### **CURRICULUM VITAE**

## The Johns Hopkins University School of Medicine

Hui Zhang September 25, 2022

### DEMOGRAPHIC AND PERSONAL INFORMATION

## **Current Appointments**

2016-present Professor, Department of Pathology, Clinical Chemistry Division, School of Medicine, Johns Hopkins University

2012-present Director, Mass Spectrometry Core Facility, Center for Biomarker Discovery and Translation, Johns Hopkins University

2011-present Joint Professor, Department of Oncology, Johns Hopkins University

2016-present Graduate Faculty of Pathobiology Graduate Training Program

2017-present Graduate Faculty of Chemical and Biomolecular Engineering Graduate Training Program

2020-present Joint Professor, Department of Urology, Johns Hopkins University

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### **Education and Training**

1985-1989 B.S. Plant Biochemistry, Peking University, Beijing, China 1989-1992 M.S. Gene and Protein Engineering, Peking University, Beijing, China 1993-1998 PhD. Biochemistry, University of Pennsylvania, Philadelphia, PA, USA

# **Professional Experience**

PhD. Candidate and research assistant, Biochemistry, University of Pennsylvania, Philadelphia, PA
Product Manager, New England Biolabs, Beverly, MA
Senior Scientist, Cell Signaling Technology, Beverly, MA
Senior Scientist, Institute for Systems Biology, Seattle, WA, USA
Assistant Professor, Department of Pathology, Johns Hopkins University, Baltimore, MD
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### **PUBLICATIONS**:

Please visit updated publications online by Google Scholar or NCBI.

## Original Research [OR]

- Bao Y, Chu R, Han J, **Zhang H**, Pan N, Gu X, Chen Z-L. Cloning and sequencing of Trichosanthin gene and its expression in Escherichia coli and tobacco plant. Sci China. 1993; 36: 669-676.
- Sheng Z-H, **Zhang H**, Barchi RL, Kallen RG. Molecular cloning and functional analysis of the promoter of rat skeletal muscle voltage-sensitive sodium channel subtype 2 (rSkM2): evidence for muscle-specific nuclear protein binding to the core promoter. DNA Cell Biol. 1994; 13: 9-23.
- **Zhang H,** Maldonado MN, Barchi RL, Kallen RG. Dual tandem promoter elements containing CCAC-like motifs from the tetrodotoxin-resistant voltage-sensitive Na<sup>+</sup> channel (rSkM2) gene can independently drive muscle-specific transcription in L6 cells. Gene Expression. 1999; 8: 85-103.
- **Zhang H**, Kolibal S, Vanderkooi JM, Cohen SA, Kallen RG. A carboxyl-terminal α-helical segment in the rat skeletal muscle voltage-dependent Na<sup>+</sup> channel is responsible for its interaction with the amino-terminus. Biochim Biophys Acta. 2000; 1467: 406-418.
- **Zhang H**, Zha X, Tan Y, Hornbeck PV, Mastrangelo AJ, Alessi DR, Polakiewicz RD, Comb MJ. Phospho-protein analysis using antibodies broadly reactive against phosphorylated motifs. J Biol Chem. 2002; 277: 39379-39387.
- Li XJ, **Zhang H**, Ranish JA, Aebersold R. Automated statistical analysis of protein abundance ratios from data generated by stable-isotope dilution and tandem mass spectrometry. Anal Chem. 2003; 75: 6648-6657.
- **Zhang H**, Li XJ, Martin DB, Aebersold R. Identification and quantification of N-linked glycoproteins using hydrazide chemistry, stable isotope labeling and mass spectrometry. Nat Biotechnol. 2003; 21: 660-666. [SI/QI]
- Desiere F, Deutsch EW, Nesvizhskii AI, Mallick P, King N, Eng JK, Aderem A, Boyle R, Brunner E, Donohoe S, Fausto N, Hafen E, Hood L, Katze MG, Kennedy KA, Kregenow F, Lee H, Lin B, Martin D, Ranish JA, Rawlings DJ, Samelson LE, Shiio Y, Watts J, Wollscheid B, Wright ME, Yan W, Yang L, Yi E. C, **Zhang H**, Aebersold R. Integration of peptide sequences obtained by high-throughput mass spectrometry with the human genome. Genome Biol. 2004; 6: R9. [SI/QI]
- **Zhang H**, Yi EC, Li XJ, Mallick P, Spratt KSK, Masselon CD, Camp II DG, Smith RD, Kemp CJ, Aebersold R. High throughput quantitative analysis of serum proteins using glycopeptide capture and LC-MS. Mol Cell Proteomics. 2005; 4: 144-155.
- Liu AY, **Zhang H**, Sorensen CM, Diamond DL. Analysis of prostate cancer using tissue specimens. J Urology. 2005: 173: 73-78.
- Rush J, Moritz A, Lee KA, Goss VL, Guo A, **Zhang H**, Zha X, Polakiewicz RD, Comb MJ. Immunoaffinity profiling of tyrosine phosphorylation in cancer cells. Nat Biotechnol. 2005; 23: 94-101.
- Chen J, Saeki F, Wiley BJ. Cang, H, Li Z, Cobb MJ, Au L, **Zhang H**, Kimmey MB, Li XD, Xia Y. Bioconjugated gold nanocages and evaluation of their potential for optical imaging and thermal therapeutic applications. Nano Lett. 2005; 5: 473-477.
- Li XJ, Yi EC, Kemp CJ, **Zhang H**, Aebersold RA software suite for the generation and comparison of peptide arrays from sets of data collected by liquid chromatography-mass spectrometry. Mol Cell Proteomics. 2005; 4: 1328-1340.
- Pan S, **Zhang H**, Rush J, Eng J, Zhang N, Patterson D, Comb MJ, Aebersold R. High-throughput proteome-screening for biomarker detection. Mol Cell Proteomics. 2005; 4: 182-190.
- Deutsch EW, Eng JK, **Zhang H**, King NL, Nesvizhskii AI, Lin B, Lee H, Yi EC, Ossola R, Aebersold R. Human plasma PeptideAtlas. Proteomics. 2005; 5: 3497-3450.
- **Zhang H**, Loriaux P, Eng J, Campbell D, Keller A, Moss P, Bonneau R, Zhang N, Zhou Y, Wollscheid B, Cooke K, Yi EC, Lee H, Peskind ER, Zhang J, Smith RD, Aebersold R. UniPep, a database for human N-linked glycosites: A Resource for Biomarker Discovery. Genome Biol. 2006; 7: R73.

- Whiteaker JR, Zhang H, Eng JK, Fang R, Piening BD, Feng L, Lorentzen TD, Schoenherr RM, Keane JF, Holzman T, Fitzgibbon M, Lin C, **Zhang H**, Cooke K, Liu T, Camp II DG, Anderson L, Watts J, Smith RD, McIntosh MW, Paulovich A. Head-to-Head Comparison of Serum Fractionation Techniques. J Proteome Res. 2007; 6: 828-36.
- Wang P, Tang H, Fitzgibbon MP, McIntosh M, Coram M, **Zhang H**, Yi E, Aebersold R. A statistical method for chromatographic alignment of LC-MS data. Biostatistics. 2007; 8: 357-67.
- **Zhang H,** Chan DW. Cancer biomarker discovery in plasma using a tissue-targeted proteomic approach. Cancer Epidem Biomar. 2007; 16:1916-1917.
- Chen J, Wang D, Xi J, Au L, Siekkinen A, Warsen A, Li Z-Y, **Zhang H**, Xia Y, Li X. Immuno gold nanocages with tailored optical properties for targeted photothermal destruction of cancer cells. Nano Lett. 2007; 7:1318-22
- **Zhang H**, Liu AY, Loriaux P, Wollscheid B, Zhou Y, Watts JD, Aebersold R. Mass spectrometric detection of tissue-derived proteins in blood. Mole Cell Proteomics. 2007; 6: 64-71.
- Zhou Y, Aebersold R, Zhang H. Isolation of N-linked glycopeptides from plasma. Anal Chem. 2007;79:5826-5837.
- Stokes MP, Rush J, Macneill J, Ren JM, Sprott K, Nardone J, Yang V, Beausoleil SA, Gygi SP, Livingstone M, **Zhang H**, Polakiewicz RD, Comb MJ. Profiling of UV-induced ATM/ATR signaling pathways. Proc Natl Acad Sci USA. 2007; 104: 19855-19860.
- Tian Y, Kelly-Spratt KS, Kemp KJ, **Zhang H.** Identification of glycoproteins from mouse skin tumors and plasma. Clin Proteomics. 2008; 4: 117-136.
- Hwang D; Zhang N, Lee H, Yi EC, **Zhang H**, Lee IY, Hood L, Aebersold R. MS-BID: a Java package for label-free LC-MS based comparative proteomic analysis. Bioinformatics. 2008; 24: 2641-2642.
- Zou Z, Ibisate M, Zhou Y, Aebersold R, Xia Y, **Zhang H.** Synthesis and evaluation of superparamagnetic silica particles for extraction of glycopeptides in the microtiter plate format. Anal Chem. 2008; 80: 1228-1234.
- Li Y, Sokoll LJ, Barker PE, **Zhang H,** Chan DW. Mass spectrometric identification of proteotypic peptides from clinically used tumor markers. Clin Proteomics. 2008; 2: 58-66.
- Letarte S, Brusniak MY, Campbell D, Eddes J, Kemp CJ, Lau H, Mueller L, Schmidt A, Shannon P, Kelly-Spratt KS, Vitek O, **Zhang H**, Aebersold R, Watts JD. Differential plasma glycoproteome of p19ARF skin cancer mouse model using the Corra label-free LC-MS proteomics platform. Clin Proteomics. 2008; 4: 105-116.
- Meany DL (Li D), Zhang Z, Sokoll LJ, **Zhang H,** Chan DW. Glycoproteomics for prostate cancer detection: changes in serum PSA glycosylation patterns. J Proteome Res. 2009; 8: 613-619.
- Tian Y, Gurley K, Meany DL (Li D), Kemp CJ, **Zhang H.** N-linked glycoproteomic analysis of formalin-fixed and paraffinembedded tissues. J Proteome Res. 2009; 8: 1657-1662.
- Li Y, Sokoll L, Rush J, Zou N, Chan DW, **Zhang H.** Targeted detection of prostate cancer proteins in serum using heavy peptide standards and MALDI-TOF/TOF. Proteom Clin Appl. 2009; 3: 597-608.
- Tian Y, Tan A, Sun X, Olson MT, Xie Z, Jinawath N, Chan DW, Shih I, Zhang Z, **Zhang H.** Quantitative proteomic analysis of ovarian cancer cells identified mitochondrial proteins associated with Paclitaxel treatment. Proteom Clin Appl. 2009; 3: 1288-1295.
- Ueda SM, Yap KL, Davidson B, Tian Y, Murthy V, Wang T-L, Visvanathan K, Kuhajda FP, Bristow RE, **Zhang H**, Shih I-M. Expression of fatty acid synthase depends on NAC1 and is associated with recurrent ovarian serous carcinomas. J Oncol. 2010; 2010: 285191.
- True LD, **Zhang H**, Yie M, Huang C-Y, Nelson PS, von Haller PD, Tjoelker LW, Kim J-S, Qian W-J, Smith RD, Ellis WJ, Liebeskind ES, Liu AY. CD90/Thy1 is over-expressed in prostate cancer-associated fibroblasts and could serve as a cancer biomarker. Modern Pathol. 2010; 23: 1346-1356.

- Tian Y, Kelly-Spratt KS, Kemp KJ, **Zhang H.** Mapping tissue-specific expression of extracellular proteome. J Proteome Res. 2010; 9: 5837-5847.
- Prakash A, Rezai T, Krastins B, Sarracino D, Athanas M, Russo P, Ross MM, **Zhang H**, Tian Y, Kulasingam V, Drabovich AP, Smith C, Batruch I, Liotta E, Petricoin EF, Diamandis EP, Chan DW, Lopez MF. A platform for establishing inter-laboratory reproducibility of selected reaction monitoring-based mass spectrometry peptide assays. J Proteome Res. 2010; 9: 6678–6688.
- Li Y, Tian Y, Rezai T, Prakash A, Lopez MF, Chan DW, **Zhang H.** Simultaneous analysis of glycosylated and sialylated PSA reveals differential distribution of glycosylated PSA isoforms in prostate cancer tissues. Anal Chem. 2011; 83: 240–245.
- Meany DL (li D), Hackler L, **Zhang H**, Chan DW. Tyramide signal amplification for antibody-overlay lectin micro-array: a strategy to improve the sensitivity of targeted glycan profiling. J Proteome Res. 2011; 10: 1425–1431.
- Baycin-Hizal D, Tian Y, Akan I, Palter K, **Zhang H,** Betenbaugh M. GlycoFly database for identification of Drosophila glycoproteomics. J Proteome Res. 2011; 10: 2777-2784.
- Baycin-Hizal D, Tian Y, Akan I, Jacobson E, Clark D, Wu A, Jampol R, Palter K, Betenbaugh M, **Zhang H.** GlycoFish: A Database of Zebrafish N-linked Glycoproteins Identified using SPEG method coupled with LC/MS. Anal Chem. 2011; 83: 5296-5303.
- Farrah T, Deutsch EW, Omenn GS, Campbell DS, Sun Z, Bletz J, Mallick P, Katz JE, Malmstroem J, Ossola R, Watts JD, Lin B, **Zhang H**, Moritz RL, Aebersold R. A High-confidence human plasma proteome reference set with estimated concentrations in PeptideAtlas. Mol Cell Proteomics. 2011; 10: M110.006353.
- Tian Y, Bova GS, **Zhang H**. Quantitative Glycoproteomic Analysis of Optimal Cutting Temperature-Embedded Frozen Tissues Identifying Glycoproteins Associated with Aggressive Prostate Cancer. Anal Chem. 2011; 83: 7013–7019.
- Hyung S, Lee M, Yu J, Shin B, Jung H, Park J, Han W, Lee K, Moon H, **Zhang H**, Aebersold R, Hwang D, Lee S, Yu M, Noh D. A serum protein profile predictive of the resistance to neoadjuvant chemotherapy in advanced breast cancers. Mol Cell Proteomics. **2011**; 10: M111.011023.
- Tian Y, Yao Z, Roden R, **Zhang H.** Identification of glycoproteins associated with different histological subtypes of ovarian tumors using quantitative glycoproteomics. Proteomics. 2011; 11: 4677-4687.
- Li Y, Tao SC, Bova GS, Liu AY, Chan DW, Zhu H, **Zhang H**. Detection and verification of glycosylation patterns of glycoproteins from clinical specimens using lectin microarrays and lectin-based immunosorbent assays. Anal Chem. 2011; 83: 8509-8516.
- Yang S, **Zhang H**. Glycan Analysis by reversible reaction to hydrazide beads and mass spectrometry. Anal Chem. 2012; 84: 2232–2238. [SI/QI]
- Tian Y, Esteva FJ, Song J, **Zhang H**. Altered expression of sialylated glycoproteins in breast cancer using hydrazide chemistry and mass spectrometry. Mol Cell Proteomics. 2012; 11: M111.011403.
- Almaraz RT, Tian Y, Bhattarcharya R, Tan E, Chen SH, Dallas MR, Chen L, Zhang Z, **Zhang H**, Konstantopoulos K, Yarema KJ. Metabolic flux increases glycoprotein sialylation: implications for cell adhesion and cancer metastasis. Mole Cell Proteomics. 2012; 11: M112.017558.
- Prakash A, Rezai T, Krastins B, Sarracino D, Athanas M, Russo P, **Zhang H**, Tian Y, Li Y, Kulasingam V, Drabovich A, Smith CR, Batruch I, Oran PE, Fredolini C, Luchini A, Liotta L, Petricoin E, Diamandis EP, Chan DW, Nelson R, Lopez AF. Inter-laboratory reproducibility of SRM assays using multiple upfront analyte enrichment strategies. J Proteome Res. 2012; 11: 3986-3995.
- Toghi Eshghi S, Li X, **Zhang H.** Targeted analyte detection by standard addition improves detection limit in MALDI mass spectrometry. Anal Chem. 2012; 84: 7626–7632.

- Baycin-Hizal D, Tabb D, Chaerkady R, Chen L, Lewis N, Nagarajan H, Sarkaria V, Kumar A, Wolozny D, Colao J, Jacobson E, Tian Y, O'Malley B, Cole B, Palsson B, **Zhang H**, Betenbaugh M. Proteomic and glycoproteomic analysis of Chinese Hamster Ovary cells. J Proteome Res. 2012; 11: 5265-5276.
- Li D, Chiu J, Chen J, **Zhang H**, Chan DW. Integrated analyses of proteins and their glycans in a magnetic beads based multiplex assay format. Clin Chem. 2013; 59: 315-324.
- Li QK, Gabrielson E, Askin F, Chan DW, **Zhang H.** Glycoproteomics using so-called 'fluid-biopsy'specimens in the discovery of lung cancer biomarkers. Promise and challenge. Proteomics Clin Appl, 2013; 7: 55-69.
- Marzinke MA, Choi CH, Chen L, Shih I-M, Chan DW, **Zhang H**. Proteomic Analysis of Temporally Stimulated Ovarian Cancer Cells for Biomarker Discovery. Mol Cell Proteomics. 2013; 12: 356-368.
- Shah P, Yang S, Sun S, Aiyetan P, Yarema K, **Zhang H.** Mass Spectrometric Analysis of Sialylated Glycans Using Solid Phase Labeling of Sialc Acids. Anal Chem. 2013; 85: 3606-3613. [SI/QI]
- Mansuy-Aubert V, Zhou QL, Xie X, Gong Z, Huang JY, Khan AR, Aubert G, Candelaria K, Thomas S, Shin DJ, Booth S, Baig SM, Bilal A, Hwang D, **Zhang H**, Lovell-Badge R, Smith SR, Awan FR, Jiang ZY. Imbalance between Neutrophil Elastase and its Inhibitor α1-Antitrypsin in Obesity Alters Insulin Sensitivity, Inflammation, and Energy Expenditure. Cell Metab. 2013; 17: 534-548.
- Yang S, Li Y, Shah P, **Zhang H**. Glycomic analysis using glycoprotein immobilization for glycan extraction. Anal Chem. 2013; 85: 5555-5561.
- Li QK, Shah P, Li Y, Aiyetan P, Chen J, Yung R, Molena D, Gabrielson E, Askin F, Chan DW, **Zhang H**. Glycoproteomic Analysis of Bronchoalveolar Lavage (BAL) Fluid Identifies Tumor-associated Glycoproteins from Lung Adenocarcinoma. J Proteome Res. 2013; 12: 3689-3696.
- Chen J, Xi J, Tian Y, Bova GS, **Zhang H.** Identification, prioritization and evaluation of glycoproteins for aggressive prostate cancer using quantitative glycoproteomics and antibody-based assays on tissue specimens. Proteomics. 2013; 13: 2268-2277.
- Yang S, Yuan W, Yang W, Zhou J, Harlan R, Edwards JL, Li S, **Zhang H**. Glycan Analysis by Isobaric Aldehyde Reactive Tags and Mass Spectrometry. Anal Chem. 2013; 85: 8188-8195. [SI/QI]
- Li D, Chiu H, **Zhang H**, Chan DW. Analysis of serum protein glycosylation by a differential lectin immunosorbent assay. Clin Proteomics. 2013; 10: 1-9.
- Chen J, Shah P, **Zhang H.** Solid phase extraction of N-linked glycopeptides using hydrazide tips. Anal Chem. 2013; 85: 10670-10674.
- Yang S, Toghi Eshghi S, Chiu H, DeVoe D, **Zhang H**. Glycomic Analysis by Glycoprotein Immobilization for Glycan Extraction and Liquid Chromatography on Microfluidic Chip. Anal Chem. 2013; 85: 10117-10125.
- Li Y, Wang X, Ao M, Gabrielson E, Askin F, **Zhang H**, Li QK. Aberrant Mucin5B expression in lung adenocarcinomas detected by iTRAQ labeling quantitative proteomics and immunohistochemistry. Clin Proteomics. 2013; 10: 15-24.
- Sun S, Zhang B, Aiyetan P, Zhou J, Shah P, Yang W, Levine DA, Zhang Z, Chan DW, **Zhang H**. Analysis of N-glycoproteins using genomic N-glycosite prediction. J Proteome Res. 2013; 12: 5609-5615.
- Chen J, Toghi Eshghi S, Bova GS, Li QK, Li X, **Zhang H.** Epithelium percentage estimation facilitates epithelial protein measurement in tissue specimens in cancer biomarker discovery. Clin Proteomics. 2013;10: 1-11.
- Sun S, Zhou J, Yang W, **Zhang H**. Inhibition of protein carbamylation in urea solution using ammonium containing buffers. Anal Biochem. 2014; 446: 76-81.
- Yang W, Zhou J, Chen L, Ao M, Sun S, Aiyetan P, Simmons A, **Zhang H**, Jackson JB. Glycoproteomic analysis identifies human glycoproteins secreted from HIV latently infected T cells and reveals their presence in HIV+ plasma. Clin

