## Yann Guitton

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4539997/publications.pdf

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52 1,325 22 34
papers citations h-index g-index

57 57 57 2084 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	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.	1.2	99
2	Essential Oils from Wild Populations of Algerian <i>Lavandula stoechas</i> L.: Composition, Chemical Variability, and <i>in vitro</i> Biological Properties. Chemistry and Biodiversity, 2011, 8, 937-953.	1.0	82
3	Auto-deconvolution and molecular networking of gas chromatography–mass spectrometry data. Nature Biotechnology, 2021, 39, 169-173.	9.4	78
4	Multidimensional NMR approaches towards highly resolved, sensitive and high-throughput quantitative metabolomics. Current Opinion in Biotechnology, 2017, 43, 49-55.	3.3	65
5	A European proposal for quality control and quality assurance of tandem mass spectral libraries. Environmental Sciences Europe, 2020, 32, .	2.6	53
6	HaloSeeker 1.0: A User-Friendly Software to Highlight Halogenated Chemicals in Nontargeted High-Resolution Mass Spectrometry Data Sets. Analytical Chemistry, 2019, 91, 3500-3507.	3.2	52
7	Differential accumulation of volatile terpene and terpene synthase mRNAs during lavender ( $<$ i>Lavandula angustifolia $<$  i> and $<$ i>L. $<$  i> x $<$ i>intermedia $<$  i>) inflorescence development. Physiologia Plantarum, 2010, 138, 150-163.	2.6	50
8	Breast Milk Lipidome Is Associated with Early Growth Trajectory in Preterm Infants. Nutrients, 2018, 10, 164.	1.7	49
9	Isolation and functional characterization of a Ï"-cadinol synthase, a new sesquiterpene synthase from Lavandula angustifolia. Plant Molecular Biology, 2014, 84, 227-241.	2.0	48
10	Single-Step Extraction Coupled with Targeted HILIC-MS/MS Approach for Comprehensive Analysis of Human Plasma Lipidome and Polar Metabolome. Metabolites, 2020, 10, 495.	1.3	46
11	Automated Detection of Natural Halogenated Compounds from LC-MS Profiles–Application to the Isolation of Bioactive Chlorinated Compounds from Marine-Derived Fungi. Analytical Chemistry, 2016, 88, 9143-9150.	<b>3.</b> 2	43
12	Rapid evaporative ionisation mass spectrometry and chemometrics for high-throughput screening of growth promoters in meat producing animals. Food Additives and Contaminants - Part A Chemistry, Analysis, Control, Exposure and Risk Assessment, 2018, 35, 900-910.	1.1	37
13	A multidimensional 1H NMR lipidomics workflow to address chemical food safety issues. Metabolomics, 2018, 14, 60.	1.4	32
14	Making complex measurements of meat composition fast: Application of rapid evaporative ionisation mass spectrometry to measuring meat quality and fraud. Meat Science, 2021, 181, 108333.	2.7	30
15	MSeasy: unsupervised and untargeted GC-MS data processing. Bioinformatics, 2012, 28, 2278-2280.	1.8	29
16	Dye residues in aquaculture products: Targeted and metabolomics mass spectrometric approaches to track their abuse. Food Chemistry, 2019, 294, 355-367.	4.2	28
17	Associations between persistent organic pollutants and endometriosis: A multiblock approach integrating metabolic and cytokine profiling. Environment International, 2022, 158, 106926.	4.8	27
18	Time Dependency of Chemodiversity and Biosynthetic Pathways: An LC-MS Metabolomic Study of Marine-Sourced Penicillium. Marine Drugs, 2016, 14, 103.	2,2	26

