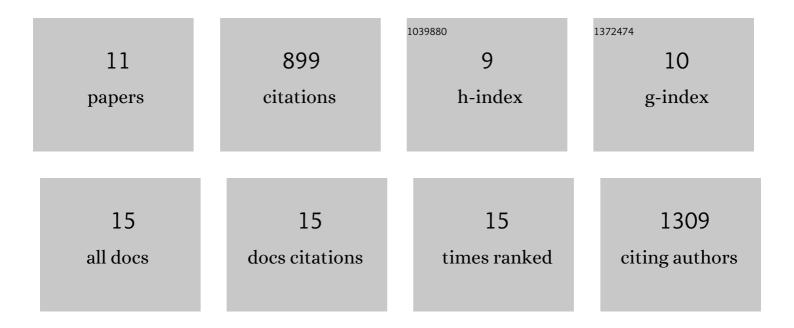
## Jiaming Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6733619/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	TMTpro reagents: a set of isobaric labeling mass tags enables simultaneous proteome-wide measurements across 16 samples. Nature Methods, 2020, 17, 399-404.	9.0	276
2	A Quantitative Tissue-Specific Landscape of Protein Redox Regulation during Aging. Cell, 2020, 180, 968-983.e24.	13.5	220
3	TMTpro-18plex: The Expanded and Complete Set of TMTpro Reagents for Sample Multiplexing. Journal of Proteome Research, 2021, 20, 2964-2972.	1.8	158
4	Proteome-wide mapping of short-lived proteins in human cells. Molecular Cell, 2021, 81, 4722-4735.e5.	4.5	64
5	Improved Monoisotopic Mass Estimation for Deeper Proteome Coverage. Journal of Proteome Research, 2021, 20, 591-598.	1.8	52
6	Investigation of Proteomic and Phosphoproteomic Responses to Signaling Network Perturbations Reveals Functional Pathway Organizations in Yeast. Cell Reports, 2019, 29, 2092-2104.e4.	2.9	41
7	Selection of Heating Temperatures Improves the Sensitivity of the Proteome Integral Solubility Alteration Assay. Journal of Proteome Research, 2020, 19, 2159-2166.	1.8	36
8	Assessing target engagement using proteome-wide solvent shift assays. ELife, 2021, 10, .	2.8	22
9	Quantitative Phosphoproteomics Revealed Glucose-Stimulated Responses of Islet Associated with Insulin Secretion. Journal of Proteome Research, 2015, 14, 4635-4646.	1.8	19
10	Systematic Synergy of Glucose and GLP-1 to Stimulate Insulin Secretion Revealed by Quantitative Phosphoproteomics. Scientific Reports, 2017, 7, 1018.	1.6	6
11	Profiling Yeast Deletion Strains Using Sample Multiplexing and Network-Based Analyses. Journal of	1.8	1