

# Weina Gao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3183316/publications.pdf>

Version: 2024-02-01

12  
papers

419  
citations

1307594

7  
h-index

1125743

13  
g-index

13  
all docs

13  
docs citations

13  
times ranked

811  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Targeting LIF-mediated paracrine interaction for pancreatic cancer therapy and monitoring. <i>Nature</i> , 2019, 569, 131-135.  | 27.8 | 287       |
| 2  | Spatial-Resolution Cell Type Proteome Profiling of Cancer Tissue by Fully Integrated Proteomics Technology. <i>Analytical Chemistry</i> , 2018, 90, 5879-5886.  | 6.5  | 35        |
| 3  | Spatial proteome profiling by immunohistochemistry-based laser capture microdissection and data-independent acquisition proteomics. <i>Analytica Chimica Acta</i> , 2020, 1127, 140-148.  | 5.4  | 23        |
| 4  | A Fully Integrated Spintip-Based Approach for Sensitive and Quantitative Profiling of Region-Resolved in Vivo Brain Glycoproteome. <i>Analytical Chemistry</i> , 2019, 91, 9181-9189.   | 6.5  | 14        |
| 5  | AutoProteome Chip System for Fully Automated and Integrated Proteomics Sample Preparation and Peptide Fractionation. <i>Analytical Chemistry</i> , 2020, 92, 8893-8900.   | 6.5  | 13        |
| 6  | An integrated strategy for high-sensitive and multi-level glycoproteome analysis from low micrograms of protein samples. <i>Journal of Chromatography A</i> , 2019, 1600, 46-54.  | 3.7  | 11        |
| 7  | Synergistic optimization of Liquid Chromatography and Mass Spectrometry parameters on Orbitrap Tribrid mass spectrometer for high efficient data-independent proteomics. <i>Journal of Mass Spectrometry</i> , 2021, 56, e4653. | 1.6  | 11        |
| 8  | Motif-dependent immune co-receptor interactome profiling by photoaffinity chemical proteomics. <i>Cell Chemical Biology</i> , 2022, 29, 1024-1036.e5.   | 5.2  | 8         |
| 9  | Fritted tip capillary column with negligible dead volume facilitated ultrasensitive and deep proteomics. <i>Analytica Chimica Acta</i> , 2022, 1201, 339615.  | 5.4  | 5         |
| 10 | Stable and EGF-Induced Temporal Interactome Profiling of CBL and CBLB Highlights Their Signaling Complex Diversity. <i>Journal of Proteome Research</i> , 2021, 20, 3709-3719.  | 3.7  | 4         |
| 11 | A fully integrated sample preparation strategy for highly sensitive intact glycoproteomics. <i>Analyst</i> , The, 2022, 147, 794-798.   | 3.5  | 4         |
| 12 | Multiomic analysis of a dried single-drop plasma sample using an integrated mass spectrometry approach. <i>Analyst</i> , The, 2020, 145, 6441-6446.   | 3.5  | 3         |