

# Clara Amid

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/734279/publications.pdf>

Version: 2024-02-01

16  
papers

1,819  
citations

840776

11  
h-index

940533

16  
g-index

17  
all docs

17  
docs citations

17  
times ranked

4114  
citing authors

#	ARTICLE	IF	CITATIONS
1	Global monitoring of antimicrobial resistance based on metagenomics analyses of urban sewage. <i>Nature Communications</i> , 2019, 10, 1124.	12.8	612
2	Minimum Information about an Uncultivated Virus Genome (MIUViG). <i>Nature Biotechnology</i> , 2019, 37, 29-37.	17.5	414
3	EBI Metagenomics in 2017: enriching the analysis of microbial communities, from sequence reads to assemblies. <i>Nucleic Acids Research</i> , 2018, 46, D726-D735.	14.5	175
4	Viral to metazoan marine plankton nucleotide sequences from the Tara Oceans expedition. <i>Scientific Data</i> , 2017, 4, 170093.	5.3	147
5	The European Nucleotide Archive in 2018. <i>Nucleic Acids Research</i> , 2019, 47, D84-D88.	14.5	103
6	The European Nucleotide Archive in 2019. <i>Nucleic Acids Research</i> , 2020, 48, D70-D76.	14.5	95
7	The European Nucleotide Archive in 2017. <i>Nucleic Acids Research</i> , 2018, 46, D36-D40.	14.5	79
8	European Nucleotide Archive in 2016. <i>Nucleic Acids Research</i> , 2017, 45, D32-D36.	14.5	68
9	Content discovery and retrieval services at the European Nucleotide Archive. <i>Nucleic Acids Research</i> , 2015, 43, D23-D29.	14.5	36
10	Biocuration of functional annotation at the European nucleotide archive. <i>Nucleic Acids Research</i> , 2016, 44, D58-D66.	14.5	28
11	The COMPARE Data Hubs. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	3.0	28
12	Value, but high costs in post-deposition data curation. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, bav126.	3.0	12
13	Identifying causative mechanisms linking early-life stress to psycho-cardio-metabolic multi-morbidity: The EarlyCause project. <i>PLoS ONE</i> , 2021, 16, e0245475.	2.5	9
14	Accelerating surveillance and research of antimicrobial resistance – an online repository for sharing of antimicrobial susceptibility data associated with whole-genome sequences. <i>Microbial Genomics</i> , 2020, 6, .	2.0	5
15	Metagenomics-Based Proficiency Test of Smoked Salmon Spiked with a Mock Community. <i>Microorganisms</i> , 2020, 8, 1861.	3.6	4
16	Comparison of sequencing methods and data processing pipelines for whole genome sequencing and minority single nucleotide variant (mSNV) analysis during an influenza A/H5N8 outbreak. <i>PLoS ONE</i> , 2020, 15, e0229326.	2.5	1