

Sule YÄ±lmaz

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4947146/publications.pdf>

Version: 2024-02-01

12
papers

6,951
citations

933447

10
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

20260
citing authors

#	ARTICLE	IF	CITATIONS
1	The PRIDE database and related tools and resources in 2019: improving support for quantification data. <i>Nucleic Acids Research</i> , 2019, 47, D442-D450.	14.5	6,449
2	Enhancement of grape seed oil extraction using a cell wall degrading enzyme cocktail. <i>Food Chemistry</i> , 2009, 115, 48-53.	8.2	129
3	MaxDIA enables library-based and library-free data-independent acquisition proteomics. <i>Nature Biotechnology</i> , 2021, 39, 1563-1573.	17.5	115
4	First Community-Wide, Comparative Cross-Linking Mass Spectrometry Study. <i>Analytical Chemistry</i> , 2019, 91, 6953-6961.	6.5	100
5	MaxQuant goes Linux. <i>Nature Methods</i> , 2018, 15, 401-401.	19.0	41
6	Cross-linked peptide identification: A computational forest of algorithms. <i>Mass Spectrometry Reviews</i> , 2018, 37, 738-749.	5.4	27
7	Xilmass: A New Approach toward the Identification of Cross-Linked Peptides. <i>Analytical Chemistry</i> , 2016, 88, 9949-9957.	6.5	25
8	Accurate and Automated High-Coverage Identification of Chemically Cross-Linked Peptides with MaxLynx. <i>Analytical Chemistry</i> , 2022, 94, 1608-1617.	6.5	19
9	Methods to Calculate Spectrum Similarity. <i>Methods in Molecular Biology</i> , 2017, 1549, 75-100.	0.9	18
10	Asn ₃ , a Reliable, Robust, and Universal Lock Mass for Improved Accuracy in LC-MS and LC-MS/MS. <i>Analytical Chemistry</i> , 2013, 85, 11054-11060.	6.5	13
11	A Pipeline for Differential Proteomics in Unsequenced Species. <i>Journal of Proteome Research</i> , 2016, 15, 1963-1970.	3.7	10
12	From challenges to perspectives. <i>EMBO Reports</i> , 2014, 15, 1010-1014.	4.5	0