





<https://bioconductor.org/packages/release/bioc/html/artMS.html>



**GitHub**

<https://github.com/biodavidjm/artMS>



# Documentation:

and everything else...

<http://artms.org>

# artMS

**Analytical R Tools  
for Mass Spectrometry**

<http://artms.org>





# artMS: Analytical R Tools for Mass Spectrometry-based Proteomics

David Jimenez-Morales<sup>1\*</sup>, Alexandre Rosa Campos<sup>2</sup>, Jeff Johnson, John Von Dollen<sup>3</sup>,  
Matthew T Wheeler<sup>1</sup>, Euan A Ashley<sup>1</sup>, Nevan J. Krogan<sup>3</sup>, Danielle L. Swaney<sup>3\*</sup>

*In preparation*

1. Stanford

2. SBP

3. UCSF

# artMS

Analytical R Tools  
for Mass Spectrometry

<http://artms.org>



<https://bioconductor.org/packages/release/bioc/html/artMS.html>



**GitHub**

<https://github.com/biodavidjm/artMS>



**Documentation:**

and everything else...

<http://artms.org>



## artMS: Analytical R Tools for Mass Spectrometry-based Proteomics

David Jimenez-Morales<sup>1\*</sup>, Alexandre Rosa Campos<sup>2</sup>, Jeff Johnson, John Von Dollen<sup>3</sup>,  
Matthew T Wheeler<sup>1</sup>, Euan A Ashley<sup>1</sup>, Nevan J. Krogan<sup>3</sup>, Danielle L. Swaney<sup>3\*</sup>

*In preparation*

1. Stanford  
2. SBP  
3. UCSF

# artMS

Analytical R Tools  
for Mass Spectrometry

<http://artms.org>



## Case Studies



## Protein Phosphorylation