- Ao MH, **Zhang H**, Sakowski L, Sharma R, Illei PB, Gabrielson E, Askin F, Li QK. The Utility of a Novel Triple Marker (combination of TTF1, Napsin A and p40) in the Subclassification of Non-small Cell Lung Cancer (NSCLC). Human Pathol. 2014; 45: 926-934.
- Liu Y, Chen J, Sethi A, Li QK, Chen L, Collins B, Gillet LCJ, Wollscheid B, **Zhang H**, Aebersold R. Glycoproteomic Analysis of Prostate Cancer Tissues by Glycoproteomic analysis of prostate cancer tissues by SWATH mass spectrometry discovers N-acylethanolamine acid amidase and protein tyrosine kinase 7 as signatures for tumor aggressiveness. Mol Cell Proteomics. 2014; 13: 1753-1768.
- Aiyetan P, Zhang B, Chen L, Zhang Z, **Zhang H.** M2Lite: An Open-source, Light-weight, Pluggable and Fast Proteome Discoverer MSF to mzIdentML Tool. J Bioinformatics. 2014; 1: 40-49.
- Wang X, Chen J, Li QK, Peskoe SB. Zhang B, Choi C, Platz EA and **Zhang H.** Overexpression of alpha (1,6) fucosyltransferase associated with aggressive prostate cancer. Glycobiology. 2014; 24: 935-944.
- Aiyetan P, Zhang B, Zhang H. XGlycScan: An open-source software for N-linked glycosite assignment, quantification and quality assessment of data from mass spectrometry-based glycoproteomic analysis. MOJ Proteomics and Bioinformatics. 2014; 1: 4-16.
- Baycin-Hizal D, Gottschalk A, Jacobson E, Mai S, Wolozny D, **Zhang H**, Krag SS, Betenbaugh MJ. Physiologic and pathophysiologic consequences of altered sialylation and glycosylation on ion channel function. Biochem Bioph Res Co. 2014; 453: 243-253.
- Tian Y, Koganti T, Yao Z, Cannon P, Shah P, Pietrovito L, Modesti A, Aiyetan P, DeLeon-Pennell K, Ma Y, Halade GV, Hicks C, **Zhang H,** Lindsey ML. Cardiac extracellular proteome profiling and membrane topology analysis using glycoproteomics. Proteom Clin Appl. 2014; 8: 595-602.
- Yang W, Shah P, Toghi Eshghi S, Yang S, Sun S. Ao M, Rubin A, Jackson, JB, **Zhang H.** Glycoform-analysis of recombinant and human deficiency virus envelope protein gp120 via higher energy C-trap dissociation and spectral-aligning strategy. Anal Chem. 2014; 86: 6959-6967.
- Toghi Eshghi S, Yang S, Wang X, Shah P, Li X, **Zhang H.** Imaging of N-Linked Glycans from Formalin-Fixed Paraffin-Embedded Tissue Sections Using MALDI Mass Spectrometry. ACS Chem Biol. 2014; 9: 2149-2156. [SI/QI]
- Rodriguez E, Chen L, Ao M, Geddes S, Gabrielson E, Askin F, **Zhang H**, Li QK. Expressions of transcript factors SALL4 and OCT4 in a subset of non-small cell lung carcinomas (NSCLC). Transl Respiratory Med. 2014; 2: 1-8.
- Zhang B. et al. & **NCI CPTAC Investigators**. Proteogenomic characterization of human colon and rectal cancer. Nature. 2014; 513: 382-387.
- Yang W, Laeyendecker O, Wendel SK, Zhang B, Sun S, Zhou J, Ao M, Moore RO, Jackson JB, **Zhang H**. Glycoproteomic Study Reveals Altered Glycoproteins from Plasma of HIV Elite Suppressors. Theronostics. 2014; 4: 1153-1163.
- Harlan R, **Zhang H**. Targeted proteomics: a bridge between discovery and validation. Expert Rev Proteomics. 2014; 11: 657-661.
- Yang S, Chen L, Sun S, Shah P, Yang W, Zhang B, Zhang Z, Chan DW, Kass DA, van Eyk JE, **Zhang H.** Glycoproteins identified from heart failure and treatment models. Proteomics. 2015; 15: 567-579.
- Shah P, Zhang B, Choi C, Yang S, Zhou J, Harlan R, Tian Y, Zhang Z, Chan DW, **Zhang H.** Tissue Proteomics Using Chemical Immobilization and Mass Spectrometry. Anal Biochem. 2015; 469: 27-33.
- Li QK, Chen L, Ao M, Chiu JH, Zhang Z, **Zhang H,** Chan DW. Serum Fucosylated Prostate-specific Antigen (PSA) Improves the Differentiation of Aggressive from Non-aggressive Prostate Cancers. Theranostics. 2015; 5: 267-276.
- Padoan A, Basso D, Malfa MA, Zambon C, Aiyetan P, Zhang H, Chiara AD, Pavanello G, Bellocco R, Chan DW, Plebani

- M. Reproducibility in urine peptidome profiling using MALDI-TOF. Proteomics. 2015; 15: 1476-1485.
- Tian Y, Choi CH, Li QK, Rahmatpanah FB, Chen X, Kim SR, Veltri R, Chia D, Zhang Z, Mercola D, **Zhang H.** Overexpression of periostin in stroma positively associated with aggressive prostate cancer. PloS one. 2015; 10: e0130333 1-10.
- Tian Y, Almaraz RT, Choi CH, Li QK, Saeui C, Li D, Shah P, Bhattacharya R, Yarema KJ, **Zhang H**. Identification of sialylated glycoproteins from metabolically oligosaccharide engineered pancreatic cells. Clin Proteomics. 2015; 12: 11-23.
- Toghi Eshghi S, Shah P, Yang W, Li X, **Zhang H**. GPQuest: A Spectral Library Matching Algorithm for Site-Specific Assignment of Spectra from Tandem Mass Spectrometric Analysis of Intact Glycopeptides. Anal Chem. 2015; 87: 5181-5188. [SI/QI]
- Li Y, Shah P, DeMarzo A, VanEyk J, Li Q, Chan DW, **Zhang H**. Identification of glycopeptides containing specific glycan motifs using a Lectin-Chemical Immobilization method. Anal Chem. 2015; 87: 4683-4687.
- Thomas S, **Zhang H**, Cotter R. Application of quantitative proteomics to the integrated analysis of the ubiquitylated and global proteomes of xenograft tumor tissues. Clin Proteomics. 2015; 12: 1-15.
- Yin B, Gao Y, Chung A, Yang S, Blake E, Kildegaard HF, Tang J, Andersen MR, **Zhang H**, Betenbaugh MJ. Glycoengineering of Chinese hamster ovary cells for enhanced erythropoietin N-glycan branching and sialylation. Biotechnol Bioeng. 2015; 112: 2343-2351.
- Sun S, **Zhang H.** Large-scale measurement of absolute protein glycosylation stoichiometry. Anal Chem. 2015; 87: 6479-6482.
- Thomas S, Harlan R, Chen J, Aiyetan P, Liu Y, Sokoll LJ, Aebersold R, Chan DW, **Zhang H**. Multiplexed targeted mass spectrometry-based assays for the quantification of N-linked glycosite-containing peptides in serum. Anal Chem., 2015; 87: 10830–10838.
- Shah P, Wang X, Yang W, Toghi Eshghi S, Sun S, Höti N, Pasay J, Rubin A, **Zhang H.** Integrated proteomic and glycoproteomic analyses of prostate cancer cells reveals glycoprotein alteration in protein abundance and glycosylation. Mol Cell Proteomics. 2015; 14: 2753-2763. [SI/QI]
- Yang S, Mishra S, Chen L, Zhou J, Chan D, Chatterjee S, **Zhang H.** Integrated glycoprotein immobilization method for glycopeptide and glycan analysis of cardiac hypertrophy. Anal Chem. 2015; 87:9671-9678.
- Tabb DL, Wang X, Carr SA, Clauser KR, Mertins P, Chambers MC, Holman JD, Wang J, Zhang B, Zimmerman LJ, Chen X, Gunawardena HP, Davies SR, Ellis MJ, Li S, Townsend RR, Boja ES, Ketchum KA, Kinsinger CR, Mesri M, Rodriguez H, Liu T, Kim S, McDermott JE, Payne SH, Petyuk VA, Rodland KD, Smith RD, Yang F, Chan DW, Zhang B, **Zhang H**, Zhang Z, Zhou J, Liebler DC. Reproducibility of Differential Proteomic Technologies in CPTAC Fractionated Xenografts. J Proteome Res. 2016; 15: 691-706.
- Zhu G, Liu Z, Epstein JI, Davis C, Christudass CS, Carter HB, Landis P, **Zhang H**, Chung J, Hewitt SM, Miller MC, Veltri RW. A novel quantitative multiplex tissue immunoblotting for biomarkers predicts a prostate cancer aggressive phenotype. Cancer EpHuidem Biomar. 2015; 24: 1864-1872.
- Kumar A, Baycin-Hizal D, Wolozny D, Ebdrup Pedersen L, Lewis NE, Heffner K, Chaerkady R, Cole RN, Shiloach J, **Zhang H**, Bowen MA, Betenbaugh MJ. Elucidation of the CHO super-ome (CHO-SO) by proteoinformatics. J Proteome Res. 2015; 14: 4687-4703.
- Yang S, Wang M, Chen L, Yin B, Song G, Turko IV, Phinney KW, Betenbaugh MJ, **Zhang H**, Li S. QUANTITY: An Isobaric Tag for Quantitative Glycomics. Sci Rep. 2015; doi: 10.1038/srep17585.
- Sun S, Zhang H. Identification and validation of atypical N-glycosylation sites. Anal Chem. 2015; 87: 11948-11951.
- Aiyetan P, Thomas S, Zhang Z, Zhang H. MRMPlus: an open source quality control and assessment tool for SRM/MRM

- Ruggles KV, Tang Z, Wang X, Grover H, Askenazi M, Teubl J, Cao S, McLellan MD, Clauser KR, Tabb DL, Mertins P, Slebos R, Erdmann-Gilmore P, Li S, Gunawardena HP, Xie L, Liu T, Zhou J, Sun S, Hoadley KA, Perou CM, Chen X, Davies SR, Maher CA, Kinsinger CR, Rodland KD, **Zhang H**, Zhang Z, Ding L, Townsend RR, Rodriguez H, Chan D, Smith RD, Liebler DC, Carr SA, Payne S, Ellis MJ, Fenyo D. An analysis of the sensitivity of proteogenomic mapping of somatic mutations and novel splicing events in cancer. Molecular & Cellular Proteomics. 2016; 15: 1060-1071.
- Hoofnagle AN, Whiteaker JR, Carr SA, Kuhn E, Liu T, Massoni SA, Thomas SN, Townsend RR, Zimmerman LJ, Boja E, Chen J, Crimmins DL, Davies SR, Gao Y, Hiltke TR, Ketchum KA, Kinsinger CR, Mesri M, Meyer MR, Qian WJ, Schoenherr RM, Scott MG, Shi T, Whiteley GR, Wrobel JA, Wu C, Ackermann BL, Aebersold R, Barnidge DR, Bunk DM, Clarke N, Fishman JB, Grant RP, Kusebauch U, Kushnir MM, Lowenthal MS, Moritz RL, Neubert H, Patterson SD, Rockwood AL, Rogers J, Singh RJ, Van Eyk JE, Wong SH, Zhang S, Chan DW, Chen X, Ellis MJ, Liebler DC, Rodland KD, Rodriguez H, Smith RD, Zhang Z, **Zhang H**, Paulovich AG. Recommendations for the Generation, Quantification, Storage, and Handling of Peptides Used for Mass Spectrometry—Based Assays. Clin Chem. 2016; 62: 48-69.
- Clark D, Mei Y, Sun S, **Zhang H**, Yang A, Mao L. Glycoproteomic approach identifies KRAS as a positive regulator of CREG1 in non-small cell lung cancer cells. Theranostics. 2016; 6: 65-77.
- Whiteaker JR, Halusa GN, Hoofnagle AN, Sharma W, MacLean B, Yan P, Wrobel JA, Kennedy J, Mani DR, Zimmerman LJ, Meyer MR, Mesri M, Boja E, Carr SA, Chan DW, Chen X, Chen J, Davies SR, Ellis MJC, Fenyö D, Hiltke T, Ketchum KA, Kinsinger C, Kuhn E, Liebler DC, Liu T, Loss M, MacCoss MJ, Qian W, Rivers R, Rodland KD, Ruggles KV, Scott MG, Smith RD, Thomas S, Townsend RR, Whiteley G, Wu C, **Zhang H**, Zhang Z, Rodriguez H, Paulovich AG. Using the CPTAC Assay Portal to identify and implement highly characterized targeted proteomics assays. Quantitative Proteomics by Mass Spectrometry. 2016; 223-226.
- Sun S, Shah P, Toghi Eshghi S, Yang W, Trikannad N, Yang S, Chen L, Aiyetan P, Höti N, Zhang Z, Chan DW, **Zhang H.** Comprehensive analysis of protein glycosylation by solid-phase extraction of N-linked glycans and glycosite-containing peptides. Nat Biotechnol. 2016; 34: 84-88. [SI/QI]
- DeLeon-Pennell KY, Tian Y, Zhang B, Cates CA, Padmanabhan Iyer R, Cannon P, Shah P, Aiyetan P, Halade GV, Ma Y, Flynn E, Zhang Z, Jin YF, **Zhang H**, Lindsey ML. CD36 Is a Matrix Metalloproteinase-9 Substrate that Stimulates Neutrophil Apoptosis and Removal during Cardiac Remodeling Circ Cardiovasc Genet. 2016; 9: 14-25.
- Lilo MT, Allison D, Wang Y, Ao M, Gabrielson E, Geddes S, **Zhang H**, Askin F, Li QL. Expression of P40 and P63 in lung cancers using fine needle aspiration cases. Understanding clinical pitfalls and limitations. J of the American Society of Cytopathology. 2016; 5: 123-132.
- Yang W, Jackson B, **Zhang H.** Identification of glycoproteins associated with HIV latently infected cells using quantitative glycoproteomics. Proteomics. 2016; 16: 1825-1948. Featured as the front cover of the issue.
- Zhang H, Liu T, Zhang Z, Payne SH, Zhang B, McDermott JE, Zhou J, Petyuk VA, Chen L, Ray D, Sun S, Yang F, Chen L, Wang J, Shah P, Cha S-W, Aiyetan P, Woo S, Tian Y, Gritsenko MA, Choi C, Monroe ME, Thomas S, Moore RJ, Yu K-H, Tabb DL, Fenyo D, Bafna V, Wang Y, Rodriguez H, Boja ES, Hiltke T, Rivers RC, Sokoll L, Zhu H, Shih I-M, Pandey A, Zhang B, Snyder MP, Levine DA, Smith RD, Chan DW, Rodland KD, and the NCI CPTAC Investigators. Integrated Proteogenomic Characterization of Human High-Grade Serous Ovarian Cancer. Cell. 2016; 166: 755-765. [SI/QI]
- Yu K, Levine DA, **Zhang H**, Chan DW, Zhang Z, Snyder M. Predicting Ovarian Cancer Patients' Clinical Response to Platinum-based Chemotherapy by their Tumor Proteomic Signatures. J Proteome Res. 2016; 15: 2455-2465.
- Mertins P. et al and **NCI CPTAC Investigators**. Proteogenomics connects somatic mutations to signaling in breast cancer. Nature. 2016; 534; 55-62.
- Jia X, Chen J, Sun S, Yang W, Yang S, Shah P, Hoti N, Veltri R, **Zhang H**. Detection of aggressive prostate cancer-associated glycoproteins in urine using glycoproteomics and mass spectrometry. Proteomics. 2016; 16: 2989-2996.

- Toghi Eshghi S, Yang W, Hu Y, Shah P, Sun S, Li X, **Zhang H**. Classification of Tandem Mass Spectra for Identification of N- and O-linked Glycopeptides. Sci Rep. 2016; 6: 37189.
- Shah P, Yang W, Sun S, Pasay J, Faraday N, **Zhang H.** Platelet glycoproteins associated with aspirin-treatment upon platelet activation. Proteomics. 2017; 17 (6): 1600199.
- Collins BC, Hunter CL, Liu Y, Schilling B, Rosenberger G, Bader SL, Chan DW, Gibson BW, Gingras AC, Held JM, Hirayama-Kurogi M, Hou G, Krisp CK, Larsen B, Lin L, Liu S, Molloy MP, Moritz RL, Ohtsuki S, Schlapbach R, Selevsek N, Thomas SN, Tzeng S, **Zhang H**, Aebersold R. Multi-laboratory assessment of reproducibility, qualitative and quantitative performance of SWATH-mass spectrometry. Nat Commun. 2017; 8: 291.
- Wang J, Ma Z, Carr SA, **Zhang H**, Zhang Z, Chan DW, Ellis MJC, Townsend RR, Smith RD, McDermott JE, Chen X, Paulovich AG, Boja ES, Mesri M, Kinsinger CR, Rodriguez H, Rodland KD, Liebler DC, Zhang B. Proteome profiling outperforms transcriptome profiling for co-expression based gene function prediction. Mol Cell Proteomics. 2017; 16: 121-134.
- Yang S, Höti N, Yang W, Liu Y, Chen L, Li S, **Zhang H.** Simultaneous analyses of N-linked and O-linked glycans of ovarian cancer cells using solid-phase chemoenzymatic method. Clinical Proteomics. 2017; 14: 3.
- Chung C, Wang Q, Yang S, Yin B, **Zhang H**, Betenbaugh M. Integrated Genome and Protein Editing Swaps α-2, 6 Sialylation for α-2, 3 Sialic Acid on Recombinant Antibodies from CHO. Biotech J. 2017; 12 (2): 1600502
- Höti N, Shah P, Hu Y, Yang S, **Zhang H**. Proteomics analyses of prostate cancer cells reveals cellular pathways associated with androgen resistance. Proteomics. 2017; 17 (6): 1600228.
- Zarif JC, Yang W, Hernandez JR, **Zhang H**, Pienta KJ. The identification of macrophage enriched glycoproteins using glycoproteomics. Mol Cell Proteomics. 2017; 16 (6), 1029-1037.
- Yang S, Zhang L, Thomas SN, Hu Y, Li S, Cipollo JF, **Zhang H**. Modification of sialic acids on solid-phase: accurate characterization of protein sialylation. Anal Chem. 2017; 16 (6): 1029-1037.
- Li QK, Shah P, Tian Y, Hu Y, Roden R, **Zhang H**, Chan DW. An integrated proteomic and glycoproteomic approach uncovers differences in glycosylation occupancy from benign and malignant epithelial ovarian tumors. Clinical Proteomics. 2017; 14 (1): 1-16.
- Zhou J, Yang W, Hu Y, Hoti N, Liu Y, Shah P, Sun S, Clark DJ, Thomas SN, **Zhang H**. Site-specific fucosylation analysis identifies glycoproteins associated with aggressive prostate cancer cell lines using tandem affinity enrichments of intact glycopeptides followed by mass spectrometry. Anal Chem. 2017; 89 (14): 7623-7630.
- Yang S, Clark D, Liu Y, Li S, **Zhang H.** High-throughput analysis of N-glycans using AutoTip via glycoprotein immobilization. Sci Rep. 2017; 7(1): 10216.
- Yang S, Chen L, Chan DW, Li QK, **Zhang H.** Protein signatures of molecular pathways in non-small cell lung carcinoma (NSCLC): comparison of glycoproteomics and global proteomics. Clinical Proteomics. 2017; 14(1): 31.
- Höti N, Yang S, Aiyetan P, Kumar B, Hu Y, Clark D, Eroglu AU, Shah P, Johnson T, Chowdery WH, **Zhang H,** Rodriguez R. Overexpression of Exportin-5 Overrides the Inhibitory Effect of miRNAs Regulation Control and Stabilize Proteins via Posttranslation Modifications in Prostate Cancer. Neoplasia. 2017; 19 (10): 817-829.
- Thomas SN, Chen L, Liu L, Hoti N, **Zhang H.** Targeted proteomic analyses of Histone H4 acetylation changes associated with homologous recombination deficient high-grade serous ovarian carcinomas. J Proteome Res. 2017; 16 (10): 3704-3710.
- Yang W, Shah P, Hu Y, Toghi Eshghi S, Sun S, Liu Y, **Zhang H**. Comparison of Enrichment Methods for Intact N-and O-linked Glycopeptides Using Strong Anion Exchange and Hydrophilic Interaction Liquid Chromatography. Anal Chem. 2017; 89: 11193-11197.

- Chung C, Wang Q, Yang S, Ponce SA, Kirsch BJ, **Zhang H**, Betenbaugh MJ. Combinatorial genome and protein engineering yields monoclonal antibodies with hypergalactosylation from CHO cells. Biotechnology and Bioengineering, 2017; 114(12): 2848-2856.
- Zhou J-Y, Chen L, Zhang B, Tian Y, Liu T, Thomas SN, Chen L, Schnaubelt M, Boja E, Hiltke T, Kinsinger CR, Rodriguez H, Davies SR, Li S, Snider JE, Erdmann-Gilmore P, Tabb DL, Townsend RR, Ellis MJ, Rodland RD, Smith RD, Carr SA, Zhang Z, Chan DW, **Zhang H**. Quality Assessments of Long-term Quantitative Proteomic Analysis of Breast Cancer Xenograft Tissues. Journal of Proteome Research, 2017; 16 (12): 4523-4530.
- Hoti N, Yang S, Hu Y, Shah P, Haffner M, and **Zhang H**. Overexpression of α (1,6) fucosyltransferase in the development of castration resistant prostate cancer cells. Prostate Cancer and Prostatic Diseases. 2018; **21**: 137–146.
- Upadhyay C, Feyznezhad R, Yang W, **Zhang H**, Zolla-Pazner S, Hioe CE. Alterations of HIV-1 envelope phenotype and antibody-mediated neutralization by signal peptide mutations. PLoS Pathogens, 2018; 14(1), e1006812.
- Uzoma I, Hu J, Cox E, Xia S, Zhou J, Rho H, Guzzo C, Paul C, Ajala O, Goodwin CR, Jeong J, Moore C, **Zhang H**, Meluh P, Blackshaw S, Matunis M, Qian J, Zhu H. Global Identification of SUMO Substrates Reveals Crosstalk between SUMOylation and Phosphorylation Promotes Cell Migration. Molecular & Cellular Proteomics. 2018; 17(5): 871-888
- Sun S, Hu Y, Jia L, Toghi Eshghi S, Liu Y, Shah P, and **Zhang H**. Site-specific Profiling of Serum Glycoproteins Using N-linked Glycan and Glycosite Analysis Reveals Atypical N-Glycosylation Sites on Albumin and Alpha-1B-glycoprotein. Analytical Chemistry. 2018; 90 (10): 6292–6299.
- Hu Y, Shah P, Clark DJ, Ao M, **Zhang H**. Reanalysis of global proteomic and phosphoproteomic data identified a large number of glycopeptides. 2018; 90 (13): 8065–8071. [SI/QI]
- Cameron S, Mix D, Ture S, Schmidt R, Mohan A, Pariser D, Stoner M, Shah P, Chen L, **Zhang H**, Field D, Modjeski K, Toth S, and Morrell C. Hypoxia and Ischemia Promote a Maladaptive Platelet Phenotype. Arteriosclerosis, Thrombosis, and Vascular Biology. 2018; ATVBAHA.118.311186.
- Clark DJ, Hu Y, Bocik W, Chen L, Schnaubelt M, Roberts R, Shah P, Whiteley G, **Zhang H**. Evaluation of NCI-7 Cell Line Panel as a Reference Material for Clinical Proteomics. Journal of Proteome Research. 2018; 17 (6), 2205-2215.
- Wang Q, Chung C-Y, Yang W, Yang G, Chough S, Chen Y, Yin B, Bhattacharya R, Hu Y, Saeui CT, Yarema KJ, Betenbaugh, **Zhang H**. Combining Butyrated ManNAc with Glycoengineered CHO Cells Improves EPO Glycan Quality and Production. Biotechnology journal 2018
- Li Chen, Bai Zhang, Michael Schnaubelt, Punit Shah, Paul Aiyetan, Daniel Chan, **Zhang H**, Zhen Zhang MS-PyCloud: An open-source, cloud computing-based pipeline for LC-MS/MS data analysis. bioRxiv 2018; page, 320887.
- Cho KC, Chen L, Hu Y, Schnaubelt M, **Zhang H**. Developing Workflow for Simultaneous Analyses of Phosphoproteomics and Glycoproteomics. ACS Chemical Biology. 2018; 14, 58-66.
- Yang G, Hu Y, Sun S, Ouyang C, Yang W, Wang Q, Betenbaugh M, **Zhang H**. Comprehensive Glycoproteomic Analysis of Chinese Hamster Ovary Cells. Analytical Chemistry. 2018; 90: 14294-14302.
- Yang W, Ao M, Hu Y, Li QK, **Zhang H**. Mapping the O-glycoproteome using site-specific extraction of O-linked glycopeptides (EXoO). Molecular systems biology. 2018; 14, e8486.
- Shaukat I, Kern JJ, Höti, **Zhang H**, Li JJ, Zheng G, Askin F, Gabrielson E, Li QK. Detection of RAS and RAS-associated alterations in primary lung adenocarcinomas. A correlation between molecular findings and tumor characteristics. Human Pathology. 2019; 84, 18-25.
- Song G, Chen L, Zhang B, Song Q, Yu Y, Moore C, Wang T-L, Shih I-M, **Zhang H**, Chan D, Zhang Z, Zhu H. Proteomewide tyrosine phosphorylation analysis reveals dysregulated signaling pathways in ovarian tumors. Molecular & Cellular Proteomics. 2019; 18: 448-460.

- Stewart AN, Tan SY, Clark DJ, **Zhang H**, Wong GW. N-Linked Glycosylation-Dependent and-Independent Mechanisms Regulating CTRP12 Cleavage, Secretion, and Stability. Biochemistry. 2019: 58: 727-741.
- Wang C, Höti N, Lih TM, Sokoll LJ, Zhang R, Zhang Z, **Zhang H**, Chan DW. Development of a glycoproteomic strategy to detect more aggressive prostate cancer using lectin-immunoassays for serum fucosylated PSA. Clinical proteomics. 2019; 16: 13.
- Clark DJ, Hu Y, Schnaubelt M, Fu Y, Ponce S, Chen S, Zhou Y, Shah P, **Zhang H**. Simple Tip-Based Sample Processing Method for Urinary Proteomic Analysis. Analytical Chemistry. 2019; 9:5517-5522.
- Wang Q, Yang G, Wang T, Yang W, Betenbaugh MJ, **Zhang H.** Characterization of intact glycopeptides reveals the impact of culture media on site-specific glycosylation of EPO-Fc fusion protein generated by CHO-GS cells. Biotechnology and Bioengineering. 2019; 116: 2303-2315.
- Toonstra, C, Hu Y, **Zhang H.** Deciphering the roles of N-glycans on collagen-platelet interactions. Journal of Proteome Research. 2019; 18: 2467-2477.
- Vasaikar S. et al. and **NCI CPTAC Investigators**. Proteogenomic Analysis of Human Colon Cancer Reveals New Therapeutic Opportunities. Cell. 2019; 177: 1035-1049.
- Sun S, Hu Y, Ao M, Shah P, Chen J, Yang W, Jia X, Tian Y, Thomas S, **Zhang H**. N-GlycositeAtlas: a database resource for mass spectrometry-based human N-linked glycoprotein and glycosylation site mapping. Clinical proteomics. 2019; 16: 1-11.
- Shu J, Dang L, Zhang D, Shah P, Chen L, **Zhang H**, Sun S. Dynamic analysis of proteomic alterations in response to N-linked glycosylation inhibition in a drug-resistant ovarian carcinoma cell line. The FEBS journal. 2019; 286: 1594-1605.
- Zhou Y, Höti N, Ao M, Zhang Z, Zhu H, Li L, Askin F, Gabrielson E, **Zhang H**. Li Qk. Expression of p16 and p53 in non-small-cell lung cancer: clinicopathological correlation and potential prognostic impact. Biomarkers in medicine. 2019; 13: 761-771.
- Wang Q, Chung CY, Yang W, Yang G, Chough S, Chen Y, Yin B, Bhattacharya R, Hu Y, Saeui CT, Yarema KJ, Betenbaugh MJ, **Zhang H**. Combining butyrated ManNAc with glycoengineered CHO cells improves EPO glycan quality and production. Biotechnology journal. 2019; 14: 1800186.
- Maria Lorna A De Leoz, David L Duewer, Adam Fung, Lily Liu, Hoi Kei Yau, Oscar Potter, Gregory O Staples, Kenichiro Furuki, Ruth Frenkel, Yunli Hu, Zoran Sosic, Peiging Zhang, Friedrich Altmann, Clemens Gru, Chun Shao, Joseph Zaia, Waltraud Evers, Stuart Pengelley, Detley Suckau, Anja Wiechmann, Anja Resemann, Wolfgang Jabs, Alain Beck, John W Froehlich, Chuncui Huang, Yan Li, Yaming Liu, Shiwei Sun, Yaojun Wang, Youngsuk Seo, Hyun Joo An, Niels-Christian Reichardt, Juan Echevarria Ruiz, Stephanie Archer-Hartmann, Parastoo Azadi, Len Bell, Zsuzsanna Lakos, Yanming An, John F Cipollo, Maja Pucic-Bakovic, Jerko Štambuk, Gordan Lauc, Xu Li, Peng George Wang, Andreas Bock, René Hennig, Erdmann Rapp, Marybeth Creskey, Terry D Cyr, Miyako Nakano, Taiki Sugiyama, Pui-King Amy Leung, Paweł Link-Lenczowski, Jolanta Jaworek, Shuang Yang, Zhang H, Tim Kelly, Song Klapoetke, Rui Cao, Jin Young Kim, Hyun Kyoung Lee, Ju Yeon Lee, Jong Shin Yoo, Sa-Rang Kim, Soo-Kyung Suh, Noortje De Haan, David Falck, Guinevere SM Lageveen-Kammeijer, Manfred Wuhrer, Robert J Emery, Radoslaw P Kozak, Li Phing Liew, Louise Royle, Paulina A Urbanowicz, Nicolle H Packer, Xiaomin Song, Arun Everest-Dass, Erika Lattová, Samanta Cajic, Kathirvel Alagesan, Daniel Kolarich, Toyin Kasali, Viv Lindo, Yuetian Chen, Kudrat Goswami, Brian Gau, Ravi Amunugama, Richard Jones, Corné JM Stroop, Koichi Kato, Hirokazu Yagi, Sachiko Kondo, CT Yuen, Akira Harazono, Xiaofeng Shi, Paula E Magnelli, Brian T Kasper, Lara Mahal, David J Harvey, Roisin O'Flaherty, Pauline M Rudd, Radka Saldova, Elizabeth S Hecht, David C Muddiman, Jichao Kang, Prachi Bhoskar, Daniele Menard, Andrew Saati, Christine Merle, Steven Mast, Sam Tep, Jennie Truong, Takashi Nishikaze, Sadanori Sekiya, Aaron Shafer, Sohei Funaoka, Masaaki Toyoda, Peter De Vreugd, Cassie Caron, Pralima Pradhan, Niclas Chiang Tan, Yehia Mechref, Sachin Patil, Jeffrey S Rohrer, Ranjan Chakrabarti, Disha Dadke, Mohammedazam Lahori, Chunxia Zou, Christopher Cairo, Bela Reiz, Randy M Whittal, Carlito B Lebrilla, Lauren Wu, Andras Guttman, Marton Szigeti, Benjamin G Kremkow, Kelvin H Lee, Carina Sihlbom, Barbara Adamczyk, Chunsheng Jin, Niclas G Karlsson, Jessica Örnros, Göran Larson, Jonas Nilsson, Bernd Meyer, Alena Wiegandt, Emy Komatsu, Helene Perreault, Edward D Bodnar, Nassur Said. NIST

- Interlaboratory Study on Glycosylation Analysis of Monoclonal Antibodies: Comparison of Results from Diverse Analytical Methods. Molecular & Cellular Proteomics. 2019; 19: 11-30.
- David J Clark, Saravana M Dhanasekaran, Francesca Petralia, Jianbo Pan, Xiaoyu Song, Yingwei Hu, Felipe da Veiga Leprevost, Boris Reva, Tung-Shing M Lih, Hui-Yin Chang, Weiping Ma, Chen Huang, Christopher J Ricketts, Lijun Chen, Azra Krek, Yize Li, Dmitry Rykunov, Qing Kay Li, Lin S Chen, Umut Ozbek, Suhas Vasaikar, Yige Wu, Seungyeul Yoo, Shrabanti Chowdhury, Matthew A Wyczalkowski, Jiayi Ji, Michael Schnaubelt, Andy Kong, Sunantha Sethuraman, Dmitry M Avtonomov, Minghui Ao, Antonio Colaprico, Song Cao, Kyung-Cho Cho, Selim Kalayci, Shiyong Ma, Wenke Liu, Kelly Ruggles, Anna Calinawan, Zeynep H Gümüş, Daniel Geiszler, Emily Kawaler, Guo Ci Teo, Bo Wen, Yuping Zhang, Sarah Keegan, Kai Li, Feng Chen, Nathan Edwards, Phillip M Pierorazio, Xi Steven Chen, Christian P Pavlovich, A Ari Hakimi, Gabriel Brominski, James J Hsieh, Andrzej Antczak, Tatiana Omelchenko, Jan Lubinski, Maciej Wiznerowicz, W Marston Linehan, Christopher R Kinsinger, Mathangi Thiagarajan, Emily S Boja, Mehdi Mesri, Tara Hiltke, Ana I Robles, Henry Rodriguez, Jiang Qian, David Fenyö, Bing Zhang, Li Ding, Eric Schadt, Arul M Chinnaiyan, Zhen Zhang, Gilbert S Omenn, Marcin Cieslik, Daniel W Chan, Alexey I Nesvizhskii, Pei Wang, **Zhang H**, and Clinical Proteomic Tumor Analysis Consortium. Integrated proteogenomic characterization of clear cell renal cell carcinoma. Cell. 2019; 179: 964-983.
- Toonstra C, Wang X, Höti N, Pan J, **Zhang H**. Expression of endoplasmic reticulum mannosyl-oligosaccharide α1, 2-mannosidase regulates tumorigenicity of prostate cancer cells. bioRxiv. 2019; 750984.
- Lih TM, Clark DJ, **Zhang H**. Omic-Sig: Utilizing Omics Data to Explore and Visualize Kinase-Substrate Interactions. BioRxiv. 2019; 746123.
- Hu Y, Ao M, Zhang H. OmicsOne: Associate Omics Data with Phenotypes in One-Click. bioRxiv. 2019; 756544.
- Pan J, Qian J, Zhang H. OmicsX: a web server for integrated OMICS analysis. bioRxiv. 2019; 755918.
- van Oostrum M, Mueller M, Klein F, Bruderer R, **Zhang H**, Pedrioli P, Reiter L, Tsapogas P, Rolink A, Wollscheid B. De novo Classification of Mouse B Cell Types using Surfaceome Proteotype Maps. Nature Communications. 2019; 10: 1-9.
- Chen SY, Dong M, Yang G, Zhou Y, Clark DJ, Lih TM, Schnaubelt M, Liu Z, **Zhang H**. Glycans, Glycosite, and Intact Glycopeptide Analysis of N-Linked Glycoproteins Using Liquid Handling Systems. Analytical chemistry. 2020; 92: 1680-1686.
- Zhou Y, Lih TSM, Yang G, Chen SY, Chen L, Chan DW, **Zhang H**, Li QK. An integrated workflow for global, glyco-, and phospho-proteomic analysis of tumor tissues. Analytical chemistry. 2020; 92: 1842-1849.
- Saeui ST, Cho KC, Dharmarha V, Nairn AV, Galizzi M, Shah SR, Gowda P, Park M, Austin M, Clarke A, Cai E, Buettner MJ, Ariss R, Moremen KW, **Zhang H**, Yarema KJ. Cell Line-, Protein-, and Sialoglycosite-Specific Control of Flux-Based Sialylation in Human Breast Cells: Implications for Cancer Progression. Frontiers in chemistry. 2020; 8.
- Jason E McDermott, Osama A Arshad, Vladislav A Petyuk, Yi Fu, Marina A Gritsenko, Therese R Clauss, Ronald J Moore, Athena A Schepmoes, Rui Zhao, Matthew E Monroe, Michael Schnaubelt, Chia-Feng Tsai, Samuel H Payne, Chen Huang, Liang-Bo Wang, Steven Foltz, Matthew Wyczalkowski, Yige Wu, Ehwang Song, Molly A Brewer, Mathangi Thiagarajan, Christopher R Kinsinger, Ana I Robles, Emily S Boja, Henry Rodriguez, Daniel W Chan, Bing Zhang, Zhen Zhang, Li Ding, Richard D Smith, Tao Liu, Karin D Rodland, and Clinical Proteomic Tumor Analysis Consortium (Jasmin Bavarva, Emily Boja, Melissa Borucki, Shuang Cai, Steven A. Carr, Daniel Chan, Xian Chen, Karl R. Clauser, Sherri R. Davies, Li Ding, Nathan J. Edwards, Kimberly Elburn, Matthew J. C. Ellis, David Fenyo, Michael A. Gillette, Linda Hannick, Tara Hiltke, Andrew Hoofnagle, Karen A. Ketchum, Christopher R. Kinsinger, Eric Kuhn, Douglas A. Levine, Daniel C. Liebler, Tao Liu, Subha Madhavan, D. R. Mani, Jason E. McDermott, Peter B. McGarvey, Philip Mertins, Mehdi Mesri, Mauricio Oberti, Akhilesh Pandey, Amanda Paulovich, Samuel H. Payne, David F. Ransohoff, Robert C. Rivers, Karin D. Rodland, Henry Rodriguez, Paul A. Rudnick, Kelly Ruggles, Melinda E. Sanders, Zhiao Shi, Ie-Ming Shih, Steven J. Skates, Robert J. C. Slebos, Richard D. Smith, Michael Snyder, Stephen E. Stein, David L. Tabb, Ratna R. Thangudu, Mathangi Thiagarajan, R. Reid Townsend, Negin Vatanian, Yue Wang, Mark Watson, Forest M. White, Jeffery Whiteaker, Gordon A. Whiteley, Bing Zhang, Zhang H, Zhen Zhang, Yingming Zhao, Heng Zhu, and Lisa J. Zimmerman). Proteogenomic

- Characterization of Ovarian HGSC Implicates Mitotic Kinases, Replication Stress in Observed Chromosomal Instability. Cell Reports Medicine. 2020; 1: 100004.
- Mertz JL, Sripathi SR, Yang X, Chen L, Esumi N, **Zhang H**, Zack DJ. Proteomic and phosphoproteomic analysis identifies novel liver-related signaling in retinal pigment epithelial cells during epithelial-mesenchymal transition. bioRxiv. 2020.
- Yang G, Höti N, Chen SY, Zhou Y, Wang Q, Betenbaugh M, **Zhang H**. One-step enrichment of intact glycopeptides from glycoengineered Chinese hamster ovary cells. Frontiers in Chemistry 2020; 8.
- N Höti, TS Lih, J Pan, Y Zhou, G Yang, A Deng, L Chen, M Dong, Yang R-B, Tu CF, Haffner HC, Li QK, **Zhang H**. A Comprehensive Analysis of FUT8 Overexpressing Prostate Cancer Cells Reveals the Role of EGFR in Castration Resistance. Cancers 2020; 12: 468.
- Cho KC, Clark DJ, Schnaubelt M, Teo GC, Leprevost FV, Bocik W, Boja ES, Hiltke T, Nesvizhskii AI, **Zhang H.** Deep Proteomics Using Two-Dimensional Data Independent Acquisition Mass Spectrometry. Analytical chemistry. 2020; 92: 4217-4225.
- Yongchao Dou, Emily A Kawaler, Daniel Cui Zhou, Marina A Gritsenko, Chen Huang, Lili Blumenberg, Alla Karpova, Vladislav A Petyuk, Sara R Savage, Shankha Satpathy, Wenke Liu, Yige Wu, Chia-Feng Tsai, Bo Wen, Zhi Li, Song Cao, Jamie Moon, Zhiao Shi, MacIntosh Cornwell, Matthew A Wyczalkowski, Rosalie K Chu, Suhas Vasaikar, Hua Zhou, Qingsong Gao, Ronald J Moore, Kai Li, Sunantha Sethuraman, Matthew E Monroe, Rui Zhao, David Heiman, Karsten Krug, Karl Clauser, Ramani Kothadia, Yosef Maruvka, Alexander R Pico, Amanda E Oliphant, Emily L Hoskins, Samuel L Pugh, Sean JI Beecroft, David W Adams, Jonathan C Jarman, Andy Kong, Hui-Yin Chang, Boris Reva, Yuxing Liao, Dmitry Rykunov, Antonio Colaprico, Xi Steven Chen, Andrzej Czekański, Marcin Jędryka, Rafal Matkowski, Maciej Wiznerowicz, Tara Hiltke, Emily Boja, Christopher R Kinsinger, Mehdi Mesri, Ana I Robles, Henry Rodriguez, David Mutch, Katherine Fuh, Matthew J Ellis, Deborah DeLair, Mathangi Thiagarajan, DR Mani, Gad Getz, Michael Noble, Alexey I Nesvizhskii, Pei Wang, Matthew L Anderson, Douglas A Levine, Richard D Smith, Samuel H Payne, Kelly V Ruggles, Karin D Rodland, Li Ding, Bing Zhang, Tao Liu, David Fenyö, and Clinical Proteomic Tumor Analysis Consortium (... Hui Zhang, Yuping Zhang, Zhen Zhang, Grace Zhao, and Hua Zhou). Proteogenomic Characterization of Endometrial Carcinoma. Cell. 2020; 180: 729-748.
- Thomas SN, Friedrich B, Schnaubelt M, Chan DW, **Zhang H**, Aebersold R. Orthogonal proteomic platforms and their implications for the stable classification of high-grade serous ovarian cancer subtypes. iScience. 2020; 101079.
- Hristova V, Sun S, **Zhang H**, Chan DW. Proteomic analysis of degradation ubiquitin signaling by ubiquitin occupancy changes responding to 26S proteasome inhibition. Clinical proteomics. 2020; 17: 1-11.
- Clark DJ, Schnaubelt M, Höti N, Hu Y, Zhou Y, Gooya M, **Zhang H**. Impact of Increased FUT8 Expression on the Extracellular Vesicle Proteome in Prostate Cancer Cells. Journal of Proteome Research. 2020; 19: 2195-2205.
- Yang W, Ao M, Song A, Xu Y, Sokoll SJ, **Zhang H**, Mass Spectrometric Mapping of Glycoproteins Modified by Tn-antigen Using Solid-phase Capture and Enzymatic Release. Analytical Chemistry. 2020; 92: 9230-9238.
- Ashley N Stewart, Hannah C Little, David J Clark, **Zhang H**, G William Wong. Protein modifications critical for myonectin/erythroferrone secretion and oligomer assembly. Biochemistry. 2020; 59: 2684-2697.
- Michael A Gillette, Shankha Satpathy, Song Cao, Saravana M Dhanasekaran, Suhas V Vasaikar, Karsten Krug, Francesca Petralia, Yize Li, Wen-Wei Liang, Boris Reva, Azra Krek, Jiayi Ji, Xiaoyu Song, Wenke Liu, Runyu Hong, Lijun Yao, Lili Blumenberg, Sara R Savage, Michael C Wendl, Bo Wen, Kai Li, Lauren C Tang, Melanie A MacMullan, Shayan C Avanessian, M Harry Kane, Chelsea J Newton, MacIntosh Cornwell, Ramani B Kothadia, Weiping Ma, Seungyeul Yoo, Rahul Mannan, Pankaj Vats, Chandan Kumar-Sinha, Emily A Kawaler, Tatiana Omelchenko, Antonio Colaprico, Yifat Geffen, Yosef E Maruvka, Felipe da Veiga Leprevost, Maciej Wiznerowicz, Zeynep H Gümüş, Rajwanth R Veluswamy, Galen Hostetter, David I Heiman, Matthew A Wyczalkowski, Tara Hiltke, Mehdi Mesri, Christopher R Kinsinger, Emily S Boja, Gilbert S Omenn, Arul M Chinnaiyan, Henry Rodriguez, Qing Kay Li, Scott D Jewell, Mathangi Thiagarajan, Gad Getz, Bing Zhang, David Fenyö, Kelly V Ruggles, Marcin P Cieslik, Ana I Robles, Karl R Clauser, Ramaswamy Govindan, Pei Wang, Alexey I Nesvizhskii, Li Ding, DR Mani, Steven

- A Carr, and Clinical Proteomic Tumor Analysis Consortium (... **Zhang H** ...). Proteogenomic characterization reveals therapeutic vulnerabilities in lung adenocarcinoma. Cell. 2020; 182: 200-225.
- Yangying Zhou, Weiming Yang, Minghui Ao, Naseruddin Höti, Edward Gabrielson, Daniel W Chan, **Zhang H**, Qing Kay Li. Proteomic Analysis of the Air-way Fluid in Lung Cancer. Detection of Periostin in Bronchoalveolar Lavage (BAL). Frontiers in Oncology. 2020; 10: 1072.
- Mingming Dong, T. Mamie Lih, Shao-Yung Chen, Kyung-Cho Cho, Rodrigo Vargas Eguez, Naseruddin Höti, Yangying Zhou, Weiming Yang, Leslie Mangold, Daniel W. Chan, Zhen Zhang, Lori J. Sokoll, Alan Partin, **Zhang H.** Urinary glycoproteins associated with aggressive prostate cancer. Theranostics. 2020; 10 (26): 11892.
- Yingwei Hu, Jianbo Pan, Punit Shah, Minghui Ao, Stefani N. Thomas, Yang Liu, Lijun Chen, Michael Schnaubelt, David J. Clark, Henry Rodriguez, Emily S Boja, Tara Hiltke, Christopher R. Kinsinger, Karin D. Rodland, Q Kay Li, Jiang Qian, Zhen Zhang, Daniel W. Chan, **Zhang H**. and the NCI CPTAC Investigators. Integrated Glycoproteomic Characterization of Human High-Grade Serous Ovarian Cancer. Cell Reports. 2020; 33(3): 108276.
- Debebe Theodros, Benjamin M Murter, John-William Sidhom, Thomas R Nirschl, David J Clark, LiJun Chen, Ada J Tam, Richard L Blosser, Zeyad R Schwen, Michael H Johnson, Phillip M Pierorazio, **Zhang H**, Sudipto Ganguly, Drew M Pardoll, Jelani C Zarif. High-Dimensional Cytometry (ExCYT) and Mass Spectrometry of myeloid infiltrate in clinically localized clear cell Renal Cell Carcinoma identifies novel potential myeloid targets for immunotherapy. Molecular & Cellular Proteomics. 2020; 19 (11): 1850-1859.
- Karsten Krug, Eric J Jaehnig, Shankha Satpathy, Lili Blumenberg, Alla Karpova, Meenakshi Anurag, George Miles, Philipp Mertins, Yifat Geffen, Lauren C Tang, David I Heiman, Song Cao, Yosef E Maruvka, Jonathan T Lei, Chen Huang, Ramani B Kothadia, Antonio Colaprico, Chet Birger, Jarey Wang, Yongchao Dou, Bo Wen, Zhiao Shi, Yuxing Liao, Maciej Wiznerowicz, Matthew A Wyczalkowski, Xi Steven Chen, Jacob J Kennedy, Amanda G Paulovich, Mathangi Thiagarajan, Christopher R Kinsinger, Tara Hiltke, Emily S Boja, Mehdi Mesri, Ana I Robles, Henry Rodriguez, Thomas F Westbrook, Li Ding, Gad Getz, Karl R Clauser, David Fenyö, Kelly V Ruggles, Bing Zhang, DR Mani, Steven A Carr, Matthew J Ellis, Michael A Gillette, Shayan C Avanessian, Shuang Cai, Daniel Chan, Xian Chen, Nathan J Edwards, Andrew N Hoofnagle, M Harry Kane, Karen A Ketchum, Eric Kuhn, Douglas A Levine, Shunqiang Li, Daniel C Liebler, Tao Liu, Jingqin Luo, Subha Madhavan, Chris Maher, Jason E McDermott, Peter B McGarvey, Mauricio Oberti, Akhilesh Pandey, Samuel H Payne, David F Ransohoff, Robert C Rivers, Karin D Rodland, Paul Rudnick, Melinda E Sanders, Kenna M Shaw, Ie-Ming Shih, Robbert JC Slebos, Richard D Smith, Michael Snyder, Stephen E Stein, David L Tabb, Ratna R Thangudu, Stefani Thomas, Yue Wang, Forest M White, Jeffrey R Whiteaker, Gordon A Whiteley, **Zhang H**, Zhen Zhang, Yingming Zhao, Heng Zhu, Lisa J Zimmerman. Proteogenomic landscape of breast cancer tumorigenesis and targeted therapy. 2020; Cell 183 (5), 1436-1456.
- Joseph L Mertz, Shisheng Sun, Bojiao Yin, Yingwei Hu, Rahul Bhattacharya, Michael J Bettenbaugh, Kevin J Yarema, **Zhang H**. Comparison of Three Glycoproteomic Methods for the Analysis of the Secretome of CHO Cells Treated with 1, 3, 4-O-Bu3ManNAc. 2020; Bioengineering 7 (4): 144.
- Jianbo Pan, Yingwei Hu, Shisheng Sun, Lijun Chen, Michael Schnaubelt, David Clark, Minghui Ao, Zhen Zhang, Daniel Chan, Jiang Qian, **Zhang H**. Glycoproteomics-based signatures for tumor subtyping and clinical outcome prediction of high-grade serous ovarian cancer. 2020; Nature communications 11 (1): 1-13.
- Yangying Zhou, T Mamie Lih, Jianbo Pan, Naseruddin Höti, Mingming Dong, Liwei Cao, Yingwei Hu, Kyung-Cho Cho, Shao-Yung Chen, Rodrigo Vargas Eguez, Edward Gabrielson, Daniel W Chan, **Zhang H**, Qing Kay Li. Proteomic signatures of 16 major types of human cancer reveal universal and cancer-type-specific proteins for the identification of potential therapeutic targets. 2020; Journal of hematology & oncology 13 (1): 1-15.
- Chitra Upadhyay, Roya Feyznezhad, Liwei Cao, Kun-Wei Chan, Kevin Liu, Weiming Yang, **Zhang H**, Jason Yolitz, James Arthos, Arthur Nadas, Xiang-Peng Kong, Susan Zolla-Pazner, Catarina E Hioe. Signal peptide of HIV-1 envelope modulates glycosylation impacting exposure of V1V2 and other epitopes. 2020; PLoS pathogens 16 (12): e1009185.
- Chen Huang, Lijun Chen, Sara R Savage, Rodrigo Vargas Eguez, Yongchao Dou, Yize Li, Felipe da Veiga Leprevost, Eric J Jaehnig, Jonathan T Lei, Bo Wen, Michael Schnaubelt, Karsten Krug, Xiaoyu Song, Marcin Cieślik, Hui-Yin Chang, Matthew A Wyczalkowski, Kai Li, Antonio Colaprico, Qing Kay Li, David J Clark, Yingwei Hu, Liwei Cao, Jianbo Pan, Yuefan Wang, Kyung-Cho Cho, Zhiao Shi, Yuxing Liao, Wen Jiang, Meenakshi Anurag, Jiayi Ji,

- Seungyeul Yoo, Daniel Cui Zhou, Wen-Wei Liang, Michael Wendl, Pankaj Vats, Steven A Carr, DR Mani, Zhen Zhang, Jiang Qian, Xi S Chen, Alexander R Pico, Pei Wang, Arul M Chinnaiyan, Karen A Ketchum, Christopher R Kinsinger, Ana I Robles, Eunkyung An, Tara Hiltke, Mehdi Mesri, Mathangi Thiagarajan, Alissa M Weaver, Andrew G Sikora, Jan Lubiński, Malgorzata Wierzbicka, Maciej Wiznerowicz, Shankha Satpathy, Michael A Gillette, George Miles, Matthew J Ellis, Gilbert S Omenn, Henry Rodriguez, Emily S Boja, Saravana M. Dhanasekaran, Li Ding, Alexey I. Nesvizhskii, Adel K.El-Naggar, Daniel W. Chan, **Zhang H,** Bing Zhang, and Clinical Proteomic Tumor Analysis Consortium. Proteogenomic insights into the biology and treatment of HPV-negative head and neck squamous cell carcinoma. 2021; 39 (3); 361-379.e16.
- Liang-Bo Wang, Alla Karpova, Marina A Gritsenko, Jennifer E Kyle, Song Cao, Yize Li, Dmitry Rykunov, Antonio Colaprico, Joseph H Rothstein, Runyu Hong, Vasileios Stathias, MacIntosh Cornwell, Francesca Petralia, Yige Wu, Boris Reva, Karsten Krug, Pietro Pugliese, Emily Kawaler, Lindsey K Olsen, Wen-Wei Liang, Xiaoyu Song, Yongchao Dou, Michael C Wendl, Wagma Caravan, Wenke Liu, Daniel Cui Zhou, Jiayi Ji, Chia-Feng Tsai, Vladislav A Petyuk, Jamie Moon, Weiping Ma, Rosalie K Chu, Karl K Weitz, Ronald J Moore, Matthew E Monroe, Rui Zhao, Xiaolu Yang, Seungyeul Yoo, Azra Krek, Alexis Demopoulos, Houxiang Zhu, Matthew A Wyczalkowski, Joshua F McMichael, Brittany L Henderson, Caleb M Lindgren, Hannah Boekweg, Shuangjia Lu, Jessika Baral, Lijun Yao, Kelly G Stratton, Lisa M Bramer, Erika Zink, Sneha P Couvillion, Kent J Bloodsworth, Shankha Satpathy, Weiva Sieh, Simina M Boca, Stephan Schürer, Feng Chen, Maciej Wiznerowicz, Karen A Ketchum, Emily S Boja, Christopher R Kinsinger, Ana I Robles, Tara Hiltke, Mathangi Thiagarajan, Alexey I Nesvizhskii, Bing Zhang, DR Mani, Michele Ceccarelli, Xi S Chen, Sandra L Cottingham, Qing Kay Li, Albert H Kim, David Fenyö, Kelly V Ruggles, Henry Rodriguez, Mehdi Mesri, Samuel H Payne, Adam C Resnick, Pei Wang, Richard D Smith, Antonio Iavarone, Milan G Chheda, Jill S Barnholtz-Sloan, Karin D Rodland, Tao Liu, Li Ding, ... Zhang H, Liwei Zhang, Zhen Zhang, Grace Zhao, Jun Zhu. Proteogenomic and metabolomic characterization of human glioblastoma. 2021; Cancer cell 39 (4): 509-528. e20.
- Jin Song, Shiyong Ma, Lori J. Sokoll, Rodrigo V. Eguez, Naseruddin Höti, **Zhang H**, Phaedre Mohr, Renu Dua, Dattatraya Patil, Kristen Douglas May, Sierra Williams, Rebecca Arnold, Martin G. Sanda, Daniel W. Chan, Zhen Zhang. A panel of selected serum protein biomarkers for the detection of aggressive prostate cancer. 2021; Theranostics 2021; 11(13):6214-6224.
- Cao, L, Huang, C, Cui Zhou, D, Hu, Y, Lih, TM, Savage, SR, Krug, K, Clark, DJ, Schnaubelt, M, Chen, L, da Veiga Leprevost, F, Eguez, RV, Yang, W, Pan, J, Wen, B, Dou, Y, Jiang, W, Liao, Y, Shi, Z, Terekhanova, NV, Cao, S, Lu, RJ, Li, Y, Liu, R, Zhu, H, Ronning, P, Wu, Y, Wyczalkowski, MA, Easwaran, H, Danilova, L, Mer, AS, Yoo, S, Wang, JM, Liu, W, Haibe-Kains, B, Thiagarajan, M, Jewell, SD, Hostetter, G, Newton, CJ, Li, QK, Roehrl, MH, Fenyo, D, Wang, P, Nesvizhskii, AI, Mani, DR, Omenn, GS, Boja, ES, Mesri, M, Robles, AI, Rodriguez, H, Bathe, OF, Chan, DW, Hruban, RH, Ding, L, Zhang, B, **Zhang, H**, Clinical Proteomic Tumor Analysis, C. Proteogenomic characterization of pancreatic ductal adenocarcinoma. Cell. 2021;184(19):5031-52 e26. Epub 2021/09/18. doi: 10.1016/j.cell.2021.08.023. PubMed PMID: 34534465.
- Chen, SY, Clark, DJ, **Zhang, H**. High-throughput analyses of glycans, glycosites, and intact glycopeptides using C4-and C18/MAX-tips and liquid handling system. Current Protocols. 2021;1(7):e186.
- Cho, K-C, Oh, S, Wang, Y, Rosenthal, LS, Na, CH, **Zhang, H**. Evaluation of the Sensitivity and Reproducibility of Targeted Proteomic Analysis Using Data-Independent Acquisition for Serum and Cerebrospinal Fluid Proteins. J Proteome Res. 2021;20(9):4284-91.
- Dong, M, Lih, T-SM, Ao, M, Hu, Y, Chen, S-Y, Eguez, RV, **Zhang, H**. Data-Independent Acquisition-Based Mass Spectrometry (DIA-MS) for Quantitative Analysis of Intact N-Linked Glycopeptides. Analytical chemistry. 2021;93(41):13774-82.
- Dong, M, Lih, T-SM, Hoti, N, Chen, S-Y, Ponce, S, Partin, A, **Zhang, H**. Development of Parallel Reaction Monitoring Assays for the Detection of Aggressive Prostate Cancer Using Urinary Glycoproteins. J Proteome Res. 2021;20(7):3590-9.
- **Zhang, H**, Ao, M, Boja, A, Schnaubelt, M, Hu, Y. OmicsOne: associate omics data with phenotypes in one-click. Clinical Proteomics. 2021;18(1):1-12.

