Ian A Lewis

List of Publications by Year in descending order

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44 papers

3,560 citations

236925 25 h-index 276875 41 g-index

52 all docs 52 docs citations

52 times ranked 5382 citing authors

#	Article	IF	CITATIONS
1	Crohn's disease therapeutic dietary intervention (CD-TDI): study protocol for a randomised controlled trial. BMJ Open Gastroenterology, 2022, 9, e000841.	2.7	O
2	Dipeptidase-1 governs renal inflammation during ischemia reperfusion injury. Science Advances, 2022, 8, eabm0142.	10.3	28
3	Colitis-associated microbiota drives changes in behaviour in male mice in the absence of inflammation. Brain, Behavior, and Immunity, 2022, 102, 266-278.	4.1	19
4	Method for absolute quantification of short chain fatty acids via reverse phase chromatography mass spectrometry. PLoS ONE, 2022, 17, e0267093.	2.5	16
5	Metabolic preference assay for rapid diagnosis of bloodstream infections. Nature Communications, 2022, 13, 2332.	12.8	20
6	Methods for Quantifying the Metabolic Boundary Fluxes of Cell Cultures in Large Cohorts by High-Resolution Hydrophilic Liquid Chromatography Mass Spectrometry. Analytical Chemistry, 2022, 94, 8874-8882.	6.5	11
7	Bacterial cyclic diguanylate signaling networks sense temperature. Nature Communications, 2021, 12, 1986.	12.8	35
8	Staphylococcus aureus induces an itaconate-dominated immunometabolic response that drives biofilm formation. Nature Communications, 2021, 12, 1399.	12.8	72
9	Unique metabolic phenotype and its transition during maturation of juvenile male germ cells. FASEB Journal, 2021, 35, e21513.	0.5	19
10	Cooperation between host immunity and the gut bacteria is essential for helminth-evoked suppression of colitis. Microbiome, 2021, 9, 186.	11,1	28
11	Untargeted Metabolomics Investigation on Selenite Reduction to Elemental Selenium by Bacillus mycoides SelTE01. Frontiers in Microbiology, 2021, 12, 711000.	3.5	6
12	Thermogenic hydrocarbon biodegradation by diverse depth-stratified microbial populations at a Scotian Basin cold seep. Nature Communications, 2020, 11, 5825.	12.8	72
13	Microbiome-derived inosine modulates response to checkpoint inhibitor immunotherapy. Science, 2020, 369, 1481-1489.	12.6	635
14	PelX is a UDP-N-acetylglucosamine C4-epimerase involved in Pel polysaccharide–dependent biofilm formation. Journal of Biological Chemistry, 2020, 295, 11949-11962.	3.4	10
15	Colitis-Induced Microbial Perturbation Promotes Postinflammatory Visceral Hypersensitivity. Cellular and Molecular Gastroenterology and Hepatology, 2020, 10, 225-244.	4.5	33
16	Optimized serial expansion of human induced pluripotent stem cells using low-density inoculation to generate clinically relevant quantities in vertical-wheel bioreactors. Stem Cells Translational Medicine, 2020, 9, 1036-1052.	3.3	40
17	Vancomycin relieves mycophenolate mofetil–induced gastrointestinal toxicity by eliminating gut bacterial β-glucuronidase activity. Science Advances, 2019, 5, eaax2358.	10.3	73
18	Metabolic potential of uncultured bacteria and archaea associated with petroleum seepage in deep-sea sediments. Nature Communications, 2019, 10, 1816.	12.8	118

