

# Package ‘Prostar’

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**Type** Package

**Title** Provides a GUI for DAPAR

**Version** 1.20.0

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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 (>= 3.6.2)

**Suggests** BiocStyle, testthat

**Imports** DAPAR (>= 1.19.16), DAPARdata (>= 1.16.0), 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** 7.1.0

**git\_url** <https://git.bioconductor.org/packages/Prostar>

**git\_branch** RELEASE\_3\_11

**git\_last\_commit** 0327129

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**Date/Publication** 2020-10-16

## R topics documented:

Prostar . . . . .	2
Index	3

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 Wieczorek

**Examples**

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

# Index

Prostar, [2](#)