Resource Summary Report

Generated by dkNET on Apr 30, 2025

ERPA

RRID:SCR_009173
Type: Tool

Proper Citation

ERPA (RRID:SCR_009173)

Resource Information

URL: http://www.mds.qmw.ac.uk/statgen/dcurtis/software.html

Proper Citation: ERPA (RRID:SCR_009173)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16,2023. Software application for non-parametric analysis (entry from Genetic Analysis Software)

Abbreviations: ERPA

Synonyms: Extended Relative Pair Analysis

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c, ms-dos

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ERPA

Resource ID: SCR_009173

Alternate IDs: nlx_154302

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250429T055313+0000

Ratings and Alerts

No rating or validation information has been found for ERPA.

No alerts have been found for ERPA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Alemayehu TG, et al. (2023) Occupational radiation exposure dose and associated factors among radiology personnel in Eastern Amhara, Ethiopia. PloS one, 18(5), e0286400.

Hammami F, et al. (2023) Analysis of a logical regulatory network reveals how Fe-S cluster biogenesis is controlled in the face of stress. microLife, 4, uqad003.

Liu JK, et al. (2019) Predicting proteome allocation, overflow metabolism, and metal requirements in a model acetogen. PLoS computational biology, 15(3), e1006848.

Tu Y, et al. (2017) Association between rs3087243 and rs231775 polymorphism within the cytotoxic T-lymphocyte antigen 4 gene and Graves' disease: a case/control study combined with meta-analyses. Oncotarget, 8(66), 110614.

Shen GQ, et al. (2007) An LRP8 variant is associated with familial and premature coronary artery disease and myocardial infarction. American journal of human genetics, 81(4), 780.

Kato H, et al. (2006) Association of single-nucleotide polymorphisms in the suppressor of cytokine signaling 2 (SOCS2) gene with type 2 diabetes in the Japanese. Genomics, 87(4), 446.

Chan VS, et al. (2006) Homozygous L-SIGN (CLEC4M) plays a protective role in SARS coronavirus infection. Nature genetics, 38(1), 38.