

Olaf Mueller

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11657281/publications.pdf>

Version: 2024-02-01

12
papers

1,362
citations

933447

10
h-index

1199594

12
g-index

14
all docs

14
docs citations

14
times ranked

2655
citing authors

#	ARTICLE	IF	CITATIONS
1	Neonatal Rhesus Macaques Have Distinct Immune Cell Transcriptional Profiles following HIV Envelope Immunization. <i>Cell Reports</i> , 2020, 30, 1553-1569.e6.	6.4	21
2	Epithelial delamination is protective during pharmaceutical-induced enteropathy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 16961-16970.	7.1	8
3	Intestinal Serum amyloid A suppresses systemic neutrophil activation and bactericidal activity in response to microbiota colonization. <i>PLoS Pathogens</i> , 2019, 15, e1007381.	4.7	54
4	High fat diet induces microbiota-dependent silencing of enteroendocrine cells. <i>ELife</i> , 2019, 8, .	6.0	73
5	Antibacterial soap use impacts skin microbial communities in rural Madagascar. <i>PLoS ONE</i> , 2018, 13, e0199899.	2.5	13
6	Conserved genomic collinearity as a source of broadly applicable, fast evolving, markers to resolve species complexes: A case study using the lichen-forming genus <i>Peltigera</i> section <i>Polydactylon</i> . <i>Molecular Phylogenetics and Evolution</i> , 2017, 117, 10-29.	2.7	30
7	Environmental influences on the skin microbiome of humans and cattle in rural Madagascar. <i>Evolution, Medicine and Public Health</i> , 2017, 2017, 144-153.	2.5	17
8	Genomic sequencing-based mutational enrichment analysis identifies motility genes in a genetically intractable gut microbe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 14127-14132.	7.1	10
9	The severity of nonalcoholic fatty liver disease is associated with gut dysbiosis and shift in the metabolic function of the gut microbiota. <i>Hepatology</i> , 2016, 63, 764-775.	7.3	1,029
10	Unisexual Reproduction of <i>Cryptococcus gattii</i> . <i>PLoS ONE</i> , 2014, 9, e111089.	2.5	20
11	High-throughput genome sequencing of lichenizing fungi to assess gene loss in the ammonium transporter/ammonia permease gene family. <i>BMC Genomics</i> , 2013, 14, 225.	2.8	41
12	Transmission of Hypervirulence Traits via Sexual Reproduction within and between Lineages of the Human Fungal Pathogen <i>Cryptococcus gattii</i> . <i>PLoS Genetics</i> , 2013, 9, e1003771.	3.5	45