

Package ‘Prostar’

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Type Package

Title Provides a GUI for DAPAR

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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, BiocManager, testthat, shinyTree, rclipboard

Imports DAPAR (>= 1.27.7), DAPARdata (>= 1.22.2), rhandsontable,
data.table, shinyjs, DT, shiny, shinyBS, shinyAce, highcharter,
htmlwidgets, webshot, R.utils, shinythemes, XML, later,
shinycssloaders, future, promises, colourpicker, shinyjqm,
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 master

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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)

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Examples

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

Index

Prostar, [2](#)