

Package ‘saseR’

August 30, 2025

Type Package

Title Scalable Aberrant Splicing and Expression Retrieval

Date 2023-06-01

Version 1.5.1

Description saseR is a highly performant and fast framework for aberrant expression and splicing analyses. The main functions are:

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biocViews DifferentialExpression, DifferentialSplicing, Regression,
GeneExpression, AlternativeSplicing, RNASeq, Sequencing,
Software

BugReports <https://github.com/statOmics/saseR/issues>

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

git_branch devel