- Li, QK, Chen, J, Hu, Y, Höti, N, Lih, T-SM, Thomas, SN, Chen, L, Roy, S, Meeker, A, Shah, P, Chen, L, Bova, GS, Zhang, B, **Zhang, H**. Proteomic characterization of primary and metastatic prostate cancer reveals reduced proteinase activity in aggressive tumors. Scientific reports. 2021;11(1):1-14.
- Yang, G, Wang, Q, Chen, L, Betenbaugh, MJ, **Zhang, H**. Glycoproteomic Characterization of FUT8 Knock-out CHO cells Reveals Roles of FUT8 in the Glycosylation. Frontiers in Chemistry. 2021;9.
- Maxwell, MJ, Arnold, A, Sweeney, H, Chen, L, Lih, T-SM, Schnaubelt, M, Eberhart, CG, Rubens, JA, **Zhang, H**, Clark, DJ, Raabe, EH. Unbiased proteomic and phosphoproteomic analysis identifies response signatures and novel susceptibilities after combined MEK and mTOR inhibition in BRAFV600E mutant glioma. Mol Cell Proteomics. 2021;20.
- Anyaogu, DC, Hansen, AH, Hoof, JB, Majewska, NI, Contesini, FJ, Paul, JT, Nielsen, KF, Hobley, TJ, Yang, S, **Zhang, H,** Betenbaugh, M, Mortensen, UH. Glycoengineering of Aspergillus nidulans to produce precursors for humanized N-glycan structures. Metabolic engineering. 2021;67:153-63.
- Mertz, JL, Sripathi, SR, Yang, X, Chen, L, Esumi, N, **Zhang, H**, Zack, DJ. Proteomic and phosphoproteomic analyses identify liver-related signaling in retinal pigment epithelial cells during EMT. Cell Reports. 2021;37(3):109866.
- Kawahara, R, Chernykh, A, Alagesan, K, Bern, M, Cao, W, Chalkley, RJ, Cheng, K, Choo, MS, Edwards, N, Goldman, R, Hoffmann, M, Hu, Y, Huang, Y, Kim, JY, Kletter, D, Liquet, B, Liu, M, Mechref, Y, Meng, B, Neelamegham, S, Nguyen-Khuong, T, Nilsson, J, Pap, A, Park, GW, Parker, BL, Pegg, CL, Penninger, JM, Phung, TK, Pioch, M, Rapp, E, Sakalli, E, Sanda, M, Schulz, BL, Scott, NE, Sofronov, G, Stadlmann, J, Vakhrushev, SY, Woo, CM, Wu, HY, Yang, P, Ying, W, **Zhang, H**, Zhang, Y, Zhao, J, Zaia, J, Haslam, SM, Palmisano, G, Yoo, JS, Larson, G, Khoo, KH, Medzihradszky, KF, Kolarich, D, Packer, NH, Thaysen-Andersen, M. Community evaluation of glycoproteomics informatics solutions reveals high-performance search strategies for serum glycopeptide analysis. Nat Methods. 2021;18(11):1304-16.
- Satpathy, S., Krug, K., Jean Beltran, PM, Savage, SR, Petralia, F, Kumar-Sinha, C, Dou, Y, Reva, B, Kane, MH, Avanessian, SC, Vasaikar, SV, Krek, A, Lei, JT, Jaehnig, EJ, Omelchenko, T, Geffen, Y, Bergstrom, EJ, Stathias, V, Christianson, KE, Heiman, DI, Cieslik, MP, Cao, S, Song, X, Ji, J, Liu, W, Li, K, Wen, B, Li, Y, Gumus, ZH, Selvan, ME, Soundararajan, R, Visal, TH, Raso, MG, Parra, ER, Babur, O, Vats, P, Anand, S, Schraink, T, Cornwell, M, Rodrigues, FM, Zhu, H, Mo, CK, Zhang, Y, da Veiga Leprevost, F, Huang, C, Chinnaiyan, AM, Wyczalkowski, MA, Omenn, GS, Newton, CJ, Schurer, S, Ruggles, KV, Fenyo, D, Jewell, SD, Thiagarajan, M, Mesri, M, Rodriguez, H, Mani, SA, Udeshi, ND, Getz, G, Suh, J, Li, QK, Hostetter, G, Paik, PK, Dhanasekaran, SM, Govindan, R, Ding, L, Robles, AI, Clauser, KR, Nesvizhskii, AI, Wang, P, Carr, SA, Zhang, B, Mani, DR, Gillette, MA, Clinical Proteomic Tumor Analysis, C (Bing Zhang, DR Mani, Michael A Gillette, Alex Green, Alfredo Molinolo, Alicia Francis, Amanda G Paulovich, Andrii Karnuta, Antonio Colaprico, Barbara Hindenach, Barbara L Pruetz, Bartosz Kubisa, Brian J Druker, Carissa Huynh, Charles A Goldthwaite Jr, Chet Birger, Christopher R Kinsinger, Corbin D Jones, Dan Rohrer, Dana R Valley, Daniel W Chan, David Chesla, Donna Hansel, Elena V Ponomareva, Elizabeth Duffy, Eric Burks, Eric E Schadt, Eugene S Fedorov, Eunkyung An, Fei Ding, George D Wilson, Harsh Batra, Zhang, H, Jennifer E Maas, Jennifer Eschbacher, Karen A Ketchum, Karin D Rodland, Katherine A Hoadley, Kei Suzuki, Ki Sung Um, Liqun Qi, Lori Bernard, Maciej Wiznerowicz, Małgorzata Wojtyś, Marcin J Domagalski, Matthew J Ellis, Maureen A Dyer, Melissa Borucki, Meenakshi Anurag, Michael J Birrer, Midie Xu, Mikhail Krotevich, Nancy Roche, Nathan J Edwards, Negin Vatanian, Neil R Mucci, Nicollette Maunganidze, Nikolay Gabrovski, Olga Potapova, Oluwole Fadare, Pamela Grady, Peter B McGarvey, Pushpa Hariharan, Ratna R Thangudu, Rebecca Montgomery, Renganayaki Pandurengan, Richard D Smith, Robert J Welsh, Sailaja Mareedu, Samuel H Payne, Sandra Cottingham, Shilpi Singh, Shirley X Tsang, Shuang Cai, Stacey Gabriel, Tao Liu, Tara Hiltke, Tanmayi Vashist, Thomas Bauer). A proteogenomic portrait of lung squamous cell carcinoma. Cell. 2021;184(16):4348-71 e40.
- Zieren, RC, Dong, L, Clark, DJ, Kuczler, MD, Horie, K, Moreno, LF, Lih, TM, Schnaubelt, M, Vermeulen, L, **Zhang, H**, de Reijke, TM, Pienta, KJ, Amend, SR. Defining candidate mRNA and protein EV biomarkers to discriminate ccRCC and pRCC from non-malignant renal cells in vitro. Med Oncol. 2021;38(9):105.
- Wang, Y, Hu, Y, Höti, N, Huang, L, **Zhang, H**. Characterization of In Vivo Protein Complexes via Chemical Cross-Linking and Mass Spectrometry. Analytical Chemistry. 2022; 94(3): 1537–1542.

- Chen, S-Y, Lih, T-SM, Li, QK, **Zhang, H**. Comparing Urinary Glycoproteins among Three Urogenital Cancers and Identifying Prostate Cancer-Specific Glycoproteins. ACS omega. 2022;7(11):9172-80.
- Cao, X, Sanchez, JC, Dinabandhu, A, Guo, C, Patel, TP, Yang, Z, Hu, MW, Chen, L, Wang, Y, Malik, D, Jee, K, Daoud, YJ, Handa, JT, **Zhang, H**, Qian, J, Montaner, S, Sodhi, A. Aqueous proteins help predict the response of patients with neovascular age-related macular degeneration to anti-VEGF therapy. J Clin Invest. 2022;132(2).
- Demirhan, D, Kumar, A, Zhu, J, Poulsen, PC, Majewska, NI, Sebastian, Y, Chaerkady, R, Yu, W, Zhu, W, Zhu, W, Zhu, Shah, P, Lekstrom, K, Cole, RN, **Zhang, H**, Betenbaugh, MJ, Bowen, MA. Comparative systeomics to elucidate physiological differences between CHO and SP2/0 cell lines. Sci Rep. 2022;12(1):3280.
- Cao L, Lih, TM, Hu Y, Schnaubelt M, Chen S-Y, Zhou Y, Guo C, Dong M, Yang W, Eguez RV, Chen L, Clark DJ, Sodhi A, Li QK, **Zhang H**. Characterization of core fucosylation via sequential enzymatic treatments of intact glycopeptides and mass spectrometry analysis. Nature Communications. 2022; 13(1):1-12.
- Wang Y, Lih TM, Chen L, Xu Y, Kuczler MD, Cao L, Pienta KJ, Amend SR, **Zhang H**. Optimized data-independent acquisition approach for proteomic analysis at single-cell level. Clinical Proteomics. 2022; 19 (1): 1-10.
- Li, QK, Lih, T-SM, Wang, Y, Hu, Y, Höti, N, Chan, DW, **Zhang, H**. Improving the detection of aggressive prostate cancer using immunohistochemical staining of protein marker panels. American journal of cancer research. 2022;12(3):1323.
- Cui Zhou D, Jayasinghe RG, Chen S, Herndon JM, Iglesia MD, Navale P, Wendl MC, Caravan W, Sato K, Storrs E, Mo CK, Liu J, Southard-Smith AN, Wu Y, Naser Al Deen N, Baer JM, Fulton RS, Wyczalkowski MA, Liu R, Fronick CC, Fulton LA, Shinkle A, Thammavong L, Zhu H, Sun H, Wang LB, Li Y, Zuo C, McMichael JF, Davies SR, Appelbaum EL, Robbins KJ, Chasnoff SE, Yang X, Reeb AN, Oh C, Serasanambati M, Lal P, Varghese R, Mashl JR, Ponce J, Terekhanova NV, Yao L, Wang F, Chen L, Schnaubelt M, Lu RJ, Schwarz JK, Puram SV, Kim AH, Song SK, Shoghi KI, Lau KS, Ju T, Chen K, Chatterjee D, Hawkins WG, **Zhang H**, Achilefu S, Chheda MG, Oh ST, Gillanders WE, Chen F, DeNardo DG, Fields RC, Ding L. Spatially restricted drivers and transitional cell populations cooperate with the microenvironment in untreated and chemo-resistant pancreatic cancer. Nat Genet. 2022;54(9):1390-405. Epub 2022/08/23. doi: 10.1038/s41588-022-01157-1. PubMed PMID: 35995947; PMCID: PMC9470535.
- Lih TM, Dong M, Mangold L, Partin A, **Zhang H**. Urinary marker panels for aggressive prostate cancer detection. Scientific reports. 2022: 12 (1): 1-9.

# Review Articles [RA]

- **Zhang H**, Yan W, Aebersold R. Chemical probes and tandem mass spectrometry: a strategy for the quantitative analysis of proteomes and subproteomes. Curr Opin Chem Biol. 2004; 8: 66-75.
- Zhang H. The Plasma Proteome: High Abundance versus Low Abundance. Expert Rev Proteomics. 2006;3:175-8
- Tian Y, Zhang H. Glycoproteomics and clinical applications. Proteom-Clin App. 2010; 4: 124-132.
- Yang S, Zhang H. Solid-phase glycan isolation for glycomics analysis. Proteomics Clin Appl. 2012; 6: 596-608.
- Li QK, Gabrielson E, **Zhang H**. Application of glycoproteomics in the discovery of biomarkers for lung cancer. Proteomics Clin Appl. 2012; 6: 244-256.
- Tian Y, Zhang H. Characterization of disease-associated N-linked glycoproteins. Proteomics. 2013; 13:504-511
- Baycin-Hizal D, Wolozny D, Colao J, Jacobson E, Tian Y, Krag SS, Betenbaugh MJ, **Zhang H.** Glyco-proteomic and Glycomic Databases. Clin Proteomics. 2014; 11: 1-10.

- Baycin-Hizal D, Gottschalk A, Jacobson E, Mai S, Wolozny D, Zhang H, Krag SS, Betenbaugh MJ. Physiologic and pathophysiologic consequences of altered sialylation and glycosylation on ion channel function. Biochemical and Biophysical Research Communications. 2014; 453: 243-253.
- Yang S, Rubin A, Toghi Eshghi S, **Zhang H**. Chemoenzymatic method for glycomics: isolation, identification, and quantitation. Proteomics. 2016; 16: 241-256.
- Bennun SV, Baycin-Hizal D, Heffner K, Can O, **Zhang H**, Betenbaugh M. Systems glycobiology: integrating glycogenomics, glycoproteomics, glycomics and other 'omics data sets to characterize cellular glycosylation processes. J Mol Bio. 2016; 428: 3337-3352.
- Thomas S, **Zhang H**. Targeted proteomic assays for the verification of global proteomics insights Expert Rev Proteomics. 2016; 13: 897-899.
- Jimenez CR, **Zhang H**, Kinsinger CR, Nice EC. The cancer proteomic landscape and the HUPO Cancer Proteome Project. Clinical Proteomics. 2018; 15 (1): 4.
- Hisashi Narimatsu, Hiroyuki Kaji, Sergey Y. Vakhrushev, Henrik Clausen, **Zhang H**, Erika Noro†, Akira Togayachi, Chiaki Nagai-Okatani, Atsushi Kuno, Xia Zou, Li Cheng, Sheng-Ce Tao, and Yangyang Sun Current Technologies for Complex Glycoproteomics and Their Applications to Biology/Disease-Driven Glycoproteomics. Journal of Proteome Research. 2018; 17, 4097-4112.
- QK Li, CP Pavlovich, **Zhang H**, CR Kinsinger, DW Chan. Challenges and opportunities in the proteomic characterization of clear cell renal cell carcinoma (ccRCC): A critical step towards the personalized care of renal cancers. Seminars in cancer biology. 2018; https://doi.org/10.1016/j.semcancer.2018.06.004.
- Dang L, Jia L, Zhi Y, Li P, Zhao T, Zhu B, Lan R, Hu Y, **Zhang H**, Sun S. Mapping human N-linked glycoproteins and glycosylation sites using mass spectrometry. TrAC Trends in Analytical Chemistry. 2019; 114: 143-150.
- Clark DJ, Zhang H. Proteomic approaches for characterizing renal cell carcinoma. Clinical Proteomics. 2020; 17(1): 1-18.
- Xu, Y, Wang, Y, Höti, N, Clark, DJ, Chen, SY, **Zhang, H**. The next "sweet" spot for pancreatic ductal adenocarcinoma: Glycoprotein for early detection. Mass Spectrometry Reviews. 2021:e21748.
- Xu, Y, **Zhang, H**. Putting the pieces together: mapping the O-glycoproteome. Current Opinion in Biotechnology. 2021;71:130-6.
- Yang, G, **Zhang, H,** Yi, W, Yan, S, Cao, L. Protein Glycosylation—Advances in Identification, Characterization and Biological Function Elucidation Using Mass Spectrometry. Frontiers in Chemistry. 2022;10.

## Book Chapters, Monographs [BC]

- Yi EC, **Zhang H,** Cooke K, Aebersold, R, Goodlett DR. Quantitative analysis proteomes and subproteomes by Isotope Coded Affinity Tags (ICAT) and solid-phase glycoprotein capture. The Proteomics Handbook. Walker JM (ed), Humana Press Totowa, NJ, 2004
- **Zhang H**, Aebersold R. Isolation of glycoproteins and identification of their N-linked glycosylation sites. Method Mol Biol. 2006; 328: 177-85.
- Tian Y, **Zhang H.** Isolation of proteins from formalin-fixed tissue for subsequent mass spectrometric analysis. Current Protocols in Molecular Biology. 2010; 10: Unit 10 26 1-7.
- Li Y, **Zhang H.** High throughput analysis of glycoproteins from plasma. Simpson RJ, Greening D (eds), Serum and Plasma Proteomics in Methods Mol Biol. 2011; 728: 125-133.
- Mertz JL, Toonstra C, Zhang H. Chemical strategies to glycoprotein analysis. 2019; 293-316.

## Editorials [ED]

- **Zhang H**, Cotter RJ. Glycoproteomics: New Technology Developments and Applications Provide Renewed Interest in Glycoproteins. *Clinical Proteomics.* 2008; 2: 1-4.
- Ellis MJ et al. & **NCI CPTAC Investigators**. Connecting Genomic Alterations to Cancer Biology with Proteomics: The NCI Clinical Proteomic Tumor Analysis Consortium. Cancer Discovery. 2013; 3 (10): 1108-1112.
- Van Eyk JE, Corrales FJ, Aebersold R, Cerciello F, Deutsch EW, Roncada P, Sanchez J-C, Yamamoto T, Yang P, **Zhang** H, Omenn GS. Highlights of the biology and disease-driven Human Proteome Project, 2015-2016. J Proteome Res. 2016; 15: 3979-3987.

# Methods and Techniques [MT]

- Tian Y, Zhou Y, Elliott S, Aebersold R, **Zhang H.** Solid-phase extraction of N-linked glycopeptides. Nat Protoc. 2007; 2: 334-339. [SI/QI]
- **Zhang H.** Glycoproteomics. Unit 24.3 in Current Protocols in Protein Science. Coligan JE, Dunn BM, Speicher DW, Wingfield PT. (eds), John Wiley Sons Inc, Hoboken, NJ, 2007
- Yang S, **Zhang H.** Glycomic analysis of glycans released from glycoproteins using chemical immobilization and mass spectrometry. Curr Protoc in Chem Biol. 2014; 6: 191-208.
- Whiteaker JR et al. & **NCI CPTAC Investigators**. CPTAC Assay Portal: a repository of targeted proteomic assays. Nature Methods. 2014; 11(7): 703-704. Steering Committee
- Yang S, Hu Y, Sokoll L, **Zhang H**. Simultaneous quantification of N- and O-glycans using a solid-phase method. Nat Protoc. 2017; 12: 1229-1244. [SI/QI]
- Mertins P, Tang LC, Krug K, Clark DJ, Gritsenko MA, Chen L, Clauser KR, Clauser TR, Shah P, Gillette MA, Petyuk VA, Thomas SN, Mani DR, Mundt F, Moore RJ, Hu Y, Zhao R, Schnaubelt M, Keshishian H, Monroe ME, Zhang Z, Udeshi ND, Mani D, Chan DW, Smith RD, **Zhang H**, Liu T and Carr SA. Reproducible workflow for multiplexed deep-scale analysis of the proteome and phosphoproteome of tumor tissues by liquid chromatography-mass spectrometry. Nature Protocols. 2018; 13: 1632-1661.
- Yang W, Song A, Ao M, Xu Y, **Zhang H.** Large-scale Mapping of Site-specific O-GalNAc Glycoproteome. Nature Protocols. 2020; July 17: 1-22.

### Other Media [OM]

The Center for Biomarker Discovery and Translation and its Role in Clinical Proteomics in Pathology. <a href="http://apps.pathology.jhu.edu/blogs/pathology/the-center-for-biomarker-discovery-and-translation-and-its-role-in-clinical-proteomics-in-pathology">http://apps.pathology.jhu.edu/blogs/pathology/the-center-for-biomarker-discovery-and-translation-and-its-role-in-clinical-proteomics-in-pathology</a>

https://www.hupo.org/2016/02/news/capture-all-and-then-release-only-the-de-n-glycosylated-peptides-for-glycoproteomics/

https://www.hupo.org/2016/06/news/clinical-proteomics-receives-impact-factor-of-3-476/

https://www.youtube.com/watch?v=m4U9bcnGtg4&feature=youtu.be

http://www.biomarkercenter.org/

## **FUNDING**

### **EXTRAMURAL** Funding

#### Current

## Research Extramural Funding

06/01/22-05/31/27 Proteogenomic Characterization of Tumor Tissues and Preclinical Models with High

Precision

U24CA271079

NIH/NCI

PI: Zhang, Chan

Notes: The overarching goal of this PCC is to comprehensively characterize additional human tumor and normal tissues as well as preclinical models provided by the NCI in order to systemically identify, prioritize, and develop targeted proteomic assays to confirm cancer-related proteins, protein modifications, and protein interaction networks and to provide public data resources for the scientific community.

04/01/19-03/31/24 Modulating Antigenic and Immunogenic Properties of HIV ENV by Altering Signal Sequence

R01AI40909-01A1

NIH/NIAID Subgrant from ICAHN School of Medicine at Mt.Sinai/NIAID

PI: Upadhyay

Notes: This study focusses on investigating the role of HIV-1 envelope signal sequence in altering its capacity to induce functional antibodies targeting the V1V2 epitopes.

09/01/10 - 3/31/23 Glycoprotein biomarkers for the early detection of aggressive prostate cancer

U01CA152813

NIH/NCI

PI: Zhang

Notes: This is a biomarker developmental laboratory (BDL) of the Early Detection Research Network (EDRN). The goal of this project is to identify glycoprotein biomarkers for the early detection of aggressive prostate cancer in tissues and urine.

08/01/16 - 04/30/27 Clinical Resources for Alcoholic Hepatitis Investigations

1R24AA025017

NIH/Alcohol Research Resource Awards

PI- Sun

Notes: The aims are to develop a clinical resource of severe alcoholic hepatitis that serves the alcohol research community.

## Previous funding in last 3 years

4/1/16 - 3/31/23 Biomarkers for the detection of early stage or low-volume ovarian cancer

Agency: NIH/NCI/U01 PI: Zhang Z, Shih I-M

Notes: Development of *in vitro* diagnostic multivariate index assay using liquid-based cervical cytology specimen and/or serum/plasma biomarkers for the detection of early stage or low-volume ovarian cancer

volume ovarian cancer

9/16/16 - 8/31/22 The comprehensive proteome characterization center at Johns Hopkins: high precision

discovery and confirmation of genoproteomic targets.

U24CA210985

NIH/NCI

PI: Chan D, Zhang H, and Zhang Z

Notes: This is a center for the Clinical Proteomic Tumor Analysis Consortium (CPTAC). The major goal of this Center is the proteomic characterization of tumors with genomic data to not only verify the genomic alterations at the protein level but also allow for the analysis of unique features that are inherent to proteins including post-translational modifications.

4/1/16 - 3/31/23 Clinical and analytical validation of cancer biomarkers

U24CA115102

NIH/NCI PI: Chan DW

Notes: This is a biomarker reference laboratory (BRL) of the Early Detection Research Network (EDRN). The major goal of this project is to conduct analytical and clinical validation studies of biomarkers

9/1/19 - 8/31/21 Discovery of Urinary Glycoprotein biomarkers for aggressive prostate cancer

Patrick C. Walsh Fund PI: Zhang H and Alan Partin

Notes: The goal of this project is to develop urinary biomarkers for aggressive prostate cancer.

05/01/18-04/30/20 Adaptive process control and advanced sensing for robust mAb glycan quality

NIIMBL

PIs: Betenbaugh M and Zhang H

Johns Hopkins University, Baltimore, MD.

instruction, JHUSOM

Notes: The objective of this project are to develop fast glycomic methods to sense mAb glycan quality for mAb production process control.

## **EDUCATIONAL ACTIVITIES** (in chronological order, earliest first by start date under each subcategory)

#### **Educational Focus**

I am very interested in teaching undergraduate and graduate courses in the areas of Biochemistry and laboratory, General Chemistry and laboratory, Bioanalytical Chemistry and laboratory, Instrumental Methods for Analytical Chemistry, Separation Methods, Protein Chemistry, and Nucleic Acids. I developed materials and lectured in courses in Fundamentals of Glycobiology, Techniques in Glycobiology, and The Role of Chromatography and Mass Spectrometry in Biological Research. In the future, I would like to develop new courses such as Structural and Functional Analysis of Glycoproteins, or Carbohydrate Chemistry and Biochemistry. Of course, it would be fun to work with students closely by teaching Seminar and Research Courses.

The Role of Chromatography and Mass Spectrometry in Biological Research, co-instructor for course

What we learn from proteome and glycoproteomic analysis. Lecturer of 4th Annual Pathology Symposium,

Fundamentals of Glycobiology, co-instructor for course ME340.709 with Dr. Gerald Hart, et al -classroom

# **Teaching**

2007, fall

2016, fall

2017, spring

#### Classroom instruction

,	ME330.804 with Dr. Robert Cotter - classroom instruction, JHUSOM
2008, fall	The Role of Chromatography and Mass Spectrometry in Biological Research, co-instructor for course
	ME330.804 with Dr. Robert Cotter - classroom instruction, JHUSOM
2009, fall	The Role of Chromatography and Mass Spectrometry in Biological Research, co-instructor for course
	ME330.804 with Dr. Robert Cotter - classroom instruction, JHUSOM
2010, fall	The Role of Chromatography and Mass Spectrometry in Biological Research, co-instructor for course
	ME330.804 with Dr. Robert Cotter - classroom instruction, JHUSOM
2012, fall	Fundamentals of Glycobiology, co-instructor for course ME340.709 with Dr. Gerald Hart, et al -classroom instruction, JHUSOM
2012, fall	Mass Spectrometry in an "Omics" World, co-instructor for course ME330.804 - classroom instruction,
	JHUSOM
2013, fall	Techniques in Glycobiology, lecturer and laboratory instructor for course ME340.710 -classroom
	instruction and laboratory instruction, JHUSOM
2014, spring	Mass Spectrometry, lecturer for course "Pharmacology Tutorials organized by Dr. Heng Zhu et al
	classroom instruction, JHUSOM
2015, spring	Fundamentals of Glycobiology, co-instructor for course ME340.709 with Dr. Gerald Hart, et al -classroom
	instruction, JHUSOM
2015, spring	Analysis of Glycoproteins, lecturer in Mass Spectrometry Class, University of Maryland, School of
	Pharmacy
2015, summer	Techniques in Glycobiology, lecturer and laboratory instructor for course ME340.710 -classroom
	instruction and laboratory instruction, JHUSOM
2016, fall	Proteomics and Cancer Biology; Glycoproteomics and Cancer Biology, lectures for graduate course, Fudan
	University, Shanghai, China

2017, summer Techniques in Glycobiology, lecturer and laboratory instructor for course ME340.710 -classroom

instruction and laboratory instruction, JHUSOM

2019, Summer K12\_Glycobiology. co-instructor with Drs. Natasha Zachara, Gerald Hart, and Ronal Schnaar -classroom

instruction, JHUSOM

### CME instruction

Regional

O4/09 Glycoproteins in Cancer Diagnosis, lecturer to Pathology Grand Rounds – CME instruction, JHUSOM Proteomics and Glycoproteomics: What We learn Beyond Genomics, lecturer to Pathology Grand Rounds

- CME instruction, Johns Hopkins University School of Medicine

## Workshops /seminars

National

2/09 Lecturer, mainly graduate or post-graduate, Glycoproteomics Short Course, 5th US HUPO Annual Conference, San Diego, CA

3/13 Organizer and lecturer, Glycoproteomics Short Course, 9th US HUPO Annual Conference, Baltimore,

### **Mentoring**

Shared publications with mentees are indicated by the publication entries in this CV

# Pre-doctoral Advisees / Mentees

Graduate	Charl	lamba
Graduate	Stuc	lemis

03/19-05/20 **Zhichen Liu**, Present Position: Graduate Student, MS; Johns Hopkins University

Awards/grants/degrees received: Master Degree, Research training; Mass spectrometry, Proteomics,

glycoproteomics, data analysis, database for glycoproteomic data. Move to position: PhD candidate, Johns Hopkins University

03/17-05/20 **Bay Xu**, Present Position: Graduate Student, MS; Johns Hopkins University

Awards/grants/degrees received: Master Degree, Research training; Proteomics and glycoproteomics.

Move to position: PhD candidate, Johns Hopkins University

03/17-12/19 Qiong Wang, Present Position: Graduate Student, PhD. Candidate; Johns Hopkins University

Awards/grants/degrees received: Ph.D degree, Research training; Proteomics and glycomics

Move to position: FDA

10/17-present Sean A. Ponce, Present Position: Graduate Student, PhD. Candidate; Johns Hopkins University

Awards/grants/degrees received: Research training; Proteomics and glycomics

02/18-06/20 Yangying Zhou, Present Position: Exchange Graduate Student, Ph.D Candidate; Johns Hopkins

University

Awards/grants/degrees received: Ph.D degree, Research training; Proteomics and glycomics

Move to position: Clinician, Xianya Hospital, Changsha, Hunan, China

04/18-present Shao-Yung Chen, Present Position: Graduate Student, PhD. Candidate; Johns Hopkins University

Awards/grants/degrees received: Research training; Proteomics and glycomics.

03/17-02/18 Rui Zhang, Present Position: Exchange Graduate Student, Ph.D Candidate; Johns Hopkins

University

Awards/grants/degrees received: Research training; Proteomics and glycomics

02/18-11/18 **Ce Wang**, Present Position: Exchange Graduate Student, Ph.D Candidate; Johns Hopkins University

Awards/grants/degrees received: Research training; Proteomics and glycomics

11/09-12/11 **Deniz Baycin-Hizal,** Ph.D, Position: Graduate Student, Ph.D Candidate; Johns Hopkins University

Awards/grants/degrees received: 11/09 Pre-doctoral training; Research training

12/11 Ph.D degree

12/11 Moved to Position: Scientist, MedImmune

6/11-05/16	<ul> <li>Shadi Toghi Eshghi, Ph.D, Position: Graduate Student, Ph.D Candidate; Johns Hopkins University Awards/grants/degrees received: Pre-doctoral training; Research training; Thesis project</li> <li>4/12 14th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD</li> <li>11/14 Travel Award. The Glycobiology for Society Annual Meeting, Honolulu, Hawaii.</li> <li>4/15 17th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD</li> <li>8/15 Named a Class of 2016 Siebel Scholar at Johns Hopkins University to receive the award from the Siebel Foundation</li> <li>3/16 Ph.D degree</li> <li>4/16 Moved to position: Scientist, Genentech</li> </ul>			
07/11-05/13	Namita Trikannad, MS, Position: Graduate Student, MS Candidate; Johns Hopkins University Awards/grants/degrees received: MS, 05/13 Pre-doctoral training; Research training 05/13 Moved to Position: Research Scientist, Assay Development and Screening Tri-Institutional Therapeutics Discovery Institute (Weill Cornell Medicine) New York			
Research Associates	3			
10/17-10/2020	Weiming Yang, Ph.D.; Present Position: Research Associate, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, mass spectrometry Move to position: Scientist, NIH, Bethesda, Maryland			
Research Assistants 7/14-present	Naseruddin Höti, Ph.D.; Present Position: Senior Research Specialist, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics			
2/17-present	<b>Lijun Chen</b> , Ph.D; Present Position: Senior Research Specialist, Johns Hopkins University Awards/grants/degrees received: Research training; Mass spectrometry			
09/17-present	Michael Schnaubelt, Research Data Analyst, Johns Hopkins University, School of Medicine, Pathology Department, Clinical Chemistry Division Awards/grants/degrees received: Research training; Proteomics and glycomics			
12/17-05/20	<b>Minghui Ao</b> , Research Specialist II, Johns Hopkins University, School of Medicine, Pathology Department, Clinical Chemistry Division Awards/grants/degrees received: Research training; Proteomics and glycomics			
05/18-01/20	Angellina Song; Present Position: Laboratory Technologist, Johns Hopkins University Awards/grants/degrees received: Research training; Mass spectrometry Move to position: Scientist, Johns Hopkins University			
10/18-06/20	Rodrigo Vargas Eguez; Present Position: Laboratory Technologist, Johns Hopkins University Awards/grants/degrees received: Research training; Mass spectrometry Move to position:			
02/11-11/17	Jered Pasay, Laboratory Technologist, Johns Hopkins University, School of Medicine, Pathology Department, Clinical Chemistry Division.  Moved to Position: Maryland Board of Pharmacology  Awards/grants/degrees received: Research training; Proteomics and glycomics  Move to position:			
06/11-05/14	Robert Harlan, Senior Research Specialist Moved to Position: Laboratory Manager, Johns Hopkins University, School of Medicine, The Center for Resources in Integrative Biology Awards/grants/degrees received: Research training; Proteomics and glycomics			

08/11-03/13 Caitlin H. Choi, Research Coordinator, Johns Hopkins University, School of Medicine, Pathology

Department, Clinical Chemistry Division

Moved to Position: MD student, ENS, MC, USNR, Class of 2017

Awards/grants/degrees received: Research training; Proteomics and glycomics

### Post-doctoral Advisees / Mentees

5/08

1/09

Post-doctoral Ac	dvisees /Mentees				
02/13-07/18	Stefani Thomas, Ph.D.; Present Position: Clinical Chemistry Fellow, Johns Hopkins University Awards/grants/degrees received: Proteomics, glycomics, mass spectrometry 4/13 15th Annual Department of Pathology Young Investigators' Day Award, Baltimore, 2/16 Young Investigator Award, Mass Spectrometry Applications to the Clinical Labs (MSACL), Palm Springs, CA Move to position: Assistant professor, University of Minnesota				
09/15-present	Yingwei Hu, Ph.D.; Present Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology 4/17 19th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD				
12/15-present	Joseph Mertz, PhD.; Present Position: Post-doctoral fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology				
03/16-present	David Clark, PhD.; Present Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology 4/17 19th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD 7/20 22d Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD				
03/17-11/19	<b>Kyung-Cho Cho</b> , Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position:				
09/17-present	<b>Jianbo Pan</b> , Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position:				
03/18-present	Tung-Shing Lih, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics				
08/18-present	Mingming Dong, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position:				
02/19-present	Liwei Cao, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics				
04/19-present	Yuefan Wang, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics				
05/05-11/06	Yong Zhou, Ph.D.; Position: Research Fellow, Scientist, Institute for Systems Biology Awards/grants/degrees received: Postdoctoral training; Promoted to Scientist 11/06 Moved to position: Scientist, Institute for Systems Biology				
11/05-09/13	Yuan Tian, Ph.D.; Position: Research Fellow, Research Associate, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training 1/08 Seed Grant from HERA Foundation 4/08 10th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD				

into Clinical Diagnostics, Seattle, WA

Seed Grant from HERA Foundation

Best poster Award, 2<sup>nd</sup> place, AACC conference on Translating Proteomic Discoveries

	4/09 10/09	11th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD Outstanding Poster Award, 3 <sup>rd</sup> place. AACC (American Association of Clinical Chemistry) - Translating Novel Biomarker to Clinical Practice, Bethesda, MD
	3/10	Travel Award, US Human Proteome Organization 6th Annual Conference in Denver, CO
	4/10	12th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD
	3/11	Travel Award, US Human Proteome Organization 7th Annual Conference in Raleigh, NC
	3/11	Promoted to a faculty as a Research Associate of Pathology at Johns Hopkins
	10/13	Moved to Position: Director of Glycoproteomics Core Facility, University of Mississippi
		Medical Center
3/07-6/11	<b>Yan Li</b> , Pl	n.D.; Position: Research Fellow, Research Associate, Johns Hopkins University
	Awards/gr	rants/degrees received: Postdoctoral training
	7/09	Prostate Cancer Fellowship Training Award from DOD-CDMRP, 7/2009-6/2011.
	9/09	Young Investigator Awards: Human Proteome Organization 8rd Annual World Congress.
	10/00	Toronto, Canada.
	10/09	Outstanding Poster Award, 1st place. AACC (American Association of Clinical Chemistry)
	9/10	- Translating Novel Biomarker to Clinical Practice. Bethesda, MD.
	9/10	Young Investigator Awards: Human Proteome Organization 9th Annual World Congress. Sydney, Australia
	11/10 Tra	vel Award of the Society of Glycobiology annual conference.
	2/11	Promoted to Faculty as a Research Associate of Pathology at Johns Hopkins
	7/11	Moved to Position: Professor of Chinese Academy of Biophysics Institute
	,	The state of the s
8/07-12/09		Ph.D.; Position: Clinical Chemistry Fellow, Johns Hopkins University
		rants/degrees received: 7/07 Postdoctoral training.
	1/10	Promoted to a faculty position as an Instructor of Pathology at Johns Hopkins.
	12/11	Promoted to Assistant Professor at Johns Hopkins
	8/12	Moved to Position: Assistant Professor of Lab Medicine and Pathology, Director of
		Clinical Chemistry, University of Minnesota Medical Center
8/10-7/12		rzinke, Ph.D.; Position: Clinical Chemistry Fellow, Johns Hopkins University
		rants/degrees received: Postdoctoral training; Proteomics, mass spectrometry training to
		er for disease biomarker discovery
	4/12	14th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD
	8/12	Moved to Position: Assistant Professor of Pathology, Johns Hopkins University
6/11-8/13	_	n Wang, Ph.D.; Position: Research Fellow, Johns Hopkins University
	-	rants/degrees received: Postdoctoral training; Glycobiology, glycomics, mass spectrometry
	11/11	Travel Award, Glycobiology Society Annual Meeting, Seattle, WA
	10/13	Moved to Position: Scientist, NIAID, NIH
9/11-6/15	Jing Chen	, Ph.D.; Position: Research Fellow, Johns Hopkins University
	Awards/gr	rants/degrees received: Postdoctoral training; Clinical proteomics, cell biology, molecular
		ycomics, mass spectrometry
	11/13	Travel Award. The Society for Glycobiology Annual Meeting, St. Petersburg, FL.
	4/15	17th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD
	7/15	Moved to position: Scientist, Agilent
2/12-9/15	Paul Aiye	tan, MS, M.D.; Position: Research Fellow, Johns Hopkins
	Awards/gr	rants/degrees received: Postdoctoral training; Proteomics, glycomics, bioinformatics
	4/12	14th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD
	10/15	Postdoctoral fellow, JHU
	3/16	Moved to position: Scientist (Research and Development), Biotech Groups' Diagnostics
		Laboratory  Convent position Picinformatics Scientist Frederick National Laboratory for
		Current position: Bioinformatics Scientist, Frederick National Laboratory for Cancer Research
		Carreet resourch
3/12-8/15	Jian-Ying	Zhou, Ph.D.; Position: Research Associate, Johns Hopkins University

