

# Package ‘assorthead’

January 25, 2026

**Version** 1.5.8

**Date** 2026-01-23

**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

**git\_url** <https://git.bioconductor.org/packages/assorthead>

**git\_branch** devel

**git\_last\_commit** 0a3e503

**git\_last\_commit\_date** 2026-01-22

**Repository** Bioconductor 3.23

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