

Package ‘Rgin’

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Title gin in R

Version 1.2.0

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Description C++ implementation of SConES.

Depends R (>= 3.5)

LinkingTo RcppEigen (>= 0.3.3.3.0)

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Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

biocViews Software, GenomeWideAssociation, SNP, GeneticVariability,
Genetics, FeatureExtraction, GraphAndNetwork, Network

Suggests knitr, rmarkdown

VignetteBuilder knitr

git_url <https://git.bioconductor.org/packages/Rgin>

git_branch RELEASE_3_8

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R topics documented: