

# Package ‘Rgin’

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**Title** gin in R

**Version** 1.6.0

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**Description** C++ implementation of SConES.

**Depends** R (>= 3.5)

**LinkingTo** RcppEigen (>= 0.3.3.5.0)

**License** MIT + file LICENSE

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.1.1

**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\_10

**git\_last\_commit** 8721092

**git\_last\_commit\_date** 2019-10-29

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**R topics documented:**