## Qi Wu

## List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

54	1,161	21	<b>32</b>
papers	citations	h-index	g-index
57	1,330 ext. citations	5.9	4.22
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
54	New GO-PEI-Au-L-Cys ZIC-HILIC composites: synthesis and selective enrichment of glycopeptides. <i>Nanoscale</i> , <b>2014</b> , 6, 5616-9	7.7	85
53	Recent advances on multidimensional liquid chromatography-mass spectrometry for proteomics: from qualitative to quantitative analysisa review. <i>Analytica Chimica Acta</i> , <b>2012</b> , 731, 1-10	6.6	79
52	Hydrophilic GO/Fe3O4/Au/PEG nanocomposites for highly selective enrichment of glycopeptides. <i>Nanoscale</i> , <b>2016</b> , 8, 4894-7	7.7	68
51	Boronate affinity monolith with a gold nanoparticle-modified hydrophilic polymer as a matrix for the highly specific capture of glycoproteins. <i>Chemistry - A European Journal</i> , <b>2014</b> , 20, 8737-43	4.8	59
50	Integrated sample pretreatment system for N-linked glycosylation site profiling with combination of hydrophilic interaction chromatography and PNGase F immobilized enzymatic reactor via a strong cation exchange precolumn. <i>Analytical Chemistry</i> , <b>2011</b> , 83, 7457-63	7.8	55
49	Epitope imprinted polyethersulfone beads by self-assembly for target protein capture from the plasma proteome. <i>Chemical Communications</i> , <b>2014</b> , 50, 9521-4	5.8	47
48	Boronic Acid-Functionalized Particles with Flexible Three-Dimensional Polymer Branch for Highly Specific Recognition of Glycoproteins. <i>ACS Applied Materials &amp; Distriction of Communication of Communication and Polymer Branch for Highly Specific Recognition of Glycoproteins. ACS Applied Materials &amp; District Recognition of Communication (Communication) and Polymer Branch for Highly Specific Recognition of Glycoproteins. ACS Applied Materials &amp; District Recognition (Communication) and Polymer Branch for Highly Specific Recognition of Glycoproteins. ACS Applied Materials &amp; District Recognition (Communication) and Polymer Branch for Highly Specific Recognition (Communication) and Polymer Branch for Highly (Communication) and Polymer Branch</i>	9.5	47
47	An efficient approach to prepare boronate core-shell polymer nanoparticles for glycoprotein recognition via combined distillation precipitation polymerization and RAFT media precipitation polymerization. <i>Chemical Communications</i> , <b>2015</b> , 51, 3896-8	5.8	44
46	1-Dodecyl-3-methylimidazolium chloride-assisted sample preparation method for efficient integral membrane proteome analysis. <i>Analytical Chemistry</i> , <b>2014</b> , 86, 7544-50	7.8	42
45	Gold nanoparticles immobilized hydrophilic monoliths with variable functional modification for highly selective enrichment and on-line deglycosylation of glycopeptides. <i>Analytica Chimica Acta</i> , <b>2015</b> , 900, 83-9	6.6	40
44	Multi-mode application of graphene quantum dots bonded silica stationary phase for high performance liquid chromatography. <i>Journal of Chromatography A</i> , <b>2017</b> , 1492, 61-69	4.5	38
43	Mass defect-based pseudo-isobaric dimethyl labeling for proteome quantification. <i>Analytical Chemistry</i> , <b>2013</b> , 85, 10658-63	7.8	38
42	Synthesis of Zwitterionic Polymer Particles via Combined Distillation Precipitation Polymerization and Click Chemistry for Highly Efficient Enrichment of Glycopeptide. <i>ACS Applied Materials &amp; Interfaces</i> , <b>2016</b> , 8, 22018-24	9.5	37
41	CHIP Regulates Aquaporin-2 Quality Control and Body Water Homeostasis. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2018</b> , 29, 936-948	12.7	36
40	Preparation of hydrophilic monolithic capillary column by in situ photo-polymerization of N-vinyl-2-pyrrolidinone and acrylamide for highly selective and sensitive enrichment of N-linked glycopeptides. <i>Talanta</i> , <b>2016</b> , 146, 225-30	6.2	35
39	Nanocellulose 3, 5-Dimethylphenylcarbamate Derivative Coated Chiral Stationary Phase: Preparation and Enantioseparation Performance. <i>Chirality</i> , <b>2016</b> , 28, 376-81	2.1	26
38	Rapid Aldosterone-Mediated Signaling in the DCT Increases Activity of the Thiazide-Sensitive NaCl Cotransporter. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2019</b> , 30, 1454-1470	12.7	25

Characterization of AQPs in Mouse, Rat, and Human Colon and Their Selective Regulation by Bile Acids. <i>Frontiers in Nutrition</i> , <b>2016</b> , 3, 46	6.2	25	