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19	nmrML: A Community Supported Open Data Standard for the Description, Storage, and Exchange of NMR Data. Analytical Chemistry, 2018, 90, 649-656.	6.5	50
20	Evidence for Regulation of Hemoglobin Metabolism and Intracellular Ionic Flux by the Plasmodium falciparum Chloroquine Resistance Transporter. Scientific Reports, 2018, 8, 13578.	3.3	24
21	Iron Sequestration in Microbiota Biofilms As A Novel Strategy for Treating Inflammatory Bowel Diseases, 2018, 24, 1493-1502.	1.9	30
22	An intact microbiota is required for the gastrointestinal toxicity of the immunosuppressant mycophenolate mofetil. Journal of Heart and Lung Transplantation, 2018, 37, 1047-1059.	0.6	59
23	Editorial overview: Recent innovations in the metabolomics revolution. Current Opinion in Biotechnology, 2017, 43, iv-vii.	6.6	7
24	Digestomics: an emerging strategy for comprehensive analysis of protein catabolism. Current Opinion in Biotechnology, 2017, 43, 134-140.	6.6	11
25	Metabolite Measurement: Pitfalls to Avoid and Practices to Follow. Annual Review of Biochemistry, 2017, 86, 277-304.	11.1	322
26	Understanding Plant Nitrogen Metabolism through Metabolomics and Computational Approaches. Plants, 2016, 5, 39.	3.5	41
27	Evolution of Fitness Cost-Neutral Mutant PfCRT Conferring P. falciparum 4-Aminoquinoline Drug Resistance Is Accompanied by Altered Parasite Metabolism and Digestive Vacuole Physiology. PLoS Pathogens, 2016, 12, e1005976.	4.7	34
28	Hierarchy in Pentose Sugar Metabolism in Clostridium acetobutylicum. Applied and Environmental Microbiology, 2015, 81, 1452-1462.	3.1	38
29	Genetic Investigation of Tricarboxylic Acid Metabolism during the Plasmodium falciparum Life Cycle. Cell Reports, 2015, 11, 164-174.	6.4	134
30	Metabolic QTL Analysis Links Chloroquine Resistance in Plasmodium falciparum to Impaired Hemoglobin Catabolism. PLoS Genetics, 2014, 10, e1004085.	3.5	73
31	Kinetic Flux Profiling Elucidates Two Independent Acetyl-CoA Biosynthetic Pathways in Plasmodium falciparum. Journal of Biological Chemistry, 2013, 288, 36338-36350.	3.4	79
32	Semiautomated Device for Batch Extraction of Metabolites from Tissue Samples. Analytical Chemistry, 2012, 84, 1809-1812.	6.5	6
33	Novel NMR and MS Approaches to Metabolomics. Methods in Pharmacology and Toxicology, 2012, , 199-230.	0.2	4
34	Relationship between treatment-seeking behaviour and artemisinin drug quality in Ghana. Malaria Journal, 2012, 11, 110.	2.3	12
35	Role of aminotransferases in glutamate metabolism of human erythrocytes. Journal of Biomolecular NMR, 2011, 49, 221-229.	2.8	46
36	NMR Method for Measuring Carbon-13 Isotopic Enrichment of Metabolites in Complex Solutions. Analytical Chemistry, 2010, 82, 4558-4563.	6.5	38

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37	Role of band 3 in regulating metabolic flux of red blood cells. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 18515-18520.	7.1	109
38	rNMR: open source software for identifying and quantifying metabolites in NMR spectra. Magnetic Resonance in Chemistry, 2009, 47, S123-6.	1.9	169
39	Metabolite identification via the Madison Metabolomics Consortium Database. Nature Biotechnology, 2008, 26, 162-164.	17.5	591
40	Stable Isotope Assisted Assignment of Elemental Compositions for Metabolomics. Analytical Chemistry, 2007, 79, 6912-6921.	6.5	90
41	Method for Determining Molar Concentrations of Metabolites in Complex Solutions from Two-Dimensional ¹ Hâ^' ¹³ C NMR Spectra. Analytical Chemistry, 2007, 79, 9385-9390.	6.5	262
42	New bioinformatics resources for metabolomics. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2007, , 157-68.	0.7	42
43	NEW BIOINFORMATICS RESOURCES FOR METABOLOMICS. , 2006, , .		43
44	Metabolomics: The Key to Unraveling the Role of the Microbiome in Visceral Pain Neurotransmission. Frontiers in Neuroscience, $0, 16, .$	2.8	3