## Julia Helmecke

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5869950/publications.pdf

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		1478505	1474206	
10	565	6	9	
papers	citations	h-index	g-index	
10	10	10	680	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	BRENDA, the ELIXIR core data resource in 2021: new developments and updates. Nucleic Acids Research, 2021, 49, D498-D508.	14.5	347
2	Enhancement of IFN $\hat{I}^3$ Production by Distinct Commensals Ameliorates Salmonella-Induced Disease. Cell Host and Microbe, 2017, 21, 682-694.e5.	11.0	91
3	Bac <i>Dive</i> in 2022: the knowledge base for standardized bacterial and archaeal data. Nucleic Acids Research, 2022, 50, D741-D746.	14.5	71
4	Convergent Loss of ABC Transporter Genes From Clostridioides difficile Genomes Is Associated With Impaired Tyrosine Uptake and p-Cresol Production. Frontiers in Microbiology, 2018, 9, 901.	3.5	30
5	MetaboMAPS: Pathway sharing and multi-omics data visualization in metabolic context. F1000Research, 2020, 9, 288.	1.6	8
6	MetaboMAPS: Pathway sharing and multi-omics data visualization in metabolic context. F1000Research, 2020, 9, 288.	1.6	8
7	The Impact of Pyroglutamate:Sulfolobus acidocaldariusHas a Growth Advantage overSaccharolobus solfataricusin Glutamate-Containing Media. Archaea, 2019, 2019, 1-9.	2.3	3
8	DSMZCellDive: Diving into high-throughput cell line data. F1000Research, 0, 11, 420.	1.6	3
9	Utilization of Phenol as Carbon Source by the Thermoacidophilic Archaeon Saccharolobus solfataricus P2 Is Limited by Oxygen Supply and the Cellular Stress Response. Frontiers in Microbiology, 2020, $11,587032$ .	3.5	2
10	The Metano Modeling Toolbox MMTB: An Intuitive, Web-Based Toolbox Introduced by Two Use Cases. Metabolites, 2021, 11, 113.	2.9	2