

Package ‘beachmat’

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Version 2.4.0

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Title Compiling Bioconductor to Handle Each Matrix Type

Encoding UTF-8

Imports methods, DelayedArray, BiocGenerics, Matrix

Suggests testthat, BiocStyle, knitr, rmarkdown, devtools

biocViews DataRepresentation, DataImport, Infrastructure

Description Provides a consistent C++ class interface for reading from and writing data to a variety of commonly used matrix types. Ordinary matrices and several sparse/dense Matrix classes are directly supported, third-party S4 classes may be supported by external linkage, while all other matrices are handled by DelayedArray block processing.

License GPL-3

NeedsCompilation yes

VignetteBuilder knitr

SystemRequirements C++11

RoxygenNote 6.1.1

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

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R topics documented: