

Package ‘assorthead’

May 1, 2026

Version 1.7.0

Date 2026-04-17

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 devel

git_last_commit 43af464

git_last_commit_date 2026-04-28

Repository Bioconductor 3.24

Date/Publication 2026-05-01

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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>

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