	Awards/grants/degrees received: Glycobiology, cell biology, mass spectrometry, proteomics, glycomics 9/15 Moved to Position: Scientist, Abbott					
3/14-8/15	Xingwang Jia, Ph.D.; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, clinical chemistry Moved to position: Clinical Chemist, 301 Hospital, Beijing, China					
6/14-10/15	Yuri Poluektov, Ph.D.; Position: Research Fellow, Johns Hopkins Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, bioinformatics, immunology 5/16 Moved to position: Fellow, FDA					
5/15-5/16	Lingquan Deng, Ph.D.; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology 12/15 Travel Award. The Society for Glycobiology Annual Meeting, San Francisco, CA 5/16 Moved to position: Scientist, GlycoMimetics					
1/11-11/16	<ul> <li>Shuang Yang, Ph.D.; Position: Research Associate, Johns Hopkins University</li> <li>Awards/grants/degrees received: Postdoctoral training; Proteomics, glycoproteomics, mass spectrometry training to build career for technology development for glycan separation and analysis</li> <li>11/11 Travel Award, Glycobiology Society Annual Meeting, Seattle, WA.</li> <li>4/12 14th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD</li> <li>9/12 Travel Award. 11th World Conference of HUPO, Boston, MA</li> <li>9/12 Best Presentation Award, 3rd place. 11th World Conference of HUPO, Boston, MA</li> <li>5/13 Best Poster Award. NIH Glycosciences Research Day, Bethesda, MD.</li> <li>6/13 Travel Award, 22nd International Symposium on Glycoconjugates, Dalian, China</li> <li>10/13 Promoted to junior faculty position as Research Associate</li> <li>11/16 Moved to position: Staff, Laboratory of Bacterial Polysaccharides, Center for Biological Evaluation and Research, Food and Drug Administration</li> </ul>					
09/11-01/17	Shisheng Sun, Ph.D.; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, mass spectrometry 9/12 Travel Award. The 11th World Conference of HUPO, Boston, MA 11/13 Travel Award. The Society for Glycobiology Annual Meeting, St. Petersburg, FL. 3/16 Travel Award. USHUPO, Boston, MA 4/16 18th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD Moved to position: Professor, Northwestern University, Xian, China					
09/15-04/17	Yang Liu, Ph.D.; Position: Research Technologist, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology Moved to position: Research Technologist, Genomic Analysis, CA.					
11/11-06/17	Punit Shah, Ph.D.; Position: Laboratory Manager, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, mass spectrometry 4/13					
06/16-06/17	Jian-Liang Zhou, Ph.D.; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Moved to position: Associate Professor of Pharmacy, Department of Traditional Chinese Medicines, Zhejiang Institute for Food and Drug Control, China					
11/11-09/17	Weiming Yang, Ph.D.; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Postdoctoral training; Proteomics, glycomics, mass spectrometry Move to position: Research Associate, Johns Hopkins University					

11/12-02/17	Lijun Chen, Ph.D; Position: Research Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Mass spectrometry
11/15-06/18	Ventzi Hristova, PhD.; Present Position: Clinical Chemistry Fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics, glycomics, and cell biology 4/17 19th Annual Department of Pathology Young Investigators' Day Award, Baltimore, MD Move to position: Scientist,
03/17-06/18	<b>Quanzi Ouyang</b> , Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position:
02/17-07/18	Christian Toonstra, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position: Postdoctoral fellow, University of Minnesota
01/17-12/18	Ganglong Yang, Ph.D.; Present Position: Research fellow, Johns Hopkins University Awards/grants/degrees received: Research training; Proteomics and glycomics Move to position:

### Thesis committees

2013-2015	David Clark, PhD candidate, University of Maryland, School of Dentistry, dissertation committee member, graduated with Ph.D
2014-2016	Chris Mitchell, PhD candidate, Johns Hopkins University, Department of Biochemical, Cellular &
	Molecular Biology, dissertation committee chair, graduated with Ph.D
2015-2018	Christopher Saeui, PhD candidate, Johns Hopkins University, Department of Biomedical Engineering,
	dissertation committee member
2015-present	William Fondrie, PhD candidate, University of Maryland, School of Dentistry, dissertation committee
	member.
2017-present	Srona Sengupta, PhD candidate, Johns Hopkins University, Immunology Program, dissertation committee member.

# Educational Program Building / Leadership

2007-present	Sponsor post-doc training grant applications and serves as advisor and sponsor of applications of
	postdoc fellowship

4/1/08 - 3/31/10	Identification of subtype-specific extracellular proteins from ovarian tumors
	LIED A Foundation

HERA Foundation

PI: Tian Y

Role: Mentor, 0%; Notes: The objective of this project is to identify extracellular proteins in different subtypes of ovarian tumor and determine the tissue-specificity.

# 7/1/09 - 6/30/11 Glycoproteomic profiling of prostate cancer for biomarker discovery

DoD-W81xwh-09-1-0136

DOD-CDMRP

PI: Li

Role: Mentor, 0%; Notes: The objective of this project is to determine the glycoprotein biomarkers to predict the outcome of the prostate cancer using glycoproteomic

## **RESEARCH ACTIVITIES**

## **Research Focus**

My research interests are centered on the structural and functional analyses of proteins and multiple protein modifications by glycosylation, phosphorylation, and acetylation. Recently, our lab has developed several novel glycoproteomic and glycomic technologies to study protein glycosylation and applied the technologies to study cell surface glycoproteins and glycoproteins secreted to body fluids. These technologies enable the identification and quantification of glycoproteins, glycosites, glycans associated with each glycosite and their occupancies. Researchers in the group participate in several national programs in characterization of glycoproteins including the Early Detection Research Network (EDRN), Clinical Proteomic Tumor Analysis Consortium (CPTAC), and Programs of Excellence in Glycosciences (PEG).

## Research Program Building / Leadership

2/07-present Mass Spectrometry Facility for the Center for Biomarker Discovery and Translation (CBDT) at Johns Hopkins: I led the development, established the Facility, serve as the director of the Facility.

## Inventions, Patents, Copyrights

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- 02/27/07 Aebersold R, **Zhang H.** Methods for quantitative proteome analysis of glycoproteins. United States patent #7,183,118.
- 04/03/07 Rush J, **Zhang H**, Zha X, Comb MJ, Tan Y. Immunoaffinity isolation of modified peptides from complex mixtures. United States patent # 7,198,896.
- 08/21/07 Comb MJ, Tan Y, **Zhang H.** Production of motif-specific and context-independent antibodies using peptide libraries as antigens. United States patent # 7,259,022.
- Rush J, **Zhang H**, Zha X, Comb MJ, Tan Y. Immunoaffinity isolation of modified peptides from complex mixtures. United States patent # 7,300,753.
- 09/14/10 Aebersold R, **Zhang H.** Affinity capture of peptides by microarray and related method. United States patent # 7,794,947.
- **Zhang H**, Li Y, Sokoll LJ, Zhang Z, Chan DW. Biomarkers for prostate cancer. United States patent #8,603,734.
- Rush J, **Zhang H**, Zha X, Comb MJ, Tan Y. Immunoaffinity isolation of modified peptides from complex mixtures. United States patent # 9,587,013.
- O3/14/17 **Zhang H**, Tian Y, Chan DW, Chen J. Biomarkers for aggressive prostate cancer. United States Patent # 9,594,086.
- 02/13/18 Li D, Chan DW, **Zhang H**, Wang X. Biomarkers for aggressive prostate cancer. United States patent # 9,891,224.
- O4/10/18 **Zhang H**, Yang S, Li S. Isobaric aldehyde-reactive tags for and analysis of glycans using same. United States patent # 9,939,444.
- 04/09/19 Li S, Zhang H, Yang S. Quaternary ammonium containing isobaric tag for quantitative glycan profiling. US Patent # 10,254,291.

## **Pending**

- 6/18/02 Comb MJ, Tan Y, **Zhang H.** Positive identification of phospho-proteins using motif-specific, context-independent antibodies coupled with database searching. WO Patent 2003107003.
- 5/21/04 Aebersold R, **Zhang H.** Compositions and methods for quantification of serum glycoproteins. WO Patent 20051142211.
- **Zhang H**, Aebersold R. Tissue-and serum-derived glycoproteins and methods of their use. WO Patent 2007047796.
- 7/25/08 **Zhang H**, Meany DL, Chan DW, Zhang Z, Li Y, Sokoll LJ. Detection of prostate cancer using PSA glycosylation patterns. Pending US application 20140193832, WO Patent 2010011357.
- **Zhang H**, Yang S. Glycan and glycopeptide capture and release using reversible hydrazone-based method. **Pending US application** 20140135235, WO Patent 2012170491.
- 9/10/13 **Zhang H** et al. Methods for quantitative analysis of glycans and glycosylation sites on glycoproteins using solid-phase extraction of glycopeptides and glycans (SPEGAG). Pending PCT application WO #2014/040,072.
- 2/10/13 **Zhang H** et al. Glycomic analysis by glycoprotein immobilization for glycan extraction and liquid chromatography on microfluidic chip. Pending PCT application WO 2014/040,066.
- 10/9/13 Veltri R, **Zhang H**, Christudass C, Liu Z, Epstein JI, Carter HB. Active surveillance biomarkers for assessing cancer patients for treatment. Provisional applications filed.
- 6/30/14 Li D, **Zhang H**, Chan DW. Serum angiogenic factors as prognostic biomarkers in prostate cancer. Pending US application 14/375,535.
- **Zhang H** et al. Solid phase extraction of global peptides, glycopeptides, and glycans using chemical immobilization in a tip. Pending PCT/US application #2014/058,087.
- **Zhang H**, Toghi Eshghi S. Mass spectrometry imaging of glycans from tissue sections and improved analyte detection method. Pending US application 14/402,478.

## SYSTEM INNOVATION AND QUALITY IMPROVEMENT ACTIVITIES

#### System Innovation Focus

In mass spectrometry core facility, we aim to develop proteomics, glycoproteomics, and mass spectrometry technologies and apply to clinical specimens by establishing standard operating procedures, automated robotic protocols, and data quality control

## System Innovation and Quality Improvement efforts within JHMI:

2013 Establishing SOPs for sample preparation for proteomics and glycoproteomics, data acquisition by mass spectrometry, and data analysis

### **ORGANIZATIONAL ACTIVITIES**

## Institutional Administrative Appointments

2015-2016 Member, Search Committee on Endowed Chairs of Bloomberg Distinguished Professorships

### **Editorial Activities**

2008 Guest Editor of 2 Special Issues of Clinical Proteomics on Glycoproteomics

2013 Guest Editor of Special Issue of Clinical Proteomics on Glycoproteomics and Glycomics 2017 Guest Editor of Special Issue of Proteomics-Clinical Applications on Glycoprotein Analysis

## Editorial Board appointments

2008-present Member, Editorial Board of Clinical Proteomics

2011-present Associate Editor of Clinical Proteomics

## Journal peer review activities

2003 - Present

6/15

Journal of Proteome Research, Molecular & Cellular Proteomics, Rapid Communications in Mass Spectrometry, Nano Letters, Nature Protocols, Nature Biotechnology, Molecular BioSystems, Expert Reviews of Proteomics, Bioinformatics, The Analyst, Mass Spectrometry Reviews, Cancer Research, Journal of Proteomics, Biomarkers, Theronostics, PLOS ONE, Analytical Chemistry, Clinical Proteomics, Proteomics, Proteomics-Clinical Applications

## Advisory Committees, Review Groups/Study Sections

2012-2015	Member of Board of Directors, United States Human Proteomic Organization (USHUPO)
2012-present	Chair of Cancer Human Proteome Project (Ca-HPP) of Human Proteomic Organization (HUPO)
2015-2019	Member of Executive Committee for World Human Proteomic Organization (HUPO) Biology/Disease
	Human Proteome Project (B/D-HPP)
2015-2018	Council Member of World Human Proteomic Organization (HUPO)
2015-present	Associate Chair of Member Connection Committee, Chinese American Society for Mass Spectrometry (
	CASMS)
2/09	Member, NIH Review Panel for Studies of Antimicrobial and Prebiotic Activity of
	Oligosaccharides (R01)
7/10	Member, Technical Evaluating Panel-Reviewer for the Interagency Agreement
11/10-07/15	Member, NIH Early Detection Research Network Standing Review Committee for Associate Membership
	applications to review application proposals three times a year
2/11	Reviewer, Netherlands Genomics Initiative-The Zenith project on Genomics and/or
	Bioinformatics
2011-2012	Reviewer, NIH small business SBIR-STTR
10/13	Member, PLCO Special Emphasis Panel/Scientific Review Group 2014/01 ZCA1 SRLB-Y (J1) B
6/14	Member, PLCO Special Emphasis Panel/Scientific Review Group 2014/10 ZCA1 TCRB-Y (01)
	S
7/14	Member, NIH IMST-B (40) P: A Resource for Biomedical Mass Spectrometry (Site Visit at
	Washington University for Renewal Application)
9/14	Reviewer, the Keck Foundation

Research Network Consortium on Beta Cell Death and Survival (HIRN-CBDS)

Member, review panel for 2015/10 ZDK1 GRB-S (O3) S - RFA-DK-14-021: Human Islet

6/15	Member, review panel for NIH IMST-M (41), Bio-Organic Biomedical Mass Spectrometry
	(renewal application by University of California at San Francisco)
7/15	Member, Technical Evaluating Panel-Reviewer for the Interagency Agreement
3/16	Member, Panel reviewer for the NIH common fund for novel and innovative tools to facilitate
	identification, tracking, manipulation, and Analysis of Glycans and their functions
6/16	Mail Reviewer, Cellular Aspects of Diabetes and Obesity [CADO] Study Section
6/16	Reviewer, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session
7/16	Reviewer, PLCO Special Emphasis Panel/Scientific Review study session
4/17	Panel reviewer for the NIH common fund for novel and innovative tools to facilitate identification,
	tracking, manipulation, and Analysis of Glycans and their functions
06/18	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
10/18	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
02/19	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
06/19	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
10/19	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session,
	Washington DC.
02/20	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session,
	Washington DC.
06/20	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
10/20	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
02/21	Reviewer, NIH Innovative Molecular Analysis Technologies (IMAT) program.
02/21	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
06/21	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
10/21	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.
06/22	Standing Member, Enabling and Bioanalytical and Imaging Technologies (EBIT) study session.

# Professional Societies

2003-present	American Society for Mass Spectrometry (ASMS)
2006-present	United States Human Proteomic Organization (USHUPO)
2009-present	Society for Glycobiology
2010-present	American Chemical Society (ACS)
2010-present	World Human Proteomic Organization (HUPO)
2015-present	America Association for Cancer Research (AACR)
2015-present	Chinese American Society for Mass Spectrometry (CASMS)

# Conference Organizer (separate into JHMI/Regional - National - International activities)

## National activities

3/08	Organizing Committee Member: 4th Conference of US Human Proteome Organization (USHUPO), Bethesda, MD
3/13 2015	Organizing Committee Member: 9th US HUPO Annual Conference, Baltimore, MD Monday Networking Night, Chinese American Society for Mass Spectrometry
08/21	Session organizer and chair: Chinese American Society for Mass Spectrometry.

## International activities

2012-2019	Organizing World Human Proteomic Organization (HUPO) Cancer Biology/Disease Human Proteome
	Project (Cancer B/D-HPP)

2018 Organizing Committee. HPLC2018, 47th International Symposium on High Performance Liquid Phase Separations and Related Techniques, Washington DC.

Session Chair (separate into JHMI/Regional – National - International activities)

## National activities

01/06	Session chair: BuzZ session. 6th Annual Plasma Proteome, San Diego, CA
03/08	Session organizer and chair: The 4th Conference of USHUPO, Bethesda, MD
03/10	Session chair: 6th Conference of USHUPO Denver CO

03/12	Session organizer and chair: 8th Conference of USHUPO, San Francisco, CA
03/13	Session organizer and chair: 9th USHUPO Annual Conference, Baltimore, MD
03/13	USHUPO short course organizer on "Glycoproteomics and glycomics", 9th US HUPO Annual
	Conference, Baltimore, MD
09/16	Session chair, Glycoproteomics - Technical limitations & Prospects. The 15th Human Proteome
	Organization World Congress (HUPO), Taipei, Taiwan
03/18	Session chair, Glycoproteomics and glycomics for clinical biomarker discovery. USHUPO2018,
	Minneapolis, MN.
07/18	Co-organizer and Session chair: HPLC2018, 47th International Symposium on High Performance
	Liquid Phase Separations and Related Techniques, Washington DC.
07/20	Session organizer and chair: Career Development. American Society for Mass Spectrometry
03/21	Session organizer and chair: Protein post-translational modifications. US Human Proteome
	Organization (USHUPO).
International a	activities
International a 09/12	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of
	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA
09/12	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of
09/12	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO,
09/12 09/13	Session leader: Cancer Proteomics and Human Proteome Project, 11 <sup>th</sup> World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12 <sup>th</sup> World Conference of HUPO, Yokohama, Japan
09/12 09/13	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO, Madrid, Spain
09/12 09/13 10/14	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO,
09/12 09/13 10/14	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO, Madrid, Spain Session chair: Proteomics. 43rd International Symposium on High Performance Liquid Phase
09/12 09/13 10/14 09/15	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO, Madrid, Spain Session chair: Proteomics. 43rd International Symposium on High Performance Liquid Phase Separations and Related Techniques, Beijing, China
09/12 09/13 10/14 09/15	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO, Madrid, Spain Session chair: Proteomics. 43rd International Symposium on High Performance Liquid Phase Separations and Related Techniques, Beijing, China Session organizer and chair: Human Cancer Proteome Project. 14th World Conference of HUPO,
09/12 09/13 10/14 09/15 09/15	Session leader: Cancer Proteomics and Human Proteome Project, 11th World Conference of HUPO, Boston, MA Session organizer: Human Cancer Proteome Project. 12th World Conference of HUPO, Yokohama, Japan Session organizer and chair: Human Cancer Proteome Project. 13th World Conference of HUPO, Madrid, Spain Session chair: Proteomics. 43rd International Symposium on High Performance Liquid Phase Separations and Related Techniques, Beijing, China Session organizer and chair: Human Cancer Proteome Project. 14th World Conference of HUPO, Vancouver, Canada

Session organizer and chair: Human Cancer Proteome Project. 16th World Conference of HUPO,

Session organizer and chair: Human Cancer Proteome Project. 17th World Conference of HUPO,

Session organizer and chair: Clinical Proteomics. 18th World Conference of HUPO, Adelaide,

# **RECOGNITION** (in chronological order, earliest first by start date under each subcategory)

Dublin, Ireland

Orlando, USA

Australia.

