

# Jacob T Nearing

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3477997/publications.pdf>

Version: 2024-02-01

12  
papers

904  
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1162367

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1199166

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15  
docs citations

15  
times ranked

961  
citing authors

#	ARTICLE	IF	CITATIONS
1	Microbiome differential abundance methods produce different results across 38 datasets. Nature Communications, 2022, 13, 342.	5.8	286
2	Plant-Based Diets and Cancer Risk: What is the Evidence?. Current Nutrition Reports, 2022, 11, 354-369.	2.1	13
3	Investigating the gut microbial community and genes in children with differing levels of change in serum asparaginase activity during pegaspargase treatment for acute lymphoblastic leukemia. Leukemia and Lymphoma, 2021, 62, 927-936.	0.6	4
4	Harnessing the plant microbiome to promote the growth of agricultural crops. Microbiological Research, 2021, 245, 126690.	2.5	84
5	An <i>in vivo</i> genome-wide shRNA screen identifies BCL6 as a targetable biomarker of paclitaxel resistance in breast cancer. Molecular Oncology, 2021, 15, 2046-2064.	2.1	5
6	Identifying biases and their potential solutions in human microbiome studies. Microbiome, 2021, 9, 113.	4.9	82
7	Investigation of the impact of commonly used medications on the oral microbiome of individuals living without major chronic conditions. PLoS ONE, 2021, 16, e0261032.	1.1	8
8	Assessing the Variation within the Oral Microbiome of Healthy Adults. MSphere, 2020, 5, .	1.3	49
9	Investigating Gut Microbial Taxa and Asparaginase Related Genes in Children Showing Different Direction of Change in Serum Asparaginase Activity Levels during Pegasparginase Treatment for Acute Lymphoblastic Leukemia. Blood, 2020, 136, 40-41.	0.6	0
10	Metagenomic Functional Shifts to Plant Induced Environmental Changes. Frontiers in Microbiology, 2019, 10, 1682.	1.5	28
11	Infectious Complications Are Associated With Alterations in the Gut Microbiome in Pediatric Patients With Acute Lymphoblastic Leukemia. Frontiers in Cellular and Infection Microbiology, 2019, 9, 28.	1.8	48
12	Denoising the Denoisers: an independent evaluation of microbiome sequence error-correction approaches. PeerJ, 2018, 6, e5364.	0.9	278