

Package ‘Prostar’

October 12, 2021

Type Package

Title Provides a GUI for DAPAR

Version 1.24.8

Date 2021-08-31

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Description This package provides a GUI interface for DAPAR.

biocViews Proteomics, MassSpectrometry, Normalization, Preprocessing,
ImmunoOncology, R.utils, GO, GUI

License Artistic-2.0

Depends R (>= 4.1.0)

Suggests BiocStyle, testthat

Imports DAPAR (>= 1.24.8), DAPARdata (>= 1.22.2), rhandsontable,
data.table, shinyjs, DT, shiny, shinyBS, shinyAce, highcharter,
htmlwidgets, webshot, R.utils, shinythemes, XML, later,
rclipboard, shinycssloaders, future, promises, stringr,
colourpicker, BiocManager, shinyjs, shinyTree, shinyWidgets,
sass, tibble

NeedsCompilation no

RoxygenNote 7.1.1

Encoding UTF-8

URL <http://www.prostar-proteomics.org/>

BugReports <https://github.com/samWieczorek/Prostar/issues>

git_url <https://git.bioconductor.org/packages/Prostar>

git_branch RELEASE_3_13

git_last_commit 5886e47

git_last_commit_date 2021-09-01

Date/Publication 2021-10-12

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Prostar

Prostar

Description

Prostar : a GUI for DAPAR

Usage

Prostar()

Value

A new window in the default internet browser

Author(s)

Samuel Wiczorek

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