

Resource Summary Report

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LRTAE

RRID:SCR_009265

Type: Tool

Proper Citation

LRTAE (RRID:SCR_009265)

Resource Information

URL: <http://ftp://linkage.rockefeller.edu/software/lrtae/>

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Description: Software application to compute a likelihood ratio test statistic that increases power to detect genetic association in the presence of phenotype, genotype, and/or haplotype misclassification errors. In addition, the program produces asymptotically unbiased estimates of frequency parameters. (entry from Genetic Analysis Software)

Abbreviations: LRTAE

Synonyms: Likelihood Ratio Test Allowing for Errors

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, ms-windows, unix, solaris, linux

Resource Name: LRTAE

Resource ID: SCR_009265

Alternate IDs: nlx_154443

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260919T195802+0000

Ratings and Alerts

No rating or validation information has been found for LRTAE.

No alerts have been found for LRTAE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Levenstien MA, et al. (2006) Are molecular haplotypes worth the time and expense? A cost-effective method for applying molecular haplotypes. PLoS genetics, 2(8), e127.