



ABRF 2017 Annual Meeting

A FORUM FOR ADVANCING TODAY'S CORE TECHNOLOGIES TO ENABLE
TOMORROW'S INNOVATIONS

sPRG:

**Development and Characterization of
a Stable-Isotope Labeled
Phosphopeptide Standard**



March 25-28, 2017

Town and Country Resort & Convention Center
San Diego, California



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Current sPRG Working Group Members

- Antonius Koller (Chair) - *Columbia University*
- Christopher Colangelo - *Primary Ion*
- David Hawke - *UT MD Anderson Cancer Center*
- Gordana Isovev - *Sciex*
- Brian C. Searle - *Proteome Software Inc.*
- Brett S Phinney - *Proteomics Core UC Davis Genome Center*
- Ryan Leib – *Stanford University*
- Bhavinkumar Patel – *Thermo Fisher*
- Allis Chien (EB Liaison) – *Stanford University*



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Past sPRG Members

- Alexander Ivanov – *Northeastern University*
- Craig P. Dufresne – *Thermo Fisher Scientific*
- Scott A. Shaffer – *University of Massachusetts Medical School*



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Mission of sPRG

- Design, Develop and Make ABRF standards available to the community
- Prepare Manuscript



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Short History of sPRG projects

- 2003: 2 digested proteins; 2 synthetic phosphopeptides
- 2007: mixture of 7 phosphorylated proteins
- 2010: 6 digested proteins; 23 synthetic phosphopeptides
- 2012: 6 digested proteins; 45 synthetic phosphopeptides; 41 synthetic modified peptides
 - <http://spectragen-informatics.com/sprg>
- 2015: 1000 stable-isotope labeled peptides
 - <http://www.jpt.com/> SpikeMix ABRF (cross-species standard)



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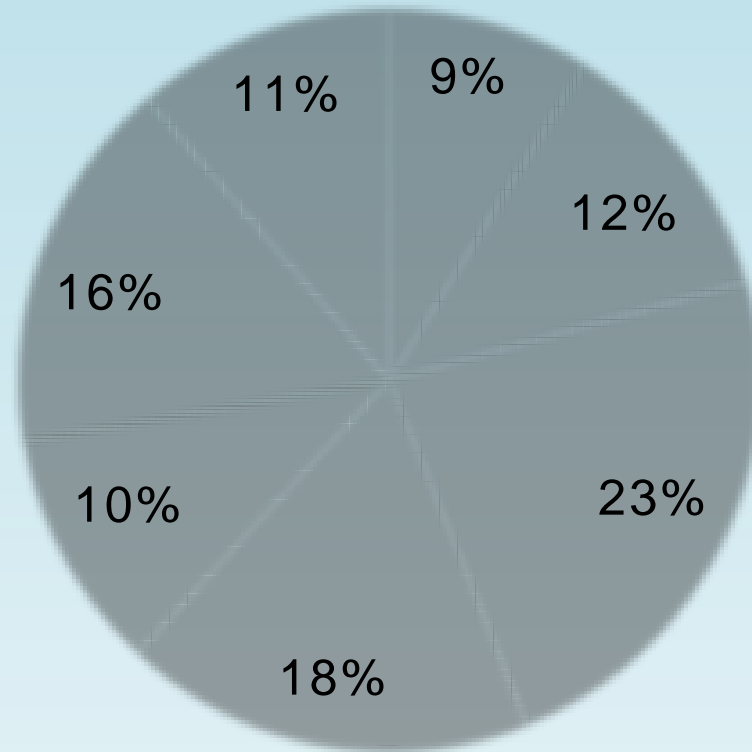
sPRG2017-2018

- **Development of 150 stable-isotope labeled phosphopeptides standard**
- Biologically relevant
- Useful for method development



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Signaling Pathways included



AMPK signaling

Death and apoptosis
signaling

EGFR/ HER signaling

Insulin/ IGF-1 signaling

mTOR signaling

PI3K/ AKT signaling

Stress (p38/ SAPK/ JNK)
signaling



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Phosphorylation site distribution

