

# Domonkos Svab

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/2104828/domonkos-svab-publications-by-year.pdf>

**Version:** 2024-04-26

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.

18  
papers

128  
citations

7  
h-index

11  
g-index

20  
ext. papers

178  
ext. citations

3.8  
avg. IF

2.55  
L-index

#	Paper	IF	Citations
18	Characterisation of new anti-O157 bacteriophages of bovine origin representing three genera.. <i>Archives of Microbiology</i> , <b>2022</b> , 204, 231	3	
17	Complete Genome Sequences of Novel Bovine T4, rv5-Like, and Dhillonviruses Effective against <i>Escherichia coli</i> O157. <i>Microbiology Resource Announcements</i> , <b>2021</b> , 10,	1.3	1
16	Novel prophage-like sequences in <i>Mycoplasma anserisalpingtonis</i> . <i>Infection, Genetics and Evolution</i> , <b>2021</b> , 92, 104886	4.5	
15	Genome Analysis of a Historical Serotype 1 Strain Carrying a Conserved Stx Prophage Region. <i>Frontiers in Microbiology</i> , <b>2020</b> , 11, 614793	5.7	0
14	R18C is a new viable P2-like bacteriophage of rabbit origin infecting <i>Citrobacter rodentium</i> and <i>Shigella sonnei</i> strains. <i>Archives of Virology</i> , <b>2019</b> , 164, 3157-3160	2.6	2
13	Complete genome sequence of C130_2, a novel myovirus infecting pathogenic <i>Escherichia coli</i> and <i>Shigella</i> strains. <i>Archives of Virology</i> , <b>2019</b> , 164, 321-324	2.6	5
12	Identification and characterization of new broad host-range rv5-like coliphages C203 and P206 directed against enterobacteria. <i>Infection, Genetics and Evolution</i> , <b>2018</b> , 64, 254-261	4.5	7
11	Identification and Characterization of T5-Like Bacteriophages Representing Two Novel Subgroups from Food Products. <i>Frontiers in Microbiology</i> , <b>2018</b> , 9, 202	5.7	24
10	Comparative Genomic and Phylogenetic Analysis of a Shiga Toxin Producing (STSS) Strain. <i>Frontiers in Cellular and Infection Microbiology</i> , <b>2017</b> , 7, 229	5.9	13
9	Cytolethal distending toxin producing <i>Escherichia coli</i> O157:H43 strain T22 represents a novel evolutionary lineage within the O157 serogroup. <i>Infection, Genetics and Evolution</i> , <b>2016</b> , 46, 110-117	4.5	1
8	Comparative analysis of the Shiga toxin converting bacteriophage first detected in <i>Shigella sonnei</i> . <i>Infection, Genetics and Evolution</i> , <b>2016</b> , 37, 150-7	4.5	23
7	A novel transducible chimeric phage from <i>Escherichia coli</i> O157:H7 Sakai strain encoding Stx1 production. <i>Infection, Genetics and Evolution</i> , <b>2015</b> , 29, 42-7	4.5	7
6	Cytolethal distending toxin A, B and C subunit proteins are necessary for the genotoxic effect of <i>Escherichia coli</i> CDT-V. <i>Acta Veterinaria Hungarica</i> , <b>2015</b> , 63, 1-10	1	7
5	Draft Genome Sequence of an <i>Escherichia coli</i> O157:H43 Strain Isolated from Cattle. <i>Genome Announcements</i> , <b>2013</b> , 1,		6
4	The long polar fimbriae operon and its flanking regions in bovine <i>Escherichia coli</i> O157:H43 and STEC O136:H12 strains. <i>Pathogens and Disease</i> , <b>2013</b> , 68, 1-7	4.2	2
3	Sequence variability of P2-like prophage genomes carrying the cytolethal distending toxin V operon in <i>Escherichia coli</i> O157. <i>Applied and Environmental Microbiology</i> , <b>2013</b> , 79, 4958-64	4.8	11
2	Allelic types of long polar fimbriae in bovine and human <i>Escherichia coli</i> O157 strains. <i>Acta Veterinaria Hungarica</i> , <b>2012</b> , 60, 1-15	1	4

- 1 Preferential ligation during TA-cloning of multitemplate PCR products--a factor causing bias in microbial community structure analysis. *Journal of Microbiological Methods*, **2011**, 85, 131-6 2.8 15