

# Package ‘Prostar’

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**Type** Package

**Title** Provides a GUI for DAPAR

**Version** 1.14.12

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**Description** This package provides a GUI interface for DAPAR.

**biocViews** ImmunoOncology, MassSpectrometry, Normalization,  
Preprocessing, R.utils Proteomics,GO, GUI

**License** Artistic-2.0

**Depends** R (>= 3.5)

**Suggests** BiocStyle, testthat

**Imports** DAPAR (>= 1.13.15), DAPARdata (>= 1.11.2), 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

**NeedsCompilation** no

**RoxygenNote** 6.0.1

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

**git\_branch** RELEASE\_3\_8

**git\_last\_commit** 2bae2f2

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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 Wieczorek

**Examples**

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

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