time-till-event traits.

**Resource Name:** GenABEL

**Proper Citation:** GenABEL (RRID:SCR_001842)

**Resource Type:** Resource, software resource, software toolkit, software library

**Keywords:** r, genome-wide association, single nucleotide polymorphism

**Resource ID:** SCR_001842

**Funding Agency:** Centre for Medical Systems Biology; Netherlands, Netherlands Genomics Initiative, Netherlands Organisation for Scientific Research, Russian Foundation for Basic Research

**References:** PMID:17384015

**Availability:** Free

**Website Status:** Last checked up

**Alternate IDs:** nlx_154328, OMICS_00234

**Alternate URLs:** [http://mga.bionet.nsc.ru/~yurii/ABEL/GenABEL/](http://mga.bionet.nsc.ru/~yurii/ABEL/GenABEL/), [https://cran.r-project.org/web/packages/GenABEL/index.html](https://cran.r-project.org/web/packages/GenABEL/index.html)
Ratings and Alerts

No rating or validation information has been found for GenABEL.
No alerts have been found for GenABEL.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 330 mentions in open access literature.

**Listed below are recent publications.** The full list is available at scicrunch.


Das RG, et al. (2019) Genome-wide association study and whole-genome sequencing identify a deletion in LRIT3 associated with canine congenital stationary night blindness. Scientific reports, 9(1), 14166.


