## Brandon Mark Gassaway, PhD

**Assistant Professor** 

E111 BNSN, Brigham Young University

Doctor of Philosophy in Cellular and Molecular Physiology

Provo. UT 84602

801-422-7193

**Education** 

brandon.gassaway@byu.edu

Yale University, New Haven, CT

Thesis Accepted with Distinction	
Certificate of Completion, Medical Research Scholars Program	
ζ	
B.S. Degree in Biochemistry	April 2011
Brigham Young University, Provo, UT	
• Magna Cum Laude with University Honors	
Research Experience	
Assistant Professor	Jan 2024-
Brigham Young University Department of Chemistry & Biochemistry	current
• Research Focus: Functionalizing protein post-translational modifications using thermal stability proteomics and phospho-amino acid orthogonal translation systems	
Post-Doctoral Fellow, Dr. Marcia Haigis	Sept 2021-
Harvard Medical School Department of Cell Biology	Dec 2023
• Research Focus: Applying novel proteomics methods to the investigation of T cell activation and function	Dec 2023
• Techniques: Phosphorylation Integrated Thermal Shift Assay, Primary murine T cell isolation and culture	
Post-Doctoral Fellow, Dr. Steven Gygi	July 2018-
Harvard Medical School Department of Cell Biology	Dec 2023
• Research Focus: Developing experimental methods and bioinformatics analyses to functionalize	

# Ph.D. Candidate, Dr. Jesse Rinehart

proteins and protein phosphorylation

affinity-purification mass spectrometry

Aug 2012-

Yale Department of Cellular and Molecular Physiology, Systems Biology Institute

May 2018

May 2018

• Project: investigating the mechanisms of hepatic insulin resistance by phosphoproteomic analysis

• Techniques: TMT-based quantitative proteomics and phosphoproteomics, protein thermal stability,

• Techniques: quantitative proteomics and phosphoproteomics, systems analysis, enzyme activity, protein purification and analysis, phosphoserine protein production using orthogonal translation systems

# Research Intern, Dr. Lisa Elkin

June 2012-Aug 2012

Bristol-Myers-Squibb Pharmaceuticals

Research Assistant, Dr. Emily Bates

· Project: development of mass spectrometry based assays for assessment of transporter inhibition • Techniques: LC-MS/MS (Agilent RapidFire® platform), small molecule lead evaluation, radioactive

substrate uptake assays

June 2009-

#### BYU Department of Chemistry and Biochemistry

April 2011

- Project: the role of inwardly rectifying potassium channels in *Drosophila* embryonic and adult development
- Techniques: phenotypic analysis, Drosophila genetics, immunohistochemistry

## **Teaching Experience**

Instructor, Dr. Jesse Rinehart and Dr. Terrence Wu	Spring 2018
Yale West Campus Analytical Core Proteomics Bootcamp 2018	
• Developed and presented curriculum for the design, execution, and data analysis of proteomics	
<ul> <li>experiments</li> <li>Designed and carried out hands-on instruction on mass spectrometer instrument maintenance</li> </ul>	
Teaching Fellow 2, Dr. Emile Boulpaep	Winter 2014
Cellular and Molecular Physiology 560	
Designed and presented review material Graded homework and exams	
<b>Teaching Fellow 2,</b> Dr. Stuart Campbell, Dr. W. Mark Saltzman	Fall 2012
Cellular and Molecular Physiology 550	Fall 2013
Designed and presented weekly review sessions using outside material and review activities	
Helped write exam questions	
Graded homework and exams	
wards, Honors, and Fellowships	
Gruber Fellow, Gruber Foundation	Sept 2011
Graduate Research Fellow, National Science Foundation	April 2011
Kagel and Blessing Scholarship, Blessing Family Foundation	Jan 2010
For non-dance majors in performing groups	
Haritaga Sahalanghin, Drigham Voung University	Aug 2005
Heritage Scholarship, Brigham Young University Four years, full tuition	Aug 2005
Tour years, run tuition	
Most Valuable Student Scholarship, National Elks Foundation	April 2005
Most Valuable Student Scholarship, National Elks Foundation	April 2005
	April 2005
Most Valuable Student Scholarship, National Elks Foundation  [Speaker. 70th Annual Conference of the American Society of Mass Spectrometry. Rigorous]	•
<b>Ieetings and Presentations</b> Speaker. 70 <sup>th</sup> Annual Conference of the American Society of Mass Spectrometry. Rigorous Benchmarking and Improvement of Phosphopeptide Identification and False Localization Rates using a	•
Ieetings and Presentations	April 2005 Nov 3, 2021
<b>Ieetings and Presentations</b> Speaker. 70 <sup>th</sup> Annual Conference of the American Society of Mass Spectrometry. Rigorous Benchmarking and Improvement of Phosphopeptide Identification and False Localization Rates using a Proteome-scale Synthetic Human Phosphoserine Library. Philadelphia, PA	Nov 3, 2021
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Poster. Yale Cellular and Molecular Physiology Department Annual Retreat. A Quantitative Oct 27, 2015
Phosphoproteomic Analysis of Acute Hepatic Insulin Resistance Reveals Novel Regulators of Insulin
Signaling. West Haven, CT

· Awarded Best Poster

Speaker. HHMI Med-into-Grad Symposium 2015 Metabolism: Fueling Translational Research. A Quantitative Phosphoproteomic Analysis of Mechanisms of Insulin Resistance Reveals a PKCε-Regulated Network. Boston, MA

Oct 9-10, 2015

Poster. Yale Systems Biology Institute Symposium & Ribbon Cutting Ceremony. A Quantitative Phosphoproteomic Analysis of Mechanisms of Insulin Resistance Reveals a PKC\varepsilon-Regulated Network. West Haven, CT

April 22, 2015

Poster. Yale Cellular and Molecular Physiology Department Annual Retreat. A Quantitative Phosphoproteomics Approach to Mechanisms of Hepatic Insulin Resistance II. Phosphoproteomic Analysis. West Haven, CT

Sep 23, 2014

Poster. MaxQuant Summer School: Computational Mass Spectrometry-Based Proteomics. Quantitative Phosphoproteomic Analysis of Acute Lipid Exposure in Rats Reveals a Direct Regulation of mTOR by AMPK. Bethesda MD.

July 21-25, 2014

Poster. Yale Cellular and Molecular Physiology Department Annual Retreat. Quantitative Phosphoproteomic Analysis of *abl-1* Knockout in *C. elegans* Embryos. West Haven, CT

Oct 28, 2013

Speaker. Mass Spectrometry in Biological Applications and New Technologies. Quantitative Phosphoproteomic Analysis of *abl-1* Knockout in *C. elegans* Embryos. West Haven, CT

June 25, 2013

## **Publications**

- **Gassaway**, **B.M.**, Li, J., Rad, R., Mintseris, J., Mohler, K., Levy, T., Aguiar, M., Beausoleil, S., Paulo, J.A., Rinehart, J., Huttlin, E.L., Gygi, S.P. (2022). A Multi-purpose, Regenerable, Proteome-scale, Human Phosphoserine Resource for Phosphoproteomics. *Nature Methods*, DOI: 10.1038/s41592-022-01638-5
- Ferguson, C. J., Urso, O., Bodrug, T., **Gassaway, B. M.,** Watson, E. R., Prabu, J. R., Lara-Gonzalez, P., Martinez-Chacin, R. C., Wu, D. Y., et al. (2022). APC7 mediates ubiquitin signaling in constitutive heterochromatin in the developing mammalian brain. *Molecular Cell*, 82(1), 90-105.e13.
- Huttlin, E.L., Bruckner, R.J., Navarrete-Perea, J., Cannon, J.R., Baltier, K., Gebreab, F., Gygi, M.P., Thornock, A., Zarraga, G., Tam, S., Szpyt, J., Gassaway, B.M., et al. (2021). Dual proteome-scale networks reveal cell-specific remodeling of the human interactome. Cell 184, 3022-3040.e28.
- Apostolidi, M., Vathiotis, I.A., Muthusamy, V., Gaule, P., **Gassaway, B.M.,** Rimm, D.L., and Rinehart, J. (2021). Targeting Pyruvate Kinase M2 Phosphorylation Reverses Aggressive Cancer Phenotypes. Cancer Research 81, 4346–4359.
- **Gassaway, B.M.,** Paulo, J.A., and Gygi, S.P. (2021). Categorization of Phosphorylation Site Behavior during the Diauxic Shift in Saccharomyces cerevisiae. Journal of Proteome Research 20, 2487–2496.
- Ringel, A.E., Drijvers, J.M., Baker, G.J., Catozzi, A., Garcia-Canaveras, J.C., **Gassaway, B.M.,** Miller, B.C., Juneja, V.R., Nguyen, T.H., Joshi, S., et al. (2020). Obesity Shapes Metabolism in the Tumor Microenvironment to Suppress Anti-Tumor Immunity. Cell 183, 1848-1866 e26.
- Wu, J., Rowart, P., Jouret, F., **Gassaway, B.M.**, Rajendran, V., Rinehart, J., and Caplan, M.J. (2020). Mechanisms involved in AMPK-mediated deposition of tight junction components to the plasma membrane. American Journal of Physiology-Cell Physiology 318, C486–C501.
- Gassaway, B.M., Cardone, R.L., Padyana, A.K., Petersen, M.C., Judd, E.T., Hayes, S., Tong, S., Barber, K.W., Apostolidi, M., Abulizi, A., et al. (2019). Distinct Hepatic PKA and CDK Signaling Pathways Control Activity-Independent Pyruvate Kinase Phosphorylation and Hepatic Glucose Production. Cell Rep 29, 3394-3404.e9.
- Codina, A., Renauer, P.A., Wang, G., Chow, R.D., Park, J.J., Ye, H., Zhang, K., Dong, M.B., Gassaway, B., Ye, L., et al. (2019). Convergent Identification and Interrogation of Tumor-Intrinsic Factors that Modulate Cancer Immunity In Vivo. Cell Syst 8, 136-151 e7.

