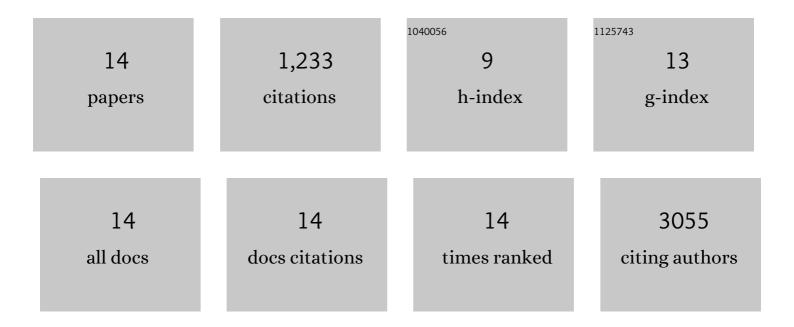
Chaochao Wu

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Preconditioning in the Rhesus Macaque Induces a Proteomic Signature Following Cerebral Ischemia that Is Associated with Neuroprotection. Translational Stroke Research, 2019, 10, 440-448. | 4.2 | 11 |
| 2 | Targeted proteomic assays for quantitation of proteins identified by proteogenomic analysis of ovarian cancer. Scientific Data, 2017, 4, 170091. | 5.3 | 29 |
| 3 | Dysfunction of PLA2G6 and CYP2C44-associated network signals imminent carcinogenesis from chronic inflammation to hepatocellular carcinoma. Journal of Molecular Cell Biology, 2017, 9, 489-503. | 3.3 | 64 |
| 4 | Quantification of mutant SPOP proteins in prostate cancer using mass spectrometry-based targeted proteomics. Journal of Translational Medicine, 2017, 15, 175. | 4.4 | 5 |
| 5 | Integrated Proteogenomic Characterization of Human High-Grade Serous Ovarian Cancer. Cell, 2016, 166, 755-765. | 28.9 | 804 |
| 6 | Recommendations for the Generation, Quantification, Storage, and Handling of Peptides Used for Mass Spectrometry–Based Assays. Clinical Chemistry, 2016, 62, 48-69. | 3.2 | 187 |
| 7 | An Optimized Informatics Pipeline for Mass Spectrometry-Based Peptidomics. Journal of the American Society for Mass Spectrometry, 2015, 26, 2002-2008. | 2.8 | 10 |
| 8 | Sensitive Targeted Quantification of ERK Phosphorylation Dynamics and Stoichiometry in Human Cells without Affinity Enrichment. Analytical Chemistry, 2015, 87, 1103-1110. | 6.5 | 32 |
| 9 | Mass Spectrometry for Biomarker Development. Biomarkers in Disease, 2015, , 17-48. | 0.1 | 1 |
| 10 | Comprehensive Quantitative Analysis of Ovarian and Breast Cancer Tumor Peptidomes. Journal of Proteome Research, 2015, 14, 422-433. | 3.7 | 26 |
| 11 | Expediting SRM Assay Development for Large-Scale Targeted Proteomics Experiments. Journal of Proteome Research, 2014, 13, 4479-4487. | 3.7 | 29 |
| 12 | Mass Spectrometry for Biomarker Development. , 2014, , 1-25. | | 0 |
| 13 | Delicate Analysis of Post-Translational Modifications on Dishevelled 3. Journal of Proteome Research, 2012, 11, 3829-3837. | 3.7 | 12 |
| 14 | Proteomic mining in the dysplastic liver of WHV/câ€ <i>myc</i> mice – insights and indicators for early hepatocarcinogenesis. FEBS Journal, 2010, 277, 4039-4053. | 4.7 | 23 |