

Package ‘alabaster’

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Title Umbrella for the Alabaster Framework

Version 1.2.0

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Description

Umbrella for the alabaster suite, providing a single-line import for all alabaster.* packages. Installing this package ensures that all known alabaster.* packages are also installed, avoiding problems with missing packages when a staging method or loading function is dynamically requested. Obviously, this comes at the cost of needing to install more packages, so advanced users and application developers may prefer to install the required alabaster.* packages individually.

Depends alabaster.base

Imports alabaster.matrix, alabaster.ranges, alabaster.se,
alabaster.sce, alabaster.spatial, alabaster.string,
alabaster.vcf, alabaster.bumpy, alabaster.mae

Suggests knitr, rmarkdown, BiocStyle

VignetteBuilder knitr

RoxygenNote 7.2.3

biocViews DataRepresentation, DataImport

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

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R topics documented: