Panagiotis A Vorkas

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4082393/publications.pdf

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22 papers

717 citations

623734 14 h-index 677142 22 g-index

22 all docs 22 docs citations

times ranked

22

1396 citing authors

#	Article	IF	CITATIONS
1	Automated Annotation of Untargeted All-Ion Fragmentation LC–MS Metabolomics Data with MetaboAnnotatoR. Analytical Chemistry, 2022, 94, 3446-3455.	6.5	18
2	Lipid profiling of mouse intestinal organoids for studying <i> APC </i> mutations. Bioscience Reports, 2021, 41, .	2.4	5
3	Characterizing the breast cancer lipidome and its interaction with the tissue microbiota. Communications Biology, 2021, 4, 1229.	4.4	17
4	Metabolic Signatures of Gestational Weight Gain and Postpartum Weight Loss in a Lifestyle Intervention Study of Overweight and Obese Women. Metabolites, 2020, 10, 498.	2.9	5
5	Tissue Multiplatform-Based Metabolomics/Metabonomics for Enhanced Metabolome Coverage. Methods in Molecular Biology, 2018, 1738, 239-260.	0.9	8
6	Expanding lipidome coverage using MS/MS-aided untargeted data-independent RPâ€"UPLCâ€"TOFâ€"MS ^E acquisition. Bioanalysis, 2018, 10, 307-319.	1.5	5
7	Application of Metabolic Profiling to Abdominal Aortic Aneurysm Research. Journal of Proteome Research, 2017, 16, 2325-2332.	3.7	10
8	Lessons from Metabonomics on the Neurobiology of Stroke. Neuroscientist, 2017, 23, 374-382.	3.5	34
9	Metabolic Phenotypes of Carotid Atherosclerotic Plaques Relate to Stroke Risk: An Exploratory Study. European Journal of Vascular and Endovascular Surgery, 2016, 52, 5-10.	1.5	32
10	Replication of LC–MS untargeted lipidomics results in patients with calcific coronary disease: An interlaboratory reproducibility study. International Journal of Cardiology, 2016, 222, 1042-1048.	1.7	13
11	Discovery of Infection Associated Metabolic Markers in Human African Trypanosomiasis. PLoS Neglected Tropical Diseases, 2015, 9, e0004200.	3.0	48
12	Untargeted UPLC-MS Profiling Pipeline to Expand Tissue Metabolome Coverage: Application to Cardiovascular Disease. Analytical Chemistry, 2015, 87, 4184-4193.	6.5	161
13	Metabolic Phenotyping of Atherosclerotic Plaques Reveals Latent Associations between Free Cholesterol and Ceramide Metabolism in Atherogenesis. Journal of Proteome Research, 2015, 14, 1389-1399.	3.7	65
14	Perturbations in fatty acid metabolism and apoptosis are manifested in calcific coronary artery disease: An exploratory lipidomic study. International Journal of Cardiology, 2015, 197, 192-199.	1.7	29
15	Optimization of metabolite extraction of human vein tissue for ultra performance liquid chromatography-mass spectrometry and nuclear magnetic resonance-based untargeted metabolic profiling. Analyst, The, 2015, 140, 7586-7597.	3.5	30
16	Chemical mapping of the colorectal cancer microenvironment via MALDI imaging mass spectrometry (MALDIâ∈MSI) reveals novel cancerâ€associated field effects. Molecular Oncology, 2014, 8, 39-49.	4.6	95
17	Systems Biology of Human Atherosclerosis. Vascular and Endovascular Surgery, 2014, 48, 5-17.	0.7	26
18	Implementation of Molecular Phenotyping Approaches in the Personalized Surgical Patient Journey. Annals of Surgery, 2012, 255, 881-889.	4.2	34

#	Article	lF	CITATION
19	A closed-tube methylation-sensitive high resolution melting assay (MS-HRMA) for the semi-quantitative determination of CST6 promoter methylation in clinical samples. BMC Cancer, 2012, 12, 486.	2.6	14
20	In-vitro Identification of Distinctive Metabolic Signatures of Intact Varicose Vein Tissue via Magic Angle Spinning Nuclear Magnetic Resonance Spectroscopy. European Journal of Vascular and Endovascular Surgery, 2012, 44, 442-450.	1.5	18
21	Mutation scanning of exon 20 of the BRCA1 gene by high-resolution melting curve analysis. Clinical Biochemistry, 2010, 43, 178-185.	1.9	13
22	PIK3CA Hotspot Mutation Scanning by a Novel and Highly Sensitive High-Resolution Small Amplicon Melting Analysis Method. Journal of Molecular Diagnostics, 2010, 12, 697-704.	2.8	37