

Package ‘Prostar’

April 12, 2022

Type Package

Title Provides a GUI for DAPAR

Version 1.26.4

Date 2022-01-22

Author Samuel Wiczorek [cre,aut],
Thomas Burger [aut],
Enora Fremy [aut]

Maintainer Samuel Wiczorek <samuel.wiczorek@cea.fr>

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 (>= 1.26.1), DAPARdata (>= 1.24.0), rhandsontable,
data.table, shinyjs, DT, shiny, shinyBS, shinyAce, highcharter,
htmlwidgets, webshot, R.utils, shinythemes, XML, later,
rclipboard, shinycssloaders, future, promises, colourpicker,
BiocManager, shinyjquery, shinyTree, shinyWidgets, sass, tibble

NeedsCompilation no

RoxygenNote 7.1.2

Encoding UTF-8

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

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

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

git_branch RELEASE_3_14

git_last_commit 8bb9825

git_last_commit_date 2022-01-21

Date/Publication 2022-04-12

R topics documented:

Prostar	2
Index	3

Prostar

Prostar

Description

Prostar : a GUI for DAPAR##'

Usage

Prostar()

Value

A new window in the default internet browser

Author(s)

Samuel Wiczorek

Examples

```
if(interactive()) {Prostar()}
```

Index

Prostar, [2](#)