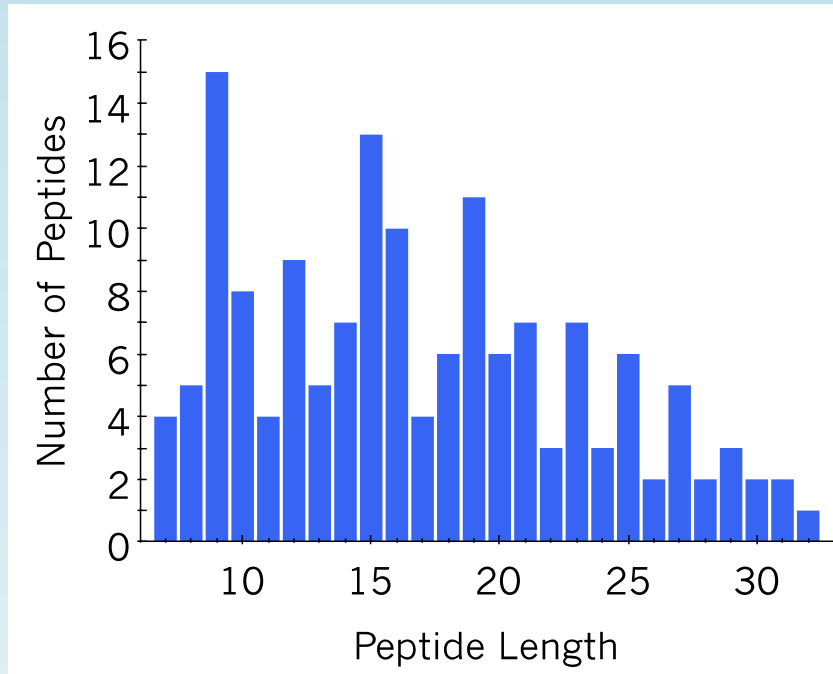
- 96 Serine phosphorylations
 - 26 Threonine phosphorylations
 - 36 Tyrosine phosphorylations
-
- 143 singly phosphorylated
 - 6 doubly phosphorylated
 - 1 triply phosphorylated



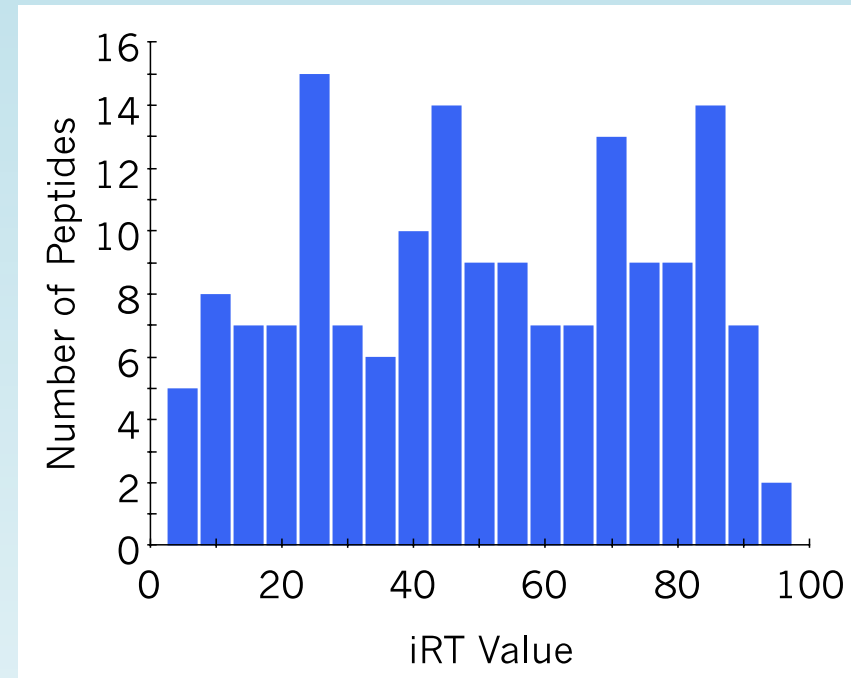
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Generating a Synthetic Phosphopeptide Standard

