Prasad Phapale

List of Publications by Year in descending order

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623734 610901 4,098 25 14 24 citations g-index h-index papers 31 31 31 6777 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Public LC-Orbitrap Tandem Mass Spectral Library for Metabolite Identification. Journal of Proteome Research, 2021, 20, 2089-2097.	3.7	18
2	Coupling proteomics and metabolomics for the unsupervised identification of protein–metabolite interactions in Chaetomium thermophilum. PLoS ONE, 2021, 16, e0254429.	2.5	5
3	SpaceM reveals metabolic states of single cells. Nature Methods, 2021, 18, 799-805.	19.0	170
4	Pharmacoâ€metabolomics opportunities in drug development and clinical research. Analytical Science Advances, 2021, 2, 611-616.	2.8	6
5	Bioaccumulation of therapeutic drugs by human gut bacteria. Nature, 2021, 597, 533-538.	27.8	159
6	Quantification of Duloxetine in the Bacterial Culture and Medium to Study Drug-gut Microbiome Interactions. Bio-protocol, 2021, 11, e4214.	0.4	2
7	Untargeted Metabolomics Workshop Report: Quality Control Considerations from Sample Preparation to Data Analysis. Journal of the American Society for Mass Spectrometry, 2020, 31, 2006-2010.	2.8	11
8	GPD1 Specifically Marks Dormant Glioma Stem Cells with a Distinct Metabolic Profile. Cell Stem Cell, 2019, 25, 241-257.e8.	11.1	66
9	3D molecular cartography using LC–MS facilitated by Optimus and 'ili software. Nature Protocols, 2018, 13, 134-154.	12.0	85
10	Curatr: a web application for creating, curating and sharing a mass spectral library. Bioinformatics, 2018, 34, 1436-1438.	4.1	14
11	Endogenous Fatty Acids Are Essential Signaling Factors of Pancreatic \hat{l}^2 -Cells and Insulin Secretion. Diabetes, 2018, 67, 1986-1998.	0.6	48
12	Capturing protein communities by structural proteomics in a thermophilic eukaryote. Molecular Systems Biology, 2017, 13, 936.	7.2	108
13	FDR-controlled metabolite annotation for high-resolution imaging mass spectrometry. Nature Methods, 2017, 14, 57-60.	19.0	314
14	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. Nature Biotechnology, 2016, 34, 828-837.	17.5	2,802
15	Regulatory Norms and Intellectual Property Rights for Biomarker Research., 2016,, 99-115.		1
16	An integrated approach for genome annotation of the eukaryotic thermophile Chaetomium thermophilum. Nucleic Acids Research, 2014, 42, 13525-13533.	14.5	55
17	Polyketide synthesis in tobacco plants transformed with a Plumbago zeylanica type III hexaketide synthase. Phytochemistry, 2014, 98, 92-100.	2.9	7
18	Expedient preparative isolation, quantification and characterization of limonoids from Neem fruits. Analytical Methods, 2013, 5, 5386.	2.7	22

#	Article	IF	CITATION
19	In vitro inhibitory effects of Wen-pi-tang-Hab-Wu-ling-san on human cytochrome P450 isoforms. Journal of Clinical Pharmacy and Therapeutics, 2011, 36, 496-503.	1.5	5
20	Analysis of Pazufloxacin Mesilate in Human Plasma and Urine by LC with Fluorescence and UV Detection, and Its Application to Pharmacokinetic Study. Chromatographia, 2010, 71, 101-106.	1.3	9
21	Rapid determination of finasteride in human plasma by UPLC–MS/MS and its application to clinical pharmacokinetic study. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 1718-1723.	2.3	20
22	Liquid chromatography–tandem mass spectrometry quantification of levosulpiride in human plasma and its application to bioequivalence study. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 2280-2285.	2.3	9
23	An Integrative Approach for Identifying a Metabolic Phenotype Predictive of Individualized Pharmacokinetics of Tacrolimus. Clinical Pharmacology and Therapeutics, 2010, 87, 426-436.	4.7	73
24	Revival of TE2A; a better chelate for Cu(II) ions than TETA?. Chemical Communications, 2010, 46, 3517.	4.1	53
25	An Improved UPLC Method for Rapid Analysis of Levofloxacin in Human Plasma. Chromatographia, 2008, 68, 187-192.	1.3	9