

# Szymon T Calus

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2862954/publications.pdf>

Version: 2024-02-01

10  
papers

3,500  
citations

1051969

10  
h-index

1526636

10  
g-index

15  
all docs

15  
docs citations

15  
times ranked

8172  
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential prevalence and host-association of antimicrobial resistance traits in disinfected and non-disinfected drinking water systems. <i>Science of the Total Environment</i> , 2020, 749, 141451.	3.9	22
2	Disinfection exhibits systematic impacts on the drinking water microbiome. <i>Microbiome</i> , 2020, 8, 42.	4.9	48
3	NanoAmpli-Seq: a workflow for amplicon sequencing for mixed microbial communities on the nanopore sequencing platform. <i>GigaScience</i> , 2018, 7, .	3.3	85
4	The distinct features of microbial "dysbiosis"™ of Crohn's™ disease do not occur to the same extent in their unaffected, genetically-linked kindred. <i>PLoS ONE</i> , 2017, 12, e0172605.	1.1	33
5	Staphylococcal species heterogeneity in the nasal microbiome following antibiotic prophylaxis revealed by tuf gene deep sequencing. <i>Microbiome</i> , 2016, 4, 63.	4.9	15
6	The effect of DNA extraction methodology on gut microbiota research applications. <i>BMC Research Notes</i> , 2016, 9, 365.	0.6	66
7	Mycobacterial Dihydrofolate Reductase Inhibitors Identified Using Chemogenomic Methods and In Vitro Validation. <i>PLoS ONE</i> , 2015, 10, e0121492.	1.1	40
8	Extensive Modulation of the Fecal Metagenome in Children With Crohn's™ Disease During Exclusive Enteral Nutrition. <i>American Journal of Gastroenterology</i> , 2015, 110, 1718-1729.	0.2	229
9	Rapid draft sequencing and real-time nanopore sequencing in a hospital outbreak of Salmonella. <i>Genome Biology</i> , 2015, 16, 114.	3.8	271
10	Reagent and laboratory contamination can critically impact sequence-based microbiome analyses. <i>BMC Biology</i> , 2014, 12, 87.	1.7	2,677