

# Package: OlinkAnalyzeVignettes (via r-universe)

June 9, 2026

**Type** Package

**Title** Vignettes for Analyzing Data using 'OlinkAnalyze'

**Version** 1.0.1

**Description** Exemplifying analysis of large-scale protein data from the 'Olink platform', primarily relative protein expression data that has been exported from 'Olink NPX Software', as well as QUANT data from 'Olink'. QUANT data is log-transformed. Materials focus on reading data, demonstrating data wrangling and quality control analysis, performing statistical analysis and generating figures to visualize the results of the statistical analysis. The goal of this package is to guide users extract biological insights from large-scale protein data run on the 'Olink platform'. More information on 'Olink' data can be found at <https://olink.com/>.

**Contact** biostattools@olink.com

**URL** <https://olink.com/>  
<https://github.com/Olink-Proteomics/OlinkRPackage>

**License** AGPL (>= 3)

**Encoding** UTF-8

**Depends** R (>= 4.1.0)

**Suggests** dplyr (>= 1.2.0), ggplot2, ggpubr, ggrepel, kableExtra, knitr, OlinkAnalyze, stringr, tidyr

**VignetteBuilder** knitr

**Config/roxygen2/version** 8.0.0

**NeedsCompilation** no

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**Repository** <https://kathy-nevola.r-universe.dev>

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open\_docs

*Open the Olink Analyze GitHub repo page*

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## Description

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*open\_docs*

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**Usage**

`open_docs()`

**Value**

None opens Olink Analyze repo page in browser

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