#### ונ

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

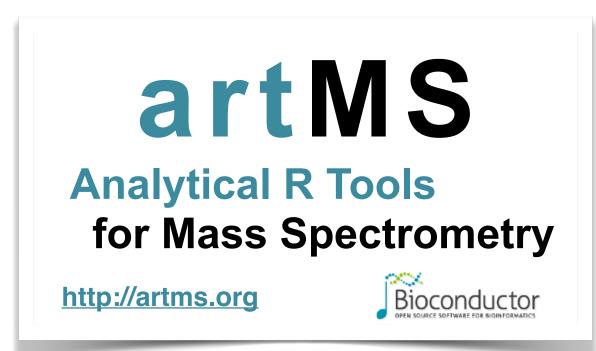
#### Provide different levels of analysis to ensure data quality, quantification, and frame results in a biological context.

### artMS

**Analytical R Tools for Mass Spectrometry** 

http://artms.org

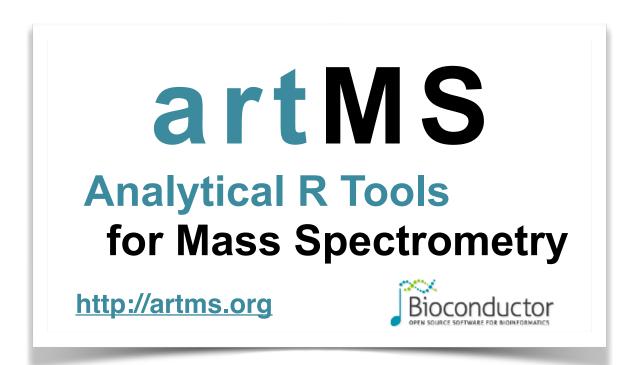




## Goals

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

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



**Quantification Methods** 

# Label Free (+/- Fractionation) SILAC