

# 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  | Heterogeneity in statin responses explained by variation in the human gut microbiome. <i>Med</i> , 2022, 3, 388-405.e6.  | 4.4  | 21        |
| 2  | From taxonomy to metabolic output: what factors define gut microbiome health?. <i>Gut Microbes</i> , 2021, 13, 1-20.   | 9.8  | 19        |
| 3  | 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        |
| 4  | Gut microbiome pattern reflects healthy ageing and predicts survival in humans. <i>Nature Metabolism</i> , 2021, 3, 274-286.   | 11.9 | 278       |
| 5  | Non-responder phenotype reveals apparent microbiome-wide antibiotic tolerance in the murine gut. <i>Communications Biology</i> , 2021, 4, 316.   | 4.4  | 2         |
| 6  | 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        |
| 7  | 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        |
| 8  | Antimicrobial Peptide against <i>Mycobacterium Tuberculosis</i> That Activates Autophagy Is an Effective Treatment for Tuberculosis. <i>Pharmaceutics</i> , 2020, 12, 1071.                  | 4.5  | 17        |
| 9  | MEMOTE for standardized genome-scale metabolic model testing. <i>Nature Biotechnology</i> , 2020, 38, 272-276.   | 17.5 | 314       |
| 10 | MICOM: Metagenome-Scale Modeling To Infer Metabolic Interactions in the Gut Microbiota. <i>MSystems</i> , 2020, 5, .   | 3.8  | 126       |
| 11 | Progressive Shifts in the Gut Microbiome Reflect Prediabetes and Diabetes Development in a Treatment-Naïve Mexican Cohort. <i>Frontiers in Endocrinology</i> , 2020, 11, 602326.             | 3.5  | 13        |
| 12 | Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019, 37, 852-857.   | 17.5 | 11,167    |
| 13 | Use and abuse of correlation analyses in microbial ecology. <i>ISME Journal</i> , 2019, 13, 2647-2655.   | 9.8  | 193       |
| 14 | 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        |
| 15 | 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       |
| 16 | Editorial: Systems Biology and the Challenge of Deciphering the Metabolic Mechanisms Underlying Cancer. <i>Frontiers in Physiology</i> , 2017, 8, 537.                                       | 2.8  | 2         |
| 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 | Effective Design of Multifunctional Peptides by Combining Compatible Functions. PLoS Computational Biology, 2016, 12, e1004786.   | 3.2 | 36        |
| 20 | Yeast Mating and Image-Based Quantification of Spatial Pattern Formation. PLoS Computational Biology, 2014, 10, e1003690.   | 3.2 | 25        |
| 21 | Cell Penetrating Peptides and Cationic Antibacterial Peptides. Journal of Biological Chemistry, 2014, 289, 14448-14457.   | 3.4 | 49        |
| 22 | Experimental and Stochastic Model Analysis of the Influence of SIC1, CLN2 and CLB5 Transcriptional Noise on the Timing Regulation of G1/S Transition in S. Cerevisiae Cell-Cycle. Biophysical Journal, 2012, 102, 228a. | 0.5 | 0         |
| 23 | Onset of Immune Senescence Defined by Unbiased Pyrosequencing of Human Immunoglobulin mRNA Repertoires. PLoS ONE, 2012, 7, e49774.  | 2.5 | 30        |
| 24 | A low number of SIC1 mRNA molecules ensures a low noise level in cell cycle progression of budding yeast. Molecular BioSystems, 2011, 7, 2804.  | 2.9 | 9         |
| 25 | What Influences DNA Replication Rate in Budding Yeast?. PLoS ONE, 2010, 5, e10203.  | 2.5 | 5         |
| 26 | What Are Poop Transplants and How Do They Work?. Frontiers for Young Minds, 0, 9, .   | 0.8 | 0         |
| 27 | Constraint-Based Reconstruction and Analyses of Metabolic Models: Open-Source Python Tools and Applications to Cancer. Frontiers in Oncology, 0, 12, .  | 2.8 | 6         |