pOmics: Tidy Proteomics Data Analysis in R

Nico Hüttmann

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Preface

This is my attempt to make proteomics data analysis in R accessible to other scientists. Took me a while to learn it, so why not share it.

Main features:

- data structure for simple loading and handling of data
- statistical analysis for proteomics data
- functional enrichment via cluster Profiler
- \bullet tidyverse friendly
- ggplot2, for publication-quality figures
- plotly, for interactive data exploration
- R Markdown, for efficient communication

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Introduction

Hi.

- 1.1 Setup R and RStudio
- 1.2 R Basics

Bye.

Getting started with R and RStudio

Overview pOmics

Importing data

Let's get started with importing our raw data into R.

The easiest way is to use the import_files() function.

Basic analysis workflow

Bonjour

5.1 A

Hola