Package: CardinalStats (via r-universe)

November 17, 2026

Type Package

Title Statistical learning for mass spectrometry imaging

Version 0.0.1

Date YYYY-MM-DD

Description Statistical modeling and machine learning methods for mass spectrometry imaging analysis in Cardinal

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Depends R (>= 4.6)

Imports methods

Suggests BiocStyle, testthat (>= 3.0.0), knitr, rmarkdown

VignetteBuilder knitr

biocViews Software, MassSpectrometry, ImagingMassSpectrometry, Clustering, Classification, Regression

URL http://www.cardinalmsi.org

BugReports https://github.com/kuwisdelu/CardinalStats/issues

Config/testthat/edition 3

Repository https://cardinalverse.r-universe.dev

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RemoteUrl https://github.com/Cardinalverse/CardinalStats

RemoteRef HEAD

RemoteSha aa7425fc0cde91d33c9c8dcfc2efe72bc8fa5732

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`CardinalStats-package` *Short description of package*

Description

Longer description of package.

Details

Detailed description of package.

See `browseVignettes("CardinalStats")` to view a user guide and vignettes of common workflows.

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