

Goals

1

Facilitate the analysis of
Mass-Spectrometry based **Proteomics** data
using the  **programming language**

2

Provide different levels of analysis to ensure **data quality, quantification,** and frame results in a **biological context.**

artMS

**Analytical R Tools
for Mass Spectrometry**

<http://artms.org>




artMS

Analytical R Tools
for Mass Spectrometry

<http://artms.org>



Goals

- 1 **Facilitate** the analysis of **Mass-Spectrometry** based **Proteomics** data using the  **programming language**
- 2 **Provide** different levels of analysis to ensure **data quality**, **quantification**, and frame results in a **biological context**.

artMS

Analytical R Tools
for Mass Spectrometry

<http://artms.org>



Quantification Methods

Label Free
SILAC

(+/- Fractionation)