

Package ‘Prostar’

March 30, 2021

Type Package

Title Provides a GUI for DAPAR

Version 1.22.8

Date 2021-03-08

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

Maintainer Samuel Wieczorek <samuel.wieczorek@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 (>= 4.0.3)

Suggests BiocStyle, testthat

Imports DAPAR (>= 1.22.6), DAPARdata (>= 1.20.0), rhandsontable,
data.table, shinyjs, DT, shiny, shinyBS, shinyAce, highcharter,
htmlwidgets, webshot, R.utils, shinythemes, XML, later,
rclipboard, shinycssloaders, future, promises, colourpicker,
BiocManager, shinyjqui, 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_12

git_last_commit e86dd05

git_last_commit_date 2021-03-24

Date/Publication 2021-03-29

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 Wieczorek

Examples

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

Index

Prostar, [2](#)