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19	Serum-based metabolomics characterization of pigs treated with ractopamine. Metabolomics, 2017, 13, 1.	1.4	26
20	Comprehensive Preterm Breast Milk Metabotype Associated with Optimal Infant Early Growth Pattern. Nutrients, 2019, 11, 528.	1.7	26
21	Cytotoxicity and mycotoxin production of shellfish-derived <i>Penicillium</i> spp., a risk for shellfish consumers. Letters in Applied Microbiology, 2013, 57, 385-392.	1.0	25
22	Non-targeted screening methodology to characterise human internal chemical exposure: Application to halogenated compounds in human milk. Talanta, 2021, 225, 121979.	2.9	25
23	Optimized characterization of short-, medium, and long-chain chlorinated paraffins in liquid chromatography-high resolution mass spectrometry. Journal of Chromatography A, 2020, 1619, 460927.	1.8	23
24	Elucidation of non-intentionally added substances migrating from polyester-polyurethane lacquers using automated LC-HRMS data processing. Analytical and Bioanalytical Chemistry, 2018, 410, 5391-5403.	1.9	22
25	Resolution of quantitative resistance to clubroot into QTL-specific metabolic modules. Journal of Experimental Botany, 2019, 70, 5375-5390.	2.4	22
26	Functional characterization of terpene synthases and chemotypic variation in three lavender species of section Stoechas. Physiologia Plantarum, 2015, 153, 43-57.	2.6	19
27	Optimization of fecal sample preparation for untargeted LC-HRMS based metabolomics. Metabolomics, 2017, 13, 1.	1.4	19
28	WiPP: Workflow for Improved Peak Picking for Gas Chromatography-Mass Spectrometry (GC-MS) Data. Metabolites, 2019, 9, 171.	1.3	19
29	Ammonium Fluoride as Suitable Additive for HILIC-Based LC-HRMS Metabolomics. Metabolites, 2019, 9, 292.	1.3	19
30	Lavender inflorescence. Plant Signaling and Behavior, 2010, 5, 749-751.	1.2	18
31	Nontargeted LC/ESI-HRMS Detection of Polyhalogenated Compounds in Marine Mammals Stranded on French Atlantic Coasts. ACS ES&T Water, 2021, 1, 309-318.	2.3	16
32	Modeling the fragmentation patterns of triacylglycerides in mass spectrometry allows the quantification of the regioisomers with a minimal number of standards. Analytica Chimica Acta, 2019, 1057, 60-69.	2.6	15
33	Simultaneous exploration of nutrients and pollutants in human milk and their impact on preterm infant growth: An integrative cross-platform approach. Environmental Research, 2020, 182, 109018.	3.7	15
34	Genome size and plastid trnK-matK markers give new insights into the evolutionary history of the genus <i>Lavandula</i> L Plant Biosystems, 2016, 150, 1216-1224.	0.8	14
35	Fungi isolated from Madagascar shrimps - investigation of the Aspergillus niger metabolism by combined LC-MS and NMR metabolomics studies. Aquaculture, 2017, 479, 750-758.	1.7	13
36	A comparative study of terpene composition in different clades of the genus Lavandula. Botany Letters, 2018, 165, 494-505.	0.7	13

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37	Cytotoxicity, Fractionation and Dereplication of Extracts of the Dinoflagellate Vulcanodinium rugosum, a Producer of Pinnatoxin G. Marine Drugs, 2013, 11, 3350-3371.	2.2	12
38	Consequences of blunting the mevalonate pathway in cancer identified by a pluri-omics approach. Cell Death and Disease, 2018, 9, 745.	2.7	12
39	Dereplication of Mammea neurophylla metabolites to isolate original 4-phenylcoumarins. Phytochemistry Letters, 2015, 11, 61-68.	0.6	10
40	Synthesis and antiproliferative activity of ligerin and new fumagillin analogs against osteosarcoma. European Journal of Medicinal Chemistry, 2014, 79, 244-250.	2.6	8
41	Successes and pitfalls in automated dereplication strategy using liquid chromatography coupled to mass spectrometry data: A CASMI 2016 experience. Phytochemistry Letters, 2017, 21, 297-305.	0.6	8
42	Untargeted Lipidomic Profiling of Dry Blood Spots Using SFC-HRMS. Metabolites, 2021, 11, 305.	1.3	8
43	From a nonâ€targeted metabolomics approach to a targeted biomarkers strategy to highlight testosterone abuse in equine. Illustration of a methodological transfer between platforms and laboratories. Drug Testing and Analysis, 2022, 14, 864-878.	1.6	8
44	Electrospray ionization and heterogeneous matrix effects in liquid chromatography/mass spectrometry based metaâ€metabolomics: A biomarker or a suppressed ion?. Rapid Communications in Mass Spectrometry, 2021, 35, e8977.	0.7	7
45	Metabolomics and lipidomics to identify biomarkers of effect related to exposure to non-dioxin-like polychlorinated biphenyls in pigs. Chemosphere, 2022, 296, 133957.	4.2	5
46	Extending the Lipidome Coverage by Combining Different Mass Spectrometric Platforms: An Innovative Strategy to Answer Chemical Food Safety Issues. Foods, 2021, 10, 1218.	1.9	4
47	Urinary metabolomic profiling from spontaneous tolerant kidney transplanted recipients shows enrichment in tryptophan-derived metabolites. EBioMedicine, 2022, 77, 103844.	2.7	4
48	Nandrolone and estradiol biomarkers identification in bovine urine applying a liquid chromatography highâ€resolution mass spectrometry metabolomics approach. Drug Testing and Analysis, 2021, , .	1.6	3
49	Successes and Pitfalls in Automated Dereplication Strategy Using Mass Spectrometry Data: a CASMI Experience. Current Metabolomics, 2017, 5, 25-34.	0.5	2
50	Combining MS/MS fragmentation, correlation and biochemical reaction networks to improve compound annotation in metabolome investigations of marine-derived Penicillium species. Planta Medica, 2014, 80, .	0.7	0
51	Automated MS/MS data annotation: CASMI experiences. Planta Medica, 2016, 81, S1-S381.	0.7	0
52	Marine halogenated compound analysis: from an R package to the isolation of new griseophenone derivatives. Planta Medica, 2016, 81, S1-S381.	0.7	0