Package 'assorthead'

November 11, 2026

Version 1.6.4

Date 2026-10-08

Title Assorted Header-Only C++ Libraries

Description Vendors an assortment of useful header-only C++ libraries.
Bioconductor packages can use these libraries in their own C++ code by LinkingTo this package without introducing any additional dependencies.
The use of a central repository avoids duplicate vendoring of libraries across multiple R packages, and enables better coordination of version updates across cohorts of interdependent C++ libraries.

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Suggests knitr, rmarkdown, BiocStyle

VignetteBuilder knitr

URL https://github.com/LTLA/assorthead

BugReports https://github.com/LTLA/assorthead/issues

Encoding UTF-8

biocViews SingleCell, QualityControl, Normalization, DataRepresentation, DataImport, DifferentialExpression, Alignment

RoxygenNote 7.3.3

git_url https://git.bioconductor.org/packages/assorthead

git_branch RELEASE_3_23

git_last_commit fb2bf57

git_last_commit_date 2026-10-07

Repository Bioconductor 3.23

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Contents

assorthead-package 2

Index **3**

`assorthead-package` *assorthead: Assorted Header-Only C++ Libraries*

Description

Vendors an assortment of useful header-only C++ libraries. Bioconductor packages can use these libraries in their own C++ code by LinkingTo this package without introducing any additional dependencies. The use of a central repository avoids duplicate vendoring of libraries across multiple R packages, and enables better coordination of version updates across cohorts of interdependent C++ libraries.

Author(s)

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See Also

Useful links:

- https://github.com/LTLA/assorthead
- Report bugs at https://github.com/LTLA/assorthead/issues

Index

assorthead *(*assorthead-package*)*, 2
assorthead-package, 2