

Package: EGSEA123 (via r-universe)

October 8, 2026

Title Easy and efficient ensemble gene set testing with EGSEA

Version 1.37.0

Date 2021-08-04

Description R package that supports the workflow article 'Easy and efficient ensemble gene set testing with EGSEA', Alhamdoosh et al. (2017), F1000Research, 6:2010.

Depends R (>= 3.4.0), EGSEA (>= 1.5.2), limma (>= 3.49.2), edgeR, illuminaio

License Artistic-2.0

Encoding UTF-8

VignetteBuilder knitr

biocViews ImmunoOncologyWorkflow, Workflow, GeneExpressionWorkflow

Suggests knitr, rmarkdown, BiocStyle

NeedsCompilation no

Workflow True

Author Monther Alhamdoosh, Charity Law, Luyi Tian, Julie Sheridan, Milica Ng and Matthew Ritchie

Maintainer Matthew Ritchie <mritchie@wehi.edu.au>

URL <https://f1000research.com/articles/6-2010>

Config/pak/sysreqs cmake libfftw3-dev make libmagick++-dev gsfonts libicu-dev libpng-dev libuv1-dev libxml2-dev libssl-dev zlib1g-dev

Repository <https://bioc.r-universe.dev>

Date/Publication 2026-04-28 12:40:26 UTC

RemoteUrl <https://github.com/bioc/EGSEA123>

RemoteRef HEAD

RemoteSha 61bf37a4e6010f0231bead98f783066a8ff69d88