

Christian Diener

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5973117/publications.pdf>

Version: 2024-02-01

27
papers

13,076
citations

623734

14
h-index

526287

27
g-index

42
all docs

42
docs citations

42
times ranked

16676
citing authors

#	ARTICLE	IF	CITATIONS
1	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019, 37, 852-857.	17.5	11,167
2	MEMOTE for standardized genome-scale metabolic model testing. <i>Nature Biotechnology</i> , 2020, 38, 272-276.	17.5	314
3	Gut microbiome pattern reflects healthy ageing and predicts survival in humans. <i>Nature Metabolism</i> , 2021, 3, 274-286.	11.9	278
4	Use and abuse of correlation analyses in microbial ecology. <i>ISME Journal</i> , 2019, 13, 2647-2655.	9.8	193
5	MICOM: Metagenome-Scale Modeling To Infer Metabolic Interactions in the Gut Microbiota. <i>MSystems</i> , 2020, 5, .	3.8	126
6	Distinct microbes, metabolites, and ecologies define the microbiome in deficient and proficient mismatch repair colorectal cancers. <i>Genome Medicine</i> , 2018, 10, 78.	8.2	107
7	Synthesis of multi-omic data and community metabolic models reveals insights into the role of hydrogen sulfide in colon cancer. <i>Methods</i> , 2018, 149, 59-68.	3.8	63
8	Cell Penetrating Peptides and Cationic Antibacterial Peptides. <i>Journal of Biological Chemistry</i> , 2014, 289, 14448-14457.	3.4	49
9	Genomic and functional characterization of a mucosal symbiont involved in early-stage colorectal cancer. <i>Cell Host and Microbe</i> , 2021, 29, 1589-1598.e6.	11.0	44
10	Effective Design of Multifunctional Peptides by Combining Compatible Functions. <i>PLoS Computational Biology</i> , 2016, 12, e1004786.	3.2	36
11	Onset of Immune Senescence Defined by Unbiased Pyrosequencing of Human Immunoglobulin mRNA Repertoires. <i>PLoS ONE</i> , 2012, 7, e49774.	2.5	30
12	Yeast Mating and Image-Based Quantification of Spatial Pattern Formation. <i>PLoS Computational Biology</i> , 2014, 10, e1003690.	3.2	25
13	Heterogeneity in statin responses explained by variation in the human gut microbiome. <i>Med</i> , 2022, 3, 388-405.e6.	4.4	21
14	From taxonomy to metabolic output: what factors define gut microbiome health?. <i>Gut Microbes</i> , 2021, 13, 1-20.	9.8	19
15	Baseline Gut Metagenomic Functional Gene Signature Associated with Variable Weight Loss Responses following a Healthy Lifestyle Intervention in Humans. <i>MSystems</i> , 2021, 6, e0096421.	3.8	19
16	Antimicrobial Peptide against Mycobacterium Tuberculosis That Activates Autophagy Is an Effective Treatment for Tuberculosis. <i>Pharmaceutics</i> , 2020, 12, 1071.	4.5	17
17	Personalized Prediction of Proliferation Rates and Metabolic Liabilities in Cancer Biopsies. <i>Frontiers in Physiology</i> , 2016, 7, 644.	2.8	16
18	The space of enzyme regulation in HeLa cells can be inferred from its intracellular metabolome. <i>Scientific Reports</i> , 2016, 6, 28415.	3.3	15

#	ARTICLE	IF	CITATIONS
19	Lettuce (<i>Lactuca sativa</i>) productivity influenced by microbial inocula under nitrogen-limited conditions in aquaponics. <i>PLoS ONE</i> , 2021, 16, e0247534.	2.5	14
20	Progressive Shifts in the Gut Microbiome Reflect Prediabetes and Diabetes Development in a Treatment-Naive Mexican Cohort. <i>Frontiers in Endocrinology</i> , 2020, 11, 602326.	3.5	13
21	A low number of SIC1 mRNA molecules ensures a low noise level in cell cycle progression of budding yeast. <i>Molecular BioSystems</i> , 2011, 7, 2804.	2.9	9
22	Constraint-Based Reconstruction and Analyses of Metabolic Models: Open-Source Python Tools and Applications to Cancer. <i>Frontiers in Oncology</i> , 0, 12, .	2.8	6
23	What Influences DNA Replication Rate in Budding Yeast?. <i>PLoS ONE</i> , 2010, 5, e10203.	2.5	5
24	Editorial: Systems Biology and the Challenge of Deciphering the Metabolic Mechanisms Underlying Cancer. <i>Frontiers in Physiology</i> , 2017, 8, 537.	2.8	2
25	Non-responder phenotype reveals apparent microbiome-wide antibiotic tolerance in the murine gut. <i>Communications Biology</i> , 2021, 4, 316.	4.4	2
26	Experimental and Stochastic Model Analysis of the Influence of SIC1, CLN2 and CLB5 Transcriptional Noise on the Timing Regulation of G1/S Transition in <i>S. Cerevisiae</i> Cell-Cycle. <i>Biophysical Journal</i> , 2012, 102, 228a.	0.5	0
27	What Are Poop Transplants and How Do They Work?. <i>Frontiers for Young Minds</i> , 0, 9, .	0.8	0