Juan Carlos GarcÃ-a-Cañaveras

List of Publications by Year in descending order

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Version: 2024-02-01

33 papers

2,237 citations

331642 21 h-index 414395 32 g-index

35 all docs 35 docs citations

35 times ranked 3994 citing authors

#	Article	IF	CITATIONS
1	MTHFD2 is a metabolic checkpoint controlling effector and regulatory TÂcell fate and function. Immunity, 2022, 55, 65-81.e9.	14.3	74
2	SHMT inhibition is effective and synergizes with methotrexate in T-cell acute lymphoblastic leukemia. Leukemia, 2021, 35, 377-388.	7.2	68
3	Reviewing the metabolome coverage provided by LC-MS: Focus on sample preparation and chromatography-A tutorial. Analytica Chimica Acta, 2021, 1147, 38-55.	5.4	40
4	Tumor Microenvironment-Derived Metabolites: A Guide to Find New Metabolic Therapeutic Targets and Biomarkers. Cancers, 2021, 13, 3230.	3.7	17
5	c-MYC Triggers Lipid Remodelling During Early Somatic Cell Reprogramming to Pluripotency. Stem Cell Reviews and Reports, 2021, 17, 2245-2261.	3.8	6
6	CAR T-Cells Depend on the Coupling of NADH Oxidation with ATP Production. Cells, 2021, 10, 2334.	4.1	7
7	Enhancing Chimeric Antigen Receptor T Cell Anti-tumor Function through Advanced Media Design. Molecular Therapy - Methods and Clinical Development, 2020, 18, 595-606.	4.1	39
8	Endothelin-1–Mediated Drug Resistance in <i>EGFR</i> -Mutant Non-Small Cell Lung Carcinoma. Cancer Research, 2020, 80, 4224-4232.	0.9	12
9	Chaperone-mediated autophagy regulates the pluripotency of embryonic stem cells. Science, 2020, 369, 397-403.	12.6	60
10	Novel media formulations to enhance Chimeric Antigen Receptor (CAR) T-cell potency and anti-tumor cell function for adoptive immunotherapy. Cytotherapy, 2020, 22, S133.	0.7	3
11	Obesity Shapes Metabolism in the Tumor Microenvironment to Suppress Anti-Tumor Immunity. Cell, 2020, 183, 1848-1866.e26.	28.9	347
12	A small molecule G6PD inhibitor reveals immune dependence on pentose phosphate pathway. Nature Chemical Biology, 2020, 16, 731-739.	8.0	101
13	Serine Catabolism Feeds NADH when Respiration Is Impaired. Cell Metabolism, 2020, 31, 809-821.e6.	16.2	118
14	Lactate dehydrogenase inhibition synergizes with IL-21 to promote CD8 ⁺ T cell stemness and antitumor immunity. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 6047-6055.	7.1	128
15	Distinct modes of mitochondrial metabolism uncouple T cell differentiation and function. Nature, 2019, 571, 403-407.	27.8	156
16	The Tumor Metabolic Microenvironment: Lessons from Lactate. Cancer Research, 2019, 79, 3155-3162.	0.9	140
17	Sparse N-way partial least squares by L1-penalization. Chemometrics and Intelligent Laboratory Systems, 2019, 185, 85-91.	3.5	7
18	LipidMS: An R Package for Lipid Annotation in Untargeted Liquid Chromatography-Data Independent Acquisition-Mass Spectrometry Lipidomics. Analytical Chemistry, 2019, 91, 836-845.	6.5	33

#	Article	lF	Citations
19	PPAR \hat{I}^3 is a nexus controlling alternative activation of macrophages via glutamine metabolism. Genes and Development, 2018, 32, 1035-1044.	5.9	84
20	A lipidomic cellâ€based assay for studying drugâ€induced phospholipidosis and steatosis. Electrophoresis, 2017, 38, 2331-2340.	2.4	18
21	RpeakChrom: Novel R package for the automated characterization and optimization of column efficiency in highâ€performance liquid chromatography analysis. Electrophoresis, 2017, 38, 2985-2995.	2.4	3
22	Liver Transplantation Biomarkers in the Metabolomics Era. Biomarkers in Disease, 2017, , 99-128.	0.1	2
23	A metabolomics cell-based approach for anticipating and investigating drug-induced liver injury. Scientific Reports, 2016, 6, 27239.	3.3	67
24	Extending metabolome coverage for untargeted metabolite profiling of adherent cultured hepatic cells. Analytical and Bioanalytical Chemistry, 2016, 408, 1217-1230.	3.7	32
25	Liver Transplantation Biomarkers in the Metabolomics Era. Biomarkers in Disease, 2016, , 1-29.	0.1	0
26	LCâ€MS untargeted metabolomic analysis of drugâ€induced hepatotoxicity in HepG2 cells. Electrophoresis, 2015, 36, 2294-2302.	2.4	32
27	In vitro/in vivo screening of oxidative homeostasis and damage to DNA, protein, and lipids using UPLC/MS-MS. Analytical and Bioanalytical Chemistry, 2014, 406, 5465-5476.	3.7	20
28	Ultra-Performance Liquid Chromatography-Mass Spectrometry Targeted Profiling of Bile Acids: Application to Serum, Liver Tissue, and Cultured Cells of Different Species. Methods in Molecular Biology, 2014, 1198, 233-247.	0.9	8
29	Metabolomics discloses donor liver biomarkers associated with early allograft dysfunction. Journal of Hepatology, 2014, 61, 564-574.	3.7	63
30	Mammalian cell metabolomics: Experimental design and sample preparation. Electrophoresis, 2013, 34, 2762-2775.	2.4	163
31	Targeted profiling of circulating and hepatic bile acids in human, mouse, and rat using a UPLC-MRM-MS-validated method. Journal of Lipid Research, 2012, 53, 2231-2241.	4.2	220
32	Chemometric approaches to improve PLSDA model outcome for predicting human non-alcoholic fatty liver disease using UPLC-MS as a metabolic profiling tool. Metabolomics, 2012, 8, 86-98.	3.0	54
33	A Comprehensive Untargeted Metabonomic Analysis of Human Steatotic Liver Tissue by RP and HILIC Chromatography Coupled to Mass Spectrometry Reveals Important Metabolic Alterations. Journal of Proteome Research, 2011, 10, 4825-4834.	3.7	114