

Package ‘Prostar’

April 15, 2020

Type Package

Title Provides a GUI for DAPAR

Version 1.18.6

Date 2020-01-22

Author Samuel Wieczorek [cre,aut],
Florence Combes [aut],
Thomas Burger [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 (>= 3.6.1)

Suggests BiocStyle, testthat

Imports DAPAR (>= 1.18.1), DAPARdata (>= 1.16.0), rhandsontable,
data.table, shinyjs, DT, shiny, shinyBS, shinyAce, highcharter
(>= 0.7.0), htmlwidgets, webshot, R.utils, shinythemes,
XML, later, rclipboard, shinycssloaders, future, promises,
colourpicker, BiocManager, shinyTree, shinyWidgets, sass,
tibble

NeedsCompilation no

RoxygenNote 6.0.1

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

git_branch RELEASE_3_10

git_last_commit d0ea961

git_last_commit_date 2020-01-28

Date/Publication 2020-04-14

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](#)