

# Liang Jin

## List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Ultra-High-Resolution IonStar Strategy Enhancing Accuracy and Precision of MS1-Based Proteomics and an Extensive Comparison with State-of-the-Art SWATH-MS in Large-Cohort Quantification. <i>Analytical Chemistry</i> , 2021, 93, 4884-4893.	6.5	14
2	Integrative Analysis of Transcriptomic and Proteomic Profiling in Inflammatory Bowel Disease Colon Biopsies. <i>Inflammatory Bowel Diseases</i> , 2019, 25, 1906-1918.	1.9	22
3	Surfactant Cocktail-Aided Extraction/Precipitation/On-Pellet Digestion Strategy Enables Efficient and Reproducible Sample Preparation for Large-Scale Quantitative Proteomics. <i>Analytical Chemistry</i> , 2018, 90, 10350-10359.	6.5	37
4	Predicted RNA Binding Proteins Pes4 and Mip6 Regulate mRNA Levels, Translation, and Localization during Sporulation in Budding Yeast. <i>Molecular and Cellular Biology</i> , 2017, 37, .	2.3	17
5	Post-transcriptional regulation in budding yeast meiosis. <i>Current Genetics</i> , 2016, 62, 313-315.	1.7	19
6	Sequestration of mRNAs Modulates the Timing of Translation during Meiosis in Budding Yeast. <i>Molecular and Cellular Biology</i> , 2015, 35, 3448-3458.	2.3	28
7	Codon Usage Bias and Determining Forces in Green Plant Mitochondrial Genomes. <i>Journal of Integrative Plant Biology</i> , 2011, 53, 324-334.	8.5	19
8	Complex Mutation and Weak Selection together Determined the Codon Usage Bias in Bryophyte Mitochondrial Genomes. <i>Journal of Integrative Plant Biology</i> , 2010, 52, 1100-1108.	8.5	12