An integrated sample pretreatment platform for quantitative N-glycoproteome analysis with combination of on-line glycopeptide enrichment, deglycosylation and dimethyl labeling. <i>Analytica Chimica Acta</i> , <b>2014</b> , 833, 1-8	6.6	23	
Graphene quantum dots functionalized Eyclodextrin and cellulose chiral stationary phases with enhanced enantioseparation performance. <i>Journal of Chromatography A</i> , <b>2019</b> , 1600, 209-218	4.5	22	
Monodisperse boronate polymeric particles synthesized by a precipitation polymerization strategy: particle formation and glycoprotein response from the standpoint of the Flory-Huggins model. <i>ACS Applied Materials &amp; Discourse (Materials &amp; Discourse)</i> 100 Materials 2014, 6, 2059-66	9.5	22	
A hydrophilic immobilized trypsin reactor with N-vinyl-2-pyrrolidinone modified polymer microparticles as matrix for highly efficient protein digestion with low peptide residue. <i>Journal of Chromatography A</i> , <b>2012</b> , 1246, 111-6	4.5	21	
Graphene quantum dots-functionalized C hydrophobic/hydrophilic stationary phase for high performance liquid chromatography. <i>Talanta</i> , <b>2019</b> , 194, 105-113	6.2	18	
Biphasic microreactor for efficient membrane protein pretreatment with a combination of formic acid assisted solubilization, on-column pH adjustment, reduction, alkylation, and tryptic digestion. <i>Analytical Chemistry</i> , <b>2013</b> , 85, 8507-12	7.8	17	
A Systems Level Analysis of Vasopressin-mediated Signaling Networks in Kidney Distal Convoluted Tubule Cells. <i>Scientific Reports</i> , <b>2015</b> , 5, 12829	4.9	17	
Ionic liquid-functionalized graphene quantum dot-bonded silica as multi-mode HPLC stationary phase with enhanced selectivity for acid compounds. <i>New Journal of Chemistry</i> , <b>2018</b> , 42, 8672-8680	3.6	16	
Activation of the metabolic sensor AMP-activated protein kinase inhibits aquaporin-2 function in kidney principal cells. <i>American Journal of Physiology - Renal Physiology</i> , <b>2016</b> , 311, F890-F900	4.3	16	
NSI and NSMT: usages of MS/MS fragment ion intensity for sensitive differential proteome detection and accurate protein fold change calculation in relative label-free proteome quantification. <i>Analyst, The</i> , <b>2012</b> , 137, 3146-53	5	14	
The murine choroid plexus epithelium expresses the 2Cl/H exchanger ClC-7 and Na/H exchanger NHE6 in the luminal membrane domain. <i>American Journal of Physiology - Cell Physiology</i> , <b>2018</b> , 314, C4	39 <sup>5</sup> <del>C</del> 44	8 <sup>12</sup>	
imFASP: An integrated approach combining in-situ filter-aided sample pretreatment with microwave-assisted protein digestion for fast and efficient proteome sample preparation. <i>Analytica Chimica Acta</i> , <b>2016</b> , 912, 58-64	6.6	11	
Prefractionation and separation by C8 stationary phase: effective strategies for integral membrane proteins analysis. <i>Talanta</i> , <b>2012</b> , 88, 567-72	6.2	11	
The thiazide sensitive sodium chloride co-transporter NCC is modulated by site-specific ubiquitylation. <i>Scientific Reports</i> , <b>2017</b> , 7, 12981	4.9	10	
A paired ions scoring algorithm based on Morpheus for simultaneous identification and quantification of proteome samples prepared by isobaric peptide termini labeling strategies. <i>Proteomics</i> , <b>2015</b> , 15, 1781-8	4.8	10	
Pseudo isobaric peptide termini labelling for relative proteome quantification by SWATH MS acquisition. <i>Analyst, The</i> , <b>2016</b> , 141, 4912-8	5	9	
Preparation and chromatographic performance of a multifunctional immobilized chiral stationary phase based on dialdehyde microcrystalline cellulose derivatives. <i>Chirality</i> , <b>2019</b> , 31, 669-681	2.1	9	
	Acids. Frontiers in Nutrition, 2016, 3, 46  An integrated sample pretreatment platform for quantitative N-glycoproteome analysis with combination of on-line glycopetide enrichment, deglycosylation and dimethyl labeling. Analytica Chimica Acta, 2014, 833, 1-8  Graphene quantum dots functionalized Bryclodextrin and cellulose chiral stationary phases with enhanced enantioseparation performance. Journal of Chromatography A, 2019, 1600, 209-218  Monodisperse boronate polymeric particles synthesized by a precipitation polymerization strategy: particle formation and glycoprotein response from the standpoint of the Flory-Huggins model. ACS Applied Materials & Ampl. Interfaces, 2014, 6, 2059-66  A hydrophillic immobilized trypsin reactor with N-vinyl-2-pyrrolidinone modified polymer microparticles as matrix for highly efficient