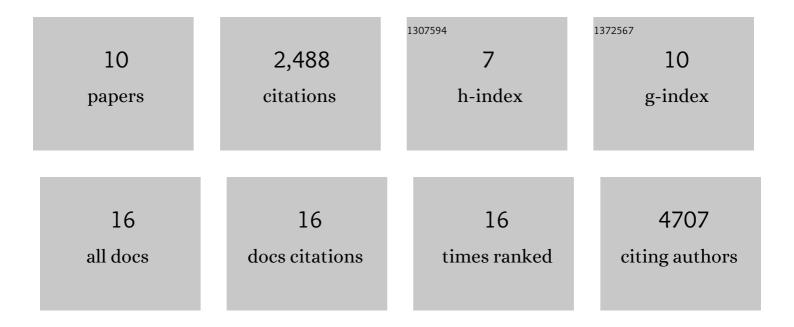
Shubham Gupta

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8856718/publications.pdf Version: 2024-02-01



SHUBHAM CUDTA

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Democratizing data-independent acquisition proteomics analysis on public cloud infrastructures via the Galaxy framework. GigaScience, 2022, 11, . | 6.4 | 13 |
| 2 | Automated Workflow for Peptide-Level Quantitation from DIA/ Data. Methods in Molecular Biology, 2021, 2228, 453-468. | 0.9 | 4 |
| 3 | DIAproteomics: A Multifunctional Data Analysis Pipeline for Data-Independent Acquisition Proteomics and Peptidomics. Journal of Proteome Research, 2021, 20, 3758-3766. | 3.7 | 17 |
| 4 | Analyzing Assay Specificity in Metabolomics Using Unique Ion Signature Simulations. Analytical Chemistry, 2021, 93, 11415-11423. | 6.5 | 1 |
| 5 | SmartPeak Automates Targeted and Quantitative Metabolomics Data Processing. Analytical Chemistry, 2020, 92, 15968-15974. | 6.5 | 23 |
| 6 | DrawAlignR: An Interactive Tool for Across Run Chromatogram Alignment Visualization. Proteomics, 2020, 20, e1900353. | 2.2 | 6 |
| 7 | Long Noncoding RNA and Predictive Model To Improve Diagnosis of Clinically Diagnosed Pulmonary Tuberculosis. Journal of Clinical Microbiology, 2020, 58, . | 3.9 | 18 |
| 8 | DIAlignR Provides Precise Retention Time Alignment Across Distant Runs in DIA and Targeted Proteomics. Molecular and Cellular Proteomics, 2019, 18, 806-817. | 3.8 | 23 |
| 9 | A Computational Tool to Detect and Avoid Redundancy in Selected Reaction Monitoring. Molecular and Cellular Proteomics, 2012, 11, 540-549. | 3.8 | 90 |
| 10 | Targeted Data Extraction of the MS/MS Spectra Generated by Data-independent Acquisition: A New Concept for Consistent and Accurate Proteome Analysis. Molecular and Cellular Proteomics, 2012, 11, 0111.016717. | 3.8 | 2,285 |