

Domonkos Sṽjb

List of Publications by Year in descending order

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papers

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citations

1162889

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20
all docs

20
docs citations

20
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citing authors

#	ARTICLE	IF	CITATIONS
1	Characterisation of new anti-O157 bacteriophages of bovine origin representing three genera. Archives of Microbiology, 2022, 204, 231.	1.0	1
2	Multiplex polymerase chain reaction typing scheme based on Escherichia coli O157:H7 Sakai prophage (Sp)-associated genes. International Journal of Infectious Diseases, 2022, 120, 68-76.	1.5	2
3	Complete Genome Sequences of Novel Bovine T4, rv5-Like, and Dhillonviruses Effective against Escherichia coli O157. Microbiology Resource Announcements, 2021, 10, .	0.3	1
4	Novel prophage-like sequences in Mycoplasma anserisalpingitidis. Infection, Genetics and Evolution, 2021, 92, 104886.	1.0	1
5	Genome Analysis of a Historical Shigella dysenteriae Serotype 1 Strain Carrying a Conserved Stx Prophage Region. Frontiers in Microbiology, 2020, 11, 614793.	1.5	1
6	R18C is a new viable P2-like bacteriophage of rabbit origin infecting Citrobacter rodentium and Shigella sonnei strains. Archives of Virology, 2019, 164, 3157-3160.	0.9	4
7	Complete genome sequence of C130_2, a novel myovirus infecting pathogenic Escherichia coli and Shigella strains. Archives of Virology, 2019, 164, 321-324.	0.9	5
8	Identification and characterization of new broad host-range rv5-like coliphages C203 and P206 directed against enterobacteria. Infection, Genetics and Evolution, 2018, 64, 254-261.	1.0	14
9	Identification and Characterization of T5-Like Bacteriophages Representing Two Novel Subgroups from Food Products. Frontiers in Microbiology, 2018, 9, 202.	1.5	39
10	Comparative Genomic and Phylogenetic Analysis of a Shiga Toxin Producing Shigella sonnei (STSS) Strain. Frontiers in Cellular and Infection Microbiology, 2017, 7, 229.	1.8	18
11	Cytolethal distending toxin producing Escherichia coli O157:H43 strain T22 represents a novel evolutionary lineage within the O157 serogroup. Infection, Genetics and Evolution, 2016, 46, 110-117.	1.0	3
12	Comparative analysis of the Shiga toxin converting bacteriophage first detected in Shigella sonnei. Infection, Genetics and Evolution, 2016, 37, 150-157.	1.0	31
13	Cytolethal distending toxin A, B and C subunit proteins are necessary for the genotoxic effect of Escherichia coli CDT-V. Acta Veterinaria Hungarica, 2015, 63, 1-10.	0.2	8
14	A novel transducible chimeric phage from Escherichia coli O157:H7 Sakai strain encoding Stx1 production. Infection, Genetics and Evolution, 2015, 29, 42-47.	1.0	12
15	Draft Genome Sequence of an Escherichia coli O157:H43 Strain Isolated from Cattle. Genome Announcements, 2013, 1, .	0.8	6
16	The long polar fimbriae operon and its flanking regions in bovine Escherichia coli O157:H43 and STEC O136:H12 strains. Pathogens and Disease, 2013, 68, 1-7.	0.8	2
17	Sequence Variability of P2-Like Prophage Genomes Carrying the Cytolethal Distending Toxin V Operon in Escherichia coli O157. Applied and Environmental Microbiology, 2013, 79, 4958-4964.	1.4	20
18	Allelic types of long polar fimbriae in bovine and human Escherichia coli O157 strains. Acta Veterinaria Hungarica, 2012, 60, 1-15.	0.2	4

#	ARTICLE	IF	CITATIONS
19	Preferential ligation during TA-cloