

Amine Ghozlane

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1278672/publications.pdf>

Version: 2024-02-01

19
papers

1,220
citations

759233

12
h-index

996975

15
g-index

22
all docs

22
docs citations

22
times ranked

2305
citing authors

#	ARTICLE	IF	CITATIONS
1	Wide diversity of methane and short-chain alkane metabolisms in uncultured archaea. <i>Nature Microbiology</i> , 2019, 4, 603-613.	13.3	187
2	Carryover effects of larval exposure to different environmental bacteria drive adult trait variation in a mosquito vector. <i>Science Advances</i> , 2017, 3, e1700585.	10.3	172
3	Bacteriocin from epidemic <i>Listeria</i> strains alters the host intestinal microbiota to favor infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 5706-5711.	7.1	166
4	Characteristics of Faecal Microbiota in Paediatric Crohn's Disease and Their Dynamic Changes During Infliximab Therapy. <i>Journal of Crohn's and Colitis</i> , 2018, 12, 337-346.	1.3	129
5	Prediction of the intestinal resistome by a three-dimensional structure-based method. <i>Nature Microbiology</i> , 2019, 4, 112-123.	13.3	129
6	Distinct systemic and mucosal immune responses during acute SARS-CoV-2 infection. <i>Nature Immunology</i> , 2021, 22, 1428-1439.	14.5	110
7	Protein flexibility in the light of structural alphabets. <i>Frontiers in Molecular Biosciences</i> , 2015, 2, 20.	3.5	71
8	Impact of the gut microbiota on the m6A epitranscriptome of mouse cecum and liver. <i>Nature Communications</i> , 2020, 11, 1344.	12.8	59
9	Mining zebrafish microbiota reveals key community-level resistance against fish pathogen infection. <i>ISME Journal</i> , 2021, 15, 702-719.	9.8	49
10	Combined bacterial and fungal intestinal microbiota analyses: Impact of storage conditions and DNA extraction protocols. <i>PLoS ONE</i> , 2018, 13, e0201174.	2.5	41
11	SHAMAN: a user-friendly website for metataxonomic analysis from raw reads to statistical analysis. <i>BMC Bioinformatics</i> , 2020, 21, 345.	2.6	41
12	Analysis of protein chameleon sequence characteristics. <i>Bioinformatics</i> , 2009, 3, 367-369.	0.5	22
13	Leucine-Rich Immune Factor APL1 Is Associated With Specific Modulation of Enteric Microbiome Taxa in the Asian Malaria Mosquito <i>Anopheles stephensi</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 306.	3.5	20
14	Exome-wide association study reveals largely distinct gene sets underlying specific resistance to dengue virus types 1 and 3 in <i>Aedes aegypti</i> . <i>PLoS Genetics</i> , 2020, 16, e1008794.	3.5	13
15	Capturing the most wanted taxa through cross-sample correlations. <i>ISME Journal</i> , 2016, 10, 2459-2467.	9.8	9
16	Title is missing!. , 2020, 16, e1008794.		0
17	Title is missing!. , 2020, 16, e1008794.		0
18	Title is missing!. , 2020, 16, e1008794.		0

#	ARTICLE	IF	CITATIONS
19	Title is missing!. , 2020, 16, e1008794.		0