

# Domonkos Svab

## List of Publications by Citations

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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	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
17	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
16	Preferential ligation during TA-cloning of multitemplate PCR products--a factor causing bias in microbial community structure analysis. <i>Journal of Microbiological Methods</i> , <b>2011</b> , 85, 131-6	2.8	15
15	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
14	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
13	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
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	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
10	Draft Genome Sequence of an <i>Escherichia coli</i> O157:H43 Strain Isolated from Cattle. <i>Genome Announcements</i> , <b>2013</b> , 1,		6
9	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
8	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
7	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
6	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
5	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
4	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
3	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
2	Novel prophage-like sequences in <i>Mycoplasma anserisalpingitidis</i> . <i>Infection, Genetics and Evolution</i> , <b>2021</b> , 92, 104886	4.5	

- 1 Characterisation of new anti-O157 bacteriophages of bovine origin representing three genera.. *Archives of Microbiology*, **2022**, 204, 231 3