



Label Free  
SILAC

# artMS

**Analytical R Tools  
for Mass Spectrometry**

<http://artms.org>



**(+/- Fraction)**

Quantification Methods

Analysis of

+Global Proteome

+AP-MS

+PostTranslational Modifications

Phosphorylation (PH)

Ubiquitination (UB)

Acetylation (AC)

(and any MaxQuant supported PTM)

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Motivation