

# Package ‘Prostar’

October 14, 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, shinyjqui, shinyTree, shinyWidgets,  
sass, tibble

**NeedsCompilation** no

**RoxygenNote** 7.1.1

**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\_13

**git\_last\_commit** 5886e47

**git\_last\_commit\_date** 2021-09-01

**Date/Publication** 2021-10-14

**R topics documented:**

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Prostar

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**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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