Peptide Length



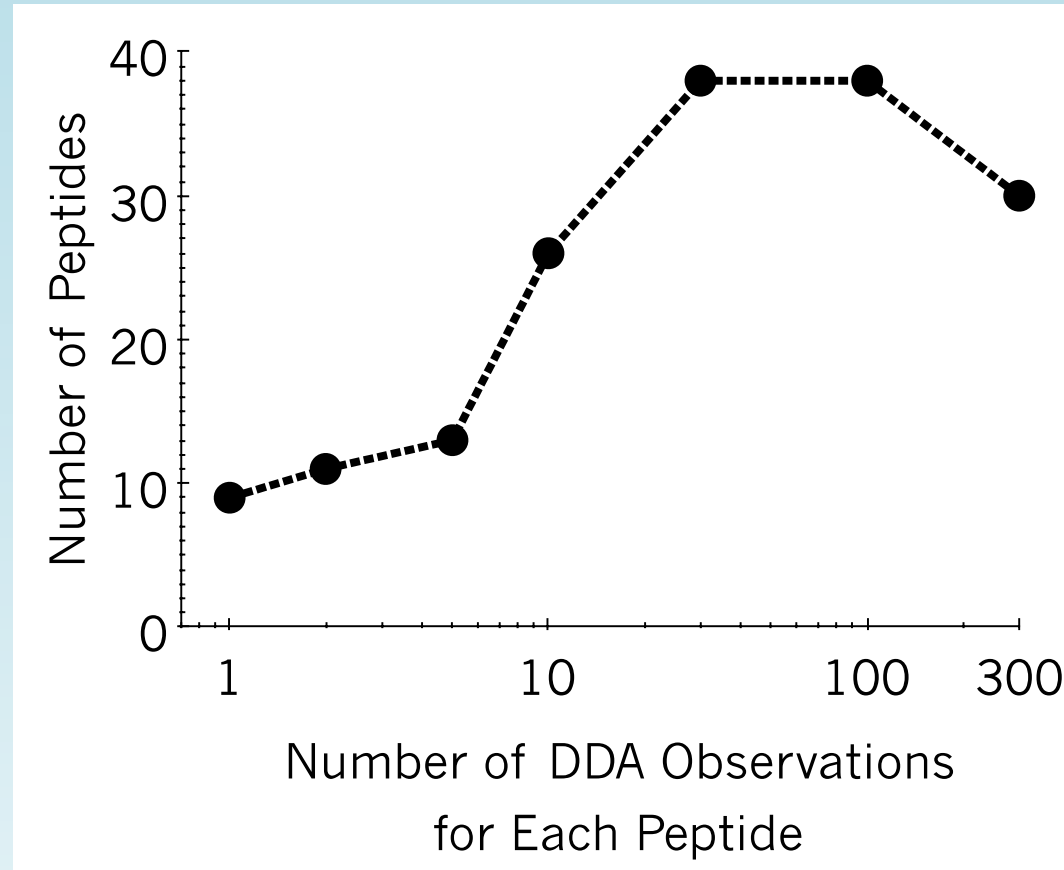
iRT Value





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of DDA Observations for each peptide





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Phosphopeptide Characterization by DIA/PRM

Peptide Type	Count	Percent
good signal	121	85.8%
low signal	6	4.3%
smear	9	6.4%
no signal	5	3.5%



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Future Study

- In late summer/fall 2017, phosphopeptides will be sent out to willing participants.

- Analysis of pure phosphopeptide standard
- Analysis of phosphopeptide standard spiked into lysate
 - TiO₂ or IMAC enrichment



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**Your participation in this study
would be greatly appreciated.**



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New members always welcome!!!!!!