## Etienne A Thévenot

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7669315/publications.pdf

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		840776	888059
17	1,750 citations	11	17
papers	citations	h-index	g-index
19	19	19	3245
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Analysis of the Human Adult Urinary Metabolome Variations with Age, Body Mass Index, and Gender by Implementing a Comprehensive Workflow for Univariate and OPLS Statistical Analyses. Journal of Proteome Research, 2015, 14, 3322-3335.	3.7	886
2	Workflow4Metabolomics: a collaborative research infrastructure for computational metabolomics. Bioinformatics, 2015, 31, 1493-1495.	4.1	333
3	Metabolomics of exhaled breath in critically ill COVID-19 patients: A pilot study. EBioMedicine, 2021, 63, 103154.	6.1	143
4	Create, run, share, publish, and reference your LC–MS, FIA–MS, GC–MS, and NMR data analysis workflows with the Workflow4Metabolomics 3.0 Galaxy online infrastructure for metabolomics. International Journal of Biochemistry and Cell Biology, 2017, 93, 89-101.	2.8	99
5	The metaRbolomics Toolbox in Bioconductor and beyond. Metabolites, 2019, 9, 200.	2.9	64
6	PhenoMeNal: processing and analysis of metabolomics data in the cloud. GigaScience, 2019, 8, .	6.4	60
7	biosigner: A New Method for the Discovery of Significant Molecular Signatures from Omics Data. Frontiers in Molecular Biosciences, 2016, 3, 26.	<b>3.</b> 5	52
8	The future of metabolomics in ELIXIR. F1000Research, 2017, 6, 1649.	1.6	19
9	Metabolic reprograming of LPS-stimulated human lung macrophages involves tryptophan metabolism and the aspartate-arginosuccinate shunt. PLoS ONE, 2020, 15, e0230813.	2.5	18
10	Multiplatform metabolomics for an integrative exploration of metabolic syndrome in older men. EBioMedicine, 2021, 69, 103440.	6.1	18
11	<i>proFIA</i> : a data preprocessing workflow for flow injection analysis coupled to high-resolution mass spectrometry. Bioinformatics, 2017, 33, 3767-3775.	4.1	14
12	The future of metabolomics in ELIXIR. F1000Research, 2017, 6, 1649.	1.6	11
13	Differential Effects of Post-Weaning Diet and Maternal Obesity on Mouse Liver and Brain Metabolomes. Nutrients, 2020, 12, 1572.	4.1	8
14	ptairMS: real-time processing and analysis of PTR-TOF-MS data for biomarker discovery in exhaled breath. Bioinformatics, 2022, 38, 1930-1937.	4.1	8
15	ProMetIS, deep phenotyping of mouse models by combined proteomics and metabolomics analysis. Scientific Data, 2021, 8, 311.	<b>5.</b> 3	6
16	Bariatric Surgery Improves the Atherogenic Profile of Circulating Methylarginines in Obese Patients: Results from a Pilot Study. Metabolites, 2021, 11, 759.	2.9	4
17	PeakForest: a multi-platform digital infrastructure for interoperable metabolite spectral data and metadata management. Metabolomics, 2022, $18, \ldots$	3.0	4