

# Mohammadali Khan Mirzaei

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/8522453/mohammadali-khan-mirzaei-publications-by-year.pdf>

**Version:** 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19  
papers

744  
citations

9  
h-index

26  
g-index

26  
ext. papers

1,056  
ext. citations

11  
avg, IF

4.88  
L-index

#	Paper	IF	Citations
19	Emerging technologies in the study of the virome. <i>Current Opinion in Virology</i> , <b>2022</b> , 54, 101231	7.5	1
18	New technologies for developing phage-based tools to manipulate the human microbiome. <i>Trends in Microbiology</i> , <b>2021</b> ,	12.4	5
17	Gut Phage Database: phage mining in the cave of wonders. <i>Signal Transduction and Targeted Therapy</i> , <b>2021</b> , 6, 193	21	2
16	Sewage and sewage-contaminated environments are the most prominent sources to isolate phages against <i>Pseudomonas aeruginosa</i> . <i>BMC Microbiology</i> , <b>2021</b> , 21, 132	4.5	3
15	Challenges of Studying the Human Virome - Relevant Emerging Technologies. <i>Trends in Microbiology</i> , <b>2021</b> , 29, 171-181	12.4	22
14	Sustainable Microbiome: a symphony orchestrated by synthetic phages. <i>Microbial Biotechnology</i> , <b>2021</b> , 14, 45-50	6.3	1
13	Improving the Inhibitory Effect of Phages against Isolated from a Burn Patient Using a Combination of Phages and Antibiotics. <i>Viruses</i> , <b>2021</b> , 13,	6.2	5
12	Differences in Gut Virome Related to Barrett Esophagus and Esophageal Adenocarcinoma. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	2
11	Bacteriophages Isolated from Stunted Children Can Regulate Gut Bacterial Communities in an Age-Specific Manner. <i>Cell Host and Microbe</i> , <b>2020</b> , 27, 199-212.e5	23.4	48
10	Global phylogeography and ancient evolution of the widespread human gut virus crAssphage. <i>Nature Microbiology</i> , <b>2019</b> , 4, 1727-1736	26.6	100
9	The prevalence of CMY-2, OXA-48 and KPC-2 genes in clinical isolates of <i>Klebsiella</i> spp. <i>Cellular and Molecular Biology</i> , <b>2018</b> , 64, 40-44	1.1	8
8	Phage Trophs in the human gut: interactions between host, bacteria and phages. <i>Nature Reviews Microbiology</i> , <b>2017</b> , 15, 397-408	22.2	184
7	The Mammalian Gut as a Matchmaker. <i>Cell Host and Microbe</i> , <b>2017</b> , 22, 726-727	23.4	2
6	Morphologically Distinct <i>Escherichia coli</i> Bacteriophages Differ in Their Efficacy and Ability to Stimulate Cytokine Release In Vitro. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 437	5.7	21
5	Adapting Drug Approval Pathways for Bacteriophage-Based Therapeutics. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 1209	5.7	94
4	Response: Commentary: Morphologically Distinct Bacteriophages Differ in Their Efficacy and Ability to Stimulate Cytokine Release. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 1974	5.7	1
3	Isolation of phages for phage therapy: a comparison of spot tests and efficiency of plating analyses for determination of host range and efficacy. <i>PLoS ONE</i> , <b>2015</b> , 10, e0118557	3.7	154

- |   |  |     |    |
|---|--|-----|----|
| 2 | Genomic, proteomic, morphological, and phylogenetic analyses of vB_EcoP_SU10, a podoviridae phage with C3 morphology. <i>PLoS ONE</i> , <b>2014</b> , 9, e116294 | 3-7 | 18 |
| 1 | Global phylogeography and ancient evolution of the widespread human gut virus crAssphage   |     | 5  |