protein digestion with low peptide residue. Journal of Chromatography A, 2012, 1246, 111-6  Graphene quantum dots-functionalized C hydrophobic/hydrophilic stationary phase for high performance liquid chromatography. Talanta, 2019, 194, 105-113  Biphasic microreactor for efficient membrane protein pretreatment with a combination of formic acid assisted solubilization, on-column pH adjustment, reduction, alkylation, and tryptic digestion. Analytical Chemistry, 2013, 85, 8507-12  A Systems Level Analysis of Vasopressin-mediated Signaling Networks in Kidney Distal Convoluted Tubule Cells. Scientific Reports, 2015, 5, 12829  Ionic liquid-Functionalized graphene quantum dot-bonded silica as multi-mode HPLC stationary phase with enhanced selectivity for acid compounds. New Journal of Chemistry, 2018, 42, 8672-8680  Activation of the metabolic sensor AMP-activated protein kinase inhibits aquaporin-2 function in kidney principal cells. American Journal of Physiology, Physiology, 2018, 311, F890-F900  NSI and NSMT: usages of MS/MS fragment ion intensity for sensitive differential proteome detection and accurate protein fold change calculation in relative label-free proteome quantification. Analyst, The, 20	Acids. Frontiers in Nutrition, 2016, 3, 46  An integrated sample pretreatment platform for quantitative N-glycoproteome analysis with combination of on-line glycopeptide enrichment, deglycosylation and dimethyl labeling. Analytica Chimica Acta, 2014, 833, 1-8  Graphene quantum dots functionalized Byclodextrin and cellulose chiral stationary phases with enhanced enantioseparation performance. Journal of Chromatography A, 2019, 1600, 209-218  4-5  Monodisperse boronate polymeric particles synthesized by a precipitation polymerization strategy: particle formation and glycoprotein response from the standpoint of the Flory-Huggins model. ACS Applied Materials & Amplied Immobilized trypsin reactor with N-vinyl-2-pyrrollidinone modified polymer microparticles as matrix for highly efficient protein digestion with low peptide residue. Journal of Chromatography A, 2012, 1246, 111-6  Graphene quantum dost-functionalized C hydrophobic/hydrophilic stationary phase for high performance liquid chromatography. Talanta, 2019, 194, 105-113  Biphasic microreactor for efficient membrane protein pretreatment with a combination of formic acid assisted solubilization, on-column pH adjustment, reduction, alkylation, and tryptic digestion. Analytical Chemistry, 2013, 85, 8507-12  A Systems Level Analysis of Vasopressin-mediated Signaling Networks in Kidney Distal Convoluted Tubule Cells. Scientific Reports, 2015, 5, 12829  Ionic liquid-functionalized graphene quantum dot-bonded silica as multi-mode HPLC stationary phase with enhanced selectivity for acid compounds. New Journal of Chemistry, 2018, 42, 8672-8680  Activation of the metabolic sensor AMP-activated protein kinase inhibits aquaporin-2 function in kindeny principal cells. American Journal of Physiology. Renal Physiology, 2016, 311, R890-F900  NSI and NSMT: usages of MS/MS fragment ion intensity for sensitive differential proteome detection and accurate protein fold change calculation in relative label-free proteome  15 quantification. Analyst, The, 2012, 137, 3146-53  The muri	Acids. Frontiers in Nutrition, 2016, 3, 46  An integrated sample pretreatment platform for quantitative N-glycoproteome analysis with combination of on-line glycopeptide enrichment, deglycosylation and dimethyl labeling. Analytica Chimica Acta, 2014, 833, 1-8  Graphene quantum dots functionalized ftyclodextrin and cellulose chiral stationary phases with enhanced enantioseparation performance. Journal of Chromatography A, 2019, 1600, 209-218  Monodisperse boronate polymeric particles synthesized by a precipitation polymerization strategy: particle formation and glycoprotein response from the standpoint of the Flory-Huggins model. ACS Applied Materials &, interfaces, 2014, 6, 2059-66  A hydrophilic Immobilized trypsin reactor with N-vinyl-2-pyrrolidinone modified polymer microparticles as matrix for highly efficient protein digestion with low peptide residue. Journal of Chromatography A, 2012, 1246, 111-6  Graphene quantum dots-functionalized C hydrophobic/hydrophilic stationary phase for high performance liquid chromatography. Talanta, 2019, 194, 105-113  Bliphasic microreactor for efficient membrane protein pretreatment with a combination of Formic acid assisted solubilization, on-column pH