

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

randomizeR

RRID:SCR_023568

Type: Tool

Proper Citation

randomizeR (RRID:SCR_023568)

Resource Information

URL: <https://www.rdocumentation.org/packages/randomizeR/versions/3.0.1>

Proper Citation: randomizeR (RRID:SCR_023568)

Description: Software R package for randomization of clinical trials. Provides sound scientific evidence for choice of randomization procedure, on the basis of well founded criteria. Includes functions for generating randomization sequences from various randomization procedures as well as various issues for assessment of randomization procedures.

Synonyms: Randomization for Clinical Trials

Resource Type: software resource, software toolkit

Keywords: randomization of clinical trials, generating randomization sequences, clinical trials, randomization sequences

Availability: Free, Available for download, Freely available

Resource Name: randomizeR

Resource ID: SCR_023568

Alternate URLs: <https://CRAN.R-project.org/package=randomizeR>

License: GPL v3

Record Creation Time: 20230516T050215+0000

Record Last Update: 20260905T133314+0000

Ratings and Alerts

No rating or validation information has been found for randomizeR.

No alerts have been found for randomizeR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Charmas M, et al. (2025) Ashwagandha Does Not Enhance the Effect of High-Intensity Interval Training on Selected Energy Metabolism Parameters in Young Healthy Men. *Nutrients*, 17(20).

Wang Y, et al. (2024) Boosting Smoking Cessation Intervention Utilization in Chinese Health Care Providers: A Randomized Controlled Trial of the "WeChat WeQuit" Medical Education Program. *Nicotine & tobacco research : official journal of the Society for Research on Nicotine and Tobacco*, 27(1), 61.

Reva ON, et al. (2005) Differentiation of regions with atypical oligonucleotide composition in bacterial genomes. *BMC bioinformatics*, 6, 251.