

Package: assorthead (via r-universe)

October 8, 2026

Version 1.7.11

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

Repository <https://bioc.r-universe.dev>

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RemoteUrl <https://github.com/bioc/assorthead>

RemoteRef HEAD

RemoteSha 8712ebd493cef14227dd4d48c99126aef61b09eb

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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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