## Awards, Honors

09/17

10/18

09/19

1997	Pre-doctoral Fellowship, American Heart Association
2003	Technology Development Award, Cell Signaling Technology
2004	Young Scientist Award, Human Proteome Organization (HUPO) 3rd Annual World Congress
2012	Elected as a Board of Director, United States Human Proteomic Organization (USHUPO)
2015	Elected as a Council Member of World Human Proteomic Organization (HUPO)

### **Invited Talks**

- 07/03 Glycopeptide profiling of serum proteins and potential application in cancer diagnosis, 94th Annual Meeting of American Association for Cancer Research, Washington, DC
- 02/04 A platform for high throughput quantitative analysis of serum proteins, IBC's 2<sup>nd</sup> Annual Biomarkers: Application of Proteomics, Microarray and Metabolite Profiling Technologies, Reston, VA
- 08/04 Glyco-biomarkers for diseases, 1st Human Disease Glycomics/Proteome Initiative (HGPI) Workshop, Osaka, Japan
- 02/05 Analysis of prostate cancer by quantitative glycoproteomics using tissue specimens, Molecular Diagnostics: New Applications and Technologies Accelerating Drug Development, Princeton, NJ
- 07/05 Analysis of prostate cancer by quantitative glycoproteomics using tissue specimens, 7th International Dalian Institute of Chemical Physics (DICP) Symposium on Separation and Detection of Biomolecules, Dalian, China
- 10/05 Detection of cancer tissue-derived proteins in blood via glycopeptide analysis and mass spectrometry, 21st Asilomar

- Conference on Mass Spectrometry, Pacific Grove, CA
- 01/06 Identification of proteotypic N-linked glycopeptides for serum biomarker discovery, 6th Annual Plasma Proteome, San Diego, CA
- 08/06 Keynote Challenges and technologies in the plasma proteome analysis for marker discovery, Chinese HUPO (CNHUPO) 4th Annual Conference, Xian, China
- 04/07 Glycopeptide analysis technology, Cell Signaling Technology, Beverly, MA
- 04/07 Selective isolation of subproteomes using covalent ligation of specific functional moieties, the Association of Biomolecular Resource Facilities (ABRF) 2007, Tampa, FL
- 05/07 Glycoproteomic analysis of breast cancer tissues and identification of glycoproteins associated with advanced cancer, The Breast Cancer Conference. Baltimore, MD
- 07/07 Identification of disease-associated glycopeptides as candidate biomarkers, Hepatitis B Foundation and the Institute for Hepatitis and Virus Research, Pennsylvania Biotechnology Center, PA
- 08/07 Glycoproteomics technologies for biomarker discovery, Northwestern University, Xian, China
- 08/07 Glycopeptides in cancer detection, Chinese HUPO (CNHUPO) 5th Annual Conference, Guangzhou, China
- 11/07 Glycoproteomics, proteomics, and mass spectrometry, Case Western Reserve University, Cleveland, OH
- 03/08 Mapping expression patterns of extracellular proteomes. 4th US HUPO Annual Conference, North Bethesda, MD
- 03/08 Targeted proteomics for cancer biomarker discovery. 2008 Early Detection Research Network (EDRN) 5th Scientific Workshop. Bethesda, MD
- 04/08 N-Glycosites as molecular signatures of extracellular proteins for different cell types and diseases, Glycobiology Interest Group, Baltimore, MD
- 05/08 The selective isolation of peptides for subproteomic analysis, Department of Anesthesiology, University of Washington, Seattle, WA
- 07/08 N-linked glycosylation and cancer detection, Department of Biochemistry Seminar Program, Johns Hopkins University, Baltimore, MD
- 02/09 Glycoproteins for pathologic prediction of prostate cancer, 3<sup>rd</sup> Annual Prostate Cancer Research Day, Department of Urology, Johns Hopkins Mt. Washington Conference Center, Baltimore, MD
- 02/09 Glycoproteomics using mass spectrometry and protein arrays, 5th US HUPO Annual Conference, San Diego, CA
- 03/09 Glycomics for prostate cancer detection, 18th Early Detection Research Network (EDRN, NCI) Steering Committee Meeting, Houston, TX
- 03/09 Proteomic analysis of ovarian tumors identified proteins associated with drug resistance, Ovarian Cancer Research Meeting, Johns Hopkins University, Baltimore, MD
- 04/09 Glycoproteins in cancer diagnosis, Pathology Grand Rounds, Johns Hopkins University, Baltimore, MD
- 07/09 Changes in glycoproteins and glycans in disease development, Northwestern University, Xian, China.
- 08/09 Glycoproteins and glycosylation patterns: a new paradigm for biomarker discovery, The 6th Early Detection Research Network (EDRN, NCI) Scientific Workshop, Bethesda, MD
- 09/09 Glycoproteomics and glycomics analyses using protein microarray, HUPO 8th Annual World Congress, Toronto, Canada
- 12/09 Altered glycosylation in breast cancer: sialylation and metastasis, The Breast Cancer Conference. Johns Hopkins University, Baltimore, MD
- 03/10 Glycoproteomics and glycomics: A New Paradigm for Biomarker Discovery, Abbott, Chicago
- 04/10 Glycoproteomics and glycomics: A New Paradigm for Molecular Analysis of Disease Progression, Glycobiology Interest Group (GIG) meeting, Baltimore, MD
- 08/10 Biomarker development laboratory, EDRN planning and steering committee meeting, Rockville, MD
- 10/10 Mass spectrometric analyses of glycoproteins and glycans, Washington-Baltimore Mass Spectrometry Discussion Group, Columbia, MD

- 11/10 Glycomics using lectin microarray and mass spectrometry, Northwestern University, Xian, China
- 12/10 Mass spectrometric analysis of proteins from FFPE- or OCT-embedded tissues, Cambridge Healthtech Institute, Providence, RI
- 03/11 Proteomics and early detection, 22nd EDRN steering committee meeting, Los Angeles, CA
- 03/11 From genomics, to proteomics, to glycoproteomics, and to glycomics, UCLA Proteomics Seminar Series, Los Angeles, CA
- 03/11 Proteomics analysis of glycosylation, 7th US HUPO Annual Conference, Raleigh, NC
- 06/11 Glycoproteins as disease biomarkers, NIH&FDA Glycosciences Research Day, Bethesda, MD
- 07/11 Identification of disease biomarkers using glycoproteomics and glycomics, Institute of Biophysics, Beijing, China
- 07/11 Glycoprotein biomarkers, Beijing Proteome Research Center, Beijing, China
- 09/11 Glycoproteomics and bioinformatics analyses of biomarkers for the early detection of aggressive prostate cancer, The 23<sup>rd</sup> EDRN steering committee meeting, Bethesda, MD
- 03/12 Targeted proteomics, The 8th Conference of USHUPO, San Francisco, CA
- 03/12 Roles of platelet glycoproteins and glycans on platelet reactivity and cardiovascular disease, Programs of Excellence in Glycosciences (PEG), San Diego, CA
- 09/12 Aberrant protein expression or PTMs associated with cancer, The 11th World Conference of HUPO, Boston, MA
- 09/12 Glycosylation changes during disease progression, New England Biolabs, Ipswich, MA
- 10/12 Integrated analysis of proteins and protein modification: Changes beyond gene expression, EDRN meeting, San Antonio, TX
- 12/12 Protein post-translational modifications in tumor tissues and cells, Cell Signaling Technology, Danvers, MA
- 01/13 Glycosylation and diseases, The Texas Medical Center, Methodist Hospital, Houston, TX
- 03/13 Clinical proteomics, The 9th US HUPO Annual Conference, Baltimore, MD
- 05/13 Glycomics analysis using solid-phase glycan extraction and mass spectrometry, The Sixth Frontiers at the Chemistry and Biology Interface Symposium, University of Maryland College Park. MD
- 05/13 Structural analysis of glycoproteins and glycans, University of Mississippi Medical Center, Jackson, MS
- 05/13 Cancer-specific glycoproteins and their roles in cancer detection, US-China Prevention Research Collaboration Meeting, Bethesda, MD
- 06/13 Structure analysis of glycoproteins using mass spectrometry, Institute of Marine & Environmental Technology, Baltimore, MD
- 07/13 Proteomic biomarker discovery for translational medicine: Team approaches, University of Maryland, School of Medicine, Baltimore, MD
- 07/13 Glycomic analysis by glycoprotein immobilization for glycan extraction and liquid chromatography on microfluidic chip, Dalian Biophysical Chemical Institute, Dalian, China
- 09/13 An integrated approach for glycoproteomics research, Shandong University and Johns Hopkins University Joined Conference, Johns Hopkins University, Baltimore, MD
- 09/13 Glycoprotein biomarkers distinguishing between aggressive and indolent forms of prostate cancer, The 26th Early Detection Research Network (EDRN) Steering Committee meeting, Seattle, WA.
- 09/13 Human proteome project in cancer, The 12th International HUPO Congress, Yokohama, Japan
- 09/13 An integrated approach of proteomics, glycoproteomics and glycomics for the structural and functional study of glycoproteins, The 12<sup>th</sup> International HUPO Congress, Yokohama, Japan
- 09/13 New HPP Initiatives: Cancer HPP, the HUPO Initiative Assembly in Kyoto, Uji Obaku Plaza, Kyoto University, Japan
- 10/13 Structural and functional analyses of glycoproteins, University of Maryland, College Park, Silver Spring, MD
- 10/13 Structural and functional analyses of glycoproteins using mass spectrometry, Glycobiology Interest Group, Johns

- Hopkins University, Baltimore, MD
- 11/13 Long-term reproducibility assessment of proteomics data generated using iTRAQ labeling and LC-MS/MS platform for large-scale quantitative proteomics, the 1st annual Clinical Proteomic Tumor Analysis Consortium (CPTAC) Scientific Symposium, Bethesda, MD
- 01/14 Glycoproteomics and glycomics approaches to study function of protein glycosylation, Georgia Institute of Technology, Atlanta, GA
- 01/14 Proteomics, glycoproteomics, and glycomics using automated sample preparation and mass spectrometry, 2014 SSI Life Science Retreat and Service Summit, Washington Conference Center, Baltimore, MD
- 02/14 Automation and clinical utility of mass spectrometry-based assays, The 4th Shimadzu International Collaborative Laboratory Forum, Ho Chi Minh city, Vietnam
- 03/14 Periostin Expression in aggressive prostate cancer, The 27th Early Detection Research Network (EDRN) Steering Committee meeting, Houston, TX
- 03/14 Glycoproteins and glycans on platelet reactivity and cardiovascular disease, Programs of Excellence in Glycosciences (PEG), Inter-PEG 2014, Washington DC
- 04/14 O-glycoproteomics, National Cancer Institute's Clinical Proteomic Tumor Analysis Consortium (CPTAC) Data Jamboree, Bethesda, MD
- 04/14 Analysis of glycoproteins using multi-omics approaches, Georgetown Univ, Washington, DC
- 05/14 Integrated Analyses of transcriptome, proteome and glycoproteome reveal over expression of fucosylated proteins in aggressive prostate cancer, The International Symposium on Clinical and Translational Medicine. Shanghai, China
- 06/14 Over expression of fucosylated proteins in aggressive prostate cancer, University of Maryland School of Medicine, Department of Pharmacology, Baltimore, MD
- 07/14 Integrated glycoproteomic analysis of prostate cancer cells determines alterations in glycoprotein expression, glycosite occupancy, glycan structures, and glycosite-specific glycosylation. Chinese National Symposium on Bio-Mass Spectrometry, Shanghai, China
- 09/14 Integrated genomic, proteomic, and glycoproteomic analyses of tumor tissues reveal alterations of glycosylation in tumor subtypes, Early Detection Research Network (EDRN) Workshop, Bethesda, MD
- 10/14 HPP Initiatives: Cancer HPP, International HUPO Congress 2014, Madrid, Spain
- 11/14 Genome, proteome, and glycoproteome analyses of ovarian tumors, National Cancer Institute's Clinical Proteomic Tumor Analysis Consortium (CPTAC) Steering committee meeting, Bethesda, MD
- 11/14 Integrated genomic, proteomic, and glycoproteomic analyses of prostate cancer cells, Annual Society Meeting of Glycobiology, Honolulu, Hawaii
- 12/14 Integrated Analyses of Genome, Proteome and Glycoproteome Reveal Altered Glycosylation in Tumor Subtypes, University of Maryland, School of Pharmacy, Baltimore, MD
- 03/15 Integrated Analyses of Genome, Proteome and Glycoproteome Reveal Altered Glycosylation in Tumor Subtypes, Beijing Proteome Research Center, Beijing, China
- 03/15 Integrated Analyses of Genome, Proteome and Glycoproteome Reveal Altered Glycosylation in Tumor Subtypes, Dalian Chemical Physical Institute, Dalian, China
- 03/15 Integrated Analyses of Genome, Proteome and Glycoproteome Reveal Altered Glycosylation in Tumor Subtypes, the 11th Annual Conference of USHUPO, Tempe, AZ
- 04/15 Structural and Functional Analysis of Glycoproteins Using Proteomics, Glycoproteomics, and Glycomics, MedImmune, Gaithersburg MD
- 09/15 Glycoproteomic and proteomic analyses reveal glycoprotein alteration in protein abundance, glycosite occupancy, glycans and glycosite-specific glycosylation, The 43rd International Symposium on High Performance Liquid Phase Separations and Related Techniques, Beijing, China
- 09/15 Comprehensive analysis of glycoproteins, The 14th World Conference of HUPO, Vancouver, Canada
- 12/15 Proteomic and Glycoproteomic Analyses Reveal Altered Protein Glycosylation, Targeted Proteomics Workshop and International Symposium, Mumbai, India

- 03/16 Data Analysis and Future Analytical Trends of Glycoproteins, Frontiers in Glycan Analysis, University of Georgia, Complex Carbohydrate Research Center, Athens, GA
- 04/16 Structural and Functional Analyses of Glycoproteins Using Mass Spectrometry, The 1st Phoenix Mini-Symposium on Frontiers of Proteomics, the National Center for Protein Sciences-Beijing (Phoenix Center) and Beijing Proteome Research Center, Beijing, China
- 06/16 Characterization of Glycoproteins for Therapeutic Development. Bristol-Myers Squibb Company, Princeton, NJ
- 06/16 Glycoprotein Analysis Using Mass Spectrometry. 2016 NIH & FDA Glycoscience Research Day. NIH Natcher Conference Center, Bethesda, MD
- 07/16 Automation of glycoprotein analysis. New England Biolabs, Ipswich, MA
- 08/16 Selection of Tandem Mass Spectra for Identification of Intact N- and O-linked Glycopeptides. The 4th workshop on Computational Proteomics, Dalian, China
- 09/16 Structural and Functional Analyses of Protein N-linked Glycosylation. University of Maryland, College Park, MD
- 09/16 Keynote Speaker, Glycoproteomics Technical limitations & Prospects. The 15th Human Proteome Organization World Congress (HUPO), Taipei, Taiwan
- 11/16 Proteomics and Glycoproteomics Research: What We Now Know. Fudan University, Shanghai, China
- 11/16 Proteomics and Glycoproteomics: What We Learn Beyond Genomics, Dalian Chemical Physical Institute, Dalian, China
- 12/16 Proteomics and Glycoproteomics: What We Learn Beyond Genomics. Pathology Grand Rounds, Johns Hopkins University, Baltimore, MD
- 01/17 Comprehensive Analysis of Protein Glycosylation by Mass Spectrometry. Glycobiology Interest Group Meeting, Baltimore, MD
- 02/17 Keynote Speaker, Genomic, Proteomic and Glycoproteomic Analyses of Tumors. The Swiss Proteomics Society and the Life Science Switzerland 2017, Zurich, Switzerland
- 02/17 Structural and Functional Analysis of Glycoproteins Using Mass Spectrometry. Department of Biochemistry and Molecular & Cellular Biology, Georgetown University, Washington DC
- 03/17 Structural and Functional Analysis of Glycoproteins Using Mass Spectrometry. Proteomic Interest Group Lecture, National Institutes of Health, Bethesda, MD
- 03/17 Methods for Quantitative Analysis of Glycoproteins. PITTCON2017, Chicago, IL
- 03/17 Proteogenomic analyses of cancer genes and proteins. USHUPO2017, San Diego, CA
- 04/17 Deciphering cell surface proteins associated with defective genomic alterations using glycoproteogenomics for targeted therapy. Sino-American Pharmaceutical Professionals Association DC chapter (SAPA-DC). Rockville, MD
- 04/17 Structural and Functional Analysis of Glycoproteins. University of Buffalo, Buffalo, NY
- 04/17 DICP Symposium on Bioanalytical methods, techniques and applications: Highly efficient Separation and Characterization of Biological Samples. Dalian, China
- 05/17 Tumor Associated Glycoproteins. NIH & FDA Glycoscience Day, National Institutes of Health, Bethesda, MD
- 08/17 Functional Analysis of Protein Glycosylation Genes. Posttranslational Modification Networks, Gordon Research Conference, The Hong Kong University of Science and Technology, Hong Kong, China
- 08/17 Functional analysis of Protein Glycosylation by Glycoproteomics. Department of Pathology and Laboratory Medicine, University of Pennsylvania, Philadelphia, PA
- 09/17 Fucosylated glycoproteins associated with advanced prostate cancer. The 16th Human Proteome Organization World Congress (HUPO), Dublin, Ireland
- 03/18 Glycoproteomics and glycomics for clinical biomarker discovery. Session Chair. USHUPO2018, Minneapolis, MN.
- 03/18 Functional analysis of protein glycosylation in cancer. Dalian Medical University, Dalian, China
- 05/18 Enrichment of Phospho- and Glyco-peptides for Simultaneous Proteomic Analyses of Multiple Protein Modifications. Inova Schar Cancer Institute, Inova Center for Personalized Health, John P. Murtha Cancer Center,

- Walter Reed National Military Medical Center, Uniformed Services University of the Health Sciences, Annandale, VA
- 06/18 Plenary Lecture, Deciphering Novel Roles of Protein Glycosylation Using Mass Spectrometry. 2<sup>nd</sup> Mass Spectrometry Symposium at the Medical College of Wisconsin, Milwaukee, WI
- 06/18 Keynote speaker, Multi-omic analysis of tumor tissues reveals a large number of glycopeptides and their association with glycosylation enzymes. GlycoT2018, Qingtao, China
- 07/18 Keynote Lecture, Co-analysis of Glycoproteomics, Phosphoproteomics, and Global Proteomics from the Same Sample. HPLC2018, 47th International Symposium on High Performance Liquid Phase Separations and Related Techniques, Washington DC
- 08/18 Multi-omic analysis of tumor tissues reveals a large number of glycopeptides and their association with glycosylation enzymes. 1st Annual North American Mass Spec Summer School, Madison, WI.
- 09/18 Multi-omic analysis of tumor tissues reveals a large number of glycopeptides and their association with glycosylation enzymes. New Jersey Mass Spectrometry Discussion Group. Bridgewater, NJ.
- 03/19 Fucosylated and urinary glycoprotein biomarkers for aggressive prostate cancer. EDRN meeting. Nashville, TN.
- 06/19 Integrated proteogenomic characterization of clear cel renal cell carcinoma. The 13th National Cancer Center International Symposium "Cancer Proteogenomics: The Force Awakens". Soul, Korea.
- 01/20 Discovery of urinary glycoproteins for aggressive prostate cancer. The 7th US-Japan Workshop on Biomarkers for Cancer early Detection. Tokyo, Japan.
- 02/20 Proteogenomic characterization of cancer. Departmental Seminar. Department of Biochemistry and Molecular Biophysics, University of Pennsylvania. Philadelphia, PA.
- 03/20 Comprehensive analysis of glycoproteins. PITTCON Conference 2020. McCormick Place. Chicago, IL.
- 08/21 Proteogenomic characterization of pancreatic ductal adenocarcinoma. Chinese American Society for Mass Spectrometry. Talk given virtually.
- 09/21 Data-independent acquisition approach for proteomic and PTM analyses at single-cell level. Think tank meeting of National Institute on Aging. Talk given virtually.
- 09/21 Proteogenomic characterization of pancreatic ductal adenocarcinoma. American Association for Cancer Research: Pancreatic Cancer. Talk given virtually.
- 11/21 Proteogenomic characterization of cancer. Department of Biochemistry and Molecular Biology. Tulane University. Talk given virtually.