adjustment, reduction, alkylation, and tryptic digestion. Analytical Chemistry, 2013, 85, 8507-12  A Systems Level Analysis of Vasopressin-mediated Signaling Networks in Kidney Distal Convoluted Tubule Cells. Scientific Reports, 2015, 5, 12829  lonic liquid-functionalized graphene quantum dot-bonded silica as multi-mode HPLC stationary phase with enhanced selectivity for acid compounds. New Journal of Chemistry, 2018, 42, 8672-8680  ACtivation of the metabolic sensor AMP-activated protein kinase inhibits aquaporin-2 function in kidney principal cells. American Journal of Physiology, 2016, 311, F890-F900  NSI and NSMT: usages of MS/MS fragment ion intensity for sensitive differential proteome quantification and accurate protein fold change calculation in relative label-free proteome quantification and separation by C8 stat

19	Partially isobaric peptide termini labeling assisted proteome quantitation based on MS and MS/MS signals. <i>Journal of Proteomics</i> , <b>2015</b> , 114, 152-60	3.9	8
18	Nanocellulose Derivative/Silica Hybrid Core-Shell Chiral Stationary Phase: Preparation and Enantioseparation Performance. <i>Molecules</i> , <b>2016</b> , 21,	4.8	8
17	Glycoprotein recognition by water-compatible core-shell polymeric submicron particles. <i>Journal of Materials Chemistry B</i> , <b>2015</b> , 3, 3927-3930	7.3	7
16	Nano-amylose-2,3-bis(3,5-dimethylphenylcarbamate)-silica hybrid sol immobilized on open tubular capillary column for capillary electrochromatography enantioseparation. <i>Electrophoresis</i> , <b>2018</b> , 39, 1086	5 <sup>3</sup> 1695	6
15	A robust and effective intact protein fractionation strategy by GO/PEI/Au/PEG nanocomposites for human plasma proteome analysis. <i>Talanta</i> , <b>2018</b> , 178, 49-56	6.2	6
14	Dysregulation of Principal Cell miRNAs Facilitates Epigenetic Regulation of AQP2 and Results in Nephrogenic Diabetes Insipidus. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2021</b> , 32, 1339-13	5 <sup>12.7</sup>	6
13	Fast MS/MS acquisition without dynamic exclusion enables precise and accurate quantification of proteome by MS/MS fragment intensity. <i>Scientific Reports</i> , <b>2016</b> , 6, 26392	4.9	5
12	Decrease of dynamic range of proteins in human plasma by ampholine immobilized polymer microspheres. <i>Analytica Chimica Acta</i> , <b>2014</b> , 826, 43-50	6.6	5
11	Genetic disruption of slc4a10 alters the capacity for cellular metabolism and vectorial ion transport in the choroid plexus epithelium. <i>Fluids and Barriers of the CNS</i> , <b>2020</b> , 17, 2	7	4
10	SDS-PAGE-free protocol for comprehensive identification of cytochrome P450 enzymes and uridine diphosphoglucuronosyl transferases in human liver microsomes. <i>Proteomics</i> , <b>2012</b> , 12, 3464-9	4.8	4
9	The Cl/HCO exchanger pendrin is downregulated during oral co-administration of exogenous mineralocorticoid and KCl in patients with primary aldosteronism. <i>Journal of Human Hypertension</i> , <b>2021</b> , 35, 837-848	2.6	4
8	Improved accuracy for label-free absolute quantification of proteome by combining the Absolute Protein EXpression profiling algorithm and summed tandem mass spectrometric total ion current. <i>Analyst, The</i> , <b>2014</b> , 139, 138-46	5	3
7	Large-Scale Proteomic Assessment of Urinary Extracellular Vesicles Highlights Their Reliability in Reflecting Protein Changes in the Kidney. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2021</b> , 32, 2195-2209	12.7	3
6	SUMOylation Landscape of Renal Cortical Collecting Duct Cells. <i>Journal of Proteome Research</i> , <b>2019</b> , 18, 3640-3648	5.6	2
5	An in vivo protein landscape of the mouse DCT during high dietary K or low dietary Na intake. <i>American Journal of Physiology - Renal Physiology</i> , <b>2021</b> , 320, F908-F921	4.3	2
4	Label-free quantification of differentially expressed proteins in mouse liver cancer cells with high and low metastasis rates by a SWATH acquisition method. <i>Science China Chemistry</i> , <b>2014</b> , 57, 718-722	7.9	1
3	Urinary proteomics for kidney dysfunction: insights and trends. <i>Expert Review of Proteomics</i> , <b>2021</b> , 18, 437-452	4.2	1
2	The Hydrogen-Coupled Oligopeptide Membrane Cotransporter Pept2 is SUMOylated in Kidney Distal Convoluted Tubule Cells. <i>Frontiers in Molecular Biosciences</i> , <b>2021</b> , 8, 790606	5.6	O

Improved Accuracy of Proteome Quantification by MS/MS Fragment Intensity. *FASEB Journal*, **2015**, 29, 567.9

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