- Samuel, V.T., Petersen, M.C., **Gassaway, B.M.,** Vatner, D.F., Rinehart, J., and Shulman, G.I. (2019). Considering the Links Between Nonalcoholic Fatty Liver Disease and Insulin Resistance: Revisiting the Role of Protein Kinase C ε. Hepatology 70, 2217–2220.
- Gassaway, B.M., Petersen, M.C., Surovtseva, Y. v, Barber, K.W., Sheetz, J.B., Aerni, H.R., Merkel, J.S., Samuel, V.T., Shulman, G.I., and Rinehart, J. (2018). PKCepsilon contributes to lipid-induced insulin resistance through cross talk with p70S6K and through previously unknown regulators of insulin signaling. Proc Natl Acad Sci U S A 115, E8996–E9005.
- D'Lima, N.G., Khitun, A., Rosenbloom, A.D., Yuan, P., **Gassaway, B.M.,** Barber, K.W., Rinehart, J., and Slavoff, S.A. (2017). Comparative Proteomics Enables Identification of Nonannotated Cold Shock Proteins in E. coli. J Proteome Res 16, 3722–3731.
- Mohler, K., Aerni, H.R., **Gassaway, B.,** Ling, J., Ibba, M., and Rinehart, J. (2017a). MS-READ: Quantitative measurement of amino acid incorporation. Biochim Biophys Acta Gen Subj 1861, 3081–3088.
- Mohler, K., Mann, R., Bullwinkle, T.J., Hopkins, K., Hwang, L., Reynolds, N.M., **Gassaway, B.,** Aerni, H.R., Rinehart, J., Polymenis, M., et al. (2017b). Editing of misaminoacylated tRNA controls the sensitivity of amino acid stress responses in Saccharomyces cerevisiae. Nucleic Acids Res 45, 3985–3996.
- Ferdaus, M.Z., Barber, K.W., Lopez-Cayuqueo, K.I., Terker, A.S., Argaiz, E.R., **Gassaway, B.M.,** Chambrey, R., Gamba, G., Rinehart, J., and McCormick, J.A. (2016). SPAK and OSR1 play essential roles in potassium homeostasis through actions on the distal convoluted tubule. J Physiol 594, 4945–4966.
- Petersen, M.C., Madiraju, A.K., **Gassaway, B.M.,** Marcel, M., Nasiri, A.R., Butrico, G., Marcucci, M.J., Zhang, D., Abulizi, A., Zhang, X.M., et al. (2016). Insulin receptor Thr1160 phosphorylation mediates lipid-induced hepatic insulin resistance. J Clin Invest 126, 4361–4371.
- Rovner, A.J., Haimovich, A.D., Katz, S.R., Li, Z., Grome, M.W., Gassaway, B.M., Amiram, M., Patel, J.R., Gallagher, R.R., Rinehart, J., et al. (2015). Recoded organisms engineered to depend on synthetic amino acids. Nature 518, 89–93.
- Sawyer, N., **Gassaway, B.M.**, Haimovich, A.D., Isaacs, F.J., Rinehart, J., and Regan, L. (2014). Designed phosphoprotein recognition in Escherichia coli. ACS Chem Biol 9, 2502–2507.
- Loffler, M.G., Birkenfeld, A.L., Philbrick, K.M., Belman, J.P., Habtemichael, E.N., Booth, C.J., Castorena, C.M., Choi, C.S., Jornayvaz, F.R., **Gassaway, B.M.,** et al. (2013). Enhanced fasting glucose turnover in mice with disrupted action of TUG protein in skeletal muscle. J Biol Chem 288, 20135–20150.
- Dahal, G.R., Rawson, J., **Gassaway, B.,** Kwok, B., Tong, Y., Ptacek, L.J., and Bates, E. (2012). An inwardly rectifying K+ channel is required for patterning. Development 139, 3653–3664.

## **Personal Information**

- Member, American Society of Mass Spectrometry
- Scoutmaster, Boy Scouts of America
- Performer, BYU International Folk Dance Ensemble
- Eagle Scout
- Conversational Proficiency in Ukrainian and Russian

2012-2015

2009-2011

2001