

# Package ‘Prostar’

November 20, 2025

**Type** Package

**Title** Provides a GUI for DAPAR

**Version** 1.43.0

**Date** 2024-01-17

**Description** This package provides a GUI interface for the DAPAR package. The package Prostar (Proteomics statistical analysis with R) is a Bioconductor distributed R package which provides all the necessary functions to analyze quantitative data from label-free proteomics experiments. Contrarily to most other similar R packages, it is endowed with rich and user-friendly graphical interfaces, so that no programming skill is required.

**biocViews** Proteomics, MassSpectrometry, Normalization, Preprocessing, Software, GUI

**License** Artistic-2.0

**Depends** R (>= 4.4.0)

**Suggests** BiocStyle, BiocManager, testthat, knitr

**Imports** DAPAR (>= 1.35.1), DAPARdata (>= 1.30.0), rhandsontable, data.table, shiny, shinyBS, shinyAce, highcharter, htmlwidgets, webshot, shinythemes, later, shinycssloaders, future, promises, shinyjs, tibble, ggplot2, gplots, shinyjs, vioplot, Biobase, DT, R.utils, RColorBrewer, XML, colourpicker, gtools, markdown, rclipboard, sass, shinyTree, shinyWidgets

**NeedsCompilation** no

**RoxygenNote** 7.3.2

**Encoding** UTF-8

**URL** <http://www.prostar-proteomics.org/>

**BugReports** <https://github.com/edyp-lab/Prostar/issues>

**VignetteBuilder** knitr

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

**git\_branch** devel

**git\_last\_commit** cfb7b6c  
**git\_last\_commit\_date** 2025-10-29  
**Repository** Bioconductor 3.23  
**Date/Publication** 2025-11-19  
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|                  |     |
|------------------|-----|
| BuildColorStyles | xxx |
|------------------|-----|

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Description

xxx

Usage

BuildColorStyles(obj)

Arguments

|     |    |
|-----|----|
| obj | xx |
|-----|----|

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|                 |     |
|-----------------|-----|
| getDataForExprs | xxx |
|-----------------|-----|

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Description

xxx

Usage

getDataForExprs(obj, digits = NULL)

Arguments

|        |     |
|--------|-----|
| obj    | xx  |
| digits | xxx |

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Prostar

*Prostar*

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**Description**

Prostar

**Usage**

Prostar()

**Value**

A new window in the default internet browser

**Author(s)**

Samuel Wieczorek

**Examples**

```
## Not run:  
  Prostar()  
  
## End(Not run)
```

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