

Resource Summary Report

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EraSOR

RRID:SCR_023544

Type: Tool

Proper Citation

EraSOR (RRID:SCR_023544)

Resource Information

URL: <https://choishingwan.gitlab.io/EraSOR/>

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Description: Software Python tool for removing bias introduced from having overlapped samples between base GWAS data and target genotype data for Polygenic Risk Score analyses. Used for adjusting inflation in PRS prediction and association statistics in presence of sample overlap or close relatedness between GWAS and target samples.

Synonyms: Erase Sample Overlap and Relatedness

Resource Type: data analysis software, data processing software, software application, software resource

Defining Citation: [DOI:10.1101/2021.12.10.472164](https://doi.org/10.1101/2021.12.10.472164)

Keywords: removing bias, overlapped samples, GWAS data, target genotype data, Polygenic Risk Score, PRS prediction and association statistics,

Availability: Free, Available for download, Freely available

Resource Name: EraSOR

Resource ID: SCR_023544

Record Creation Time: 20230504T050216+0000

Record Last Update: 20260829T182828+0000

Ratings and Alerts

No rating or validation information has been found for EraSOR.

No alerts have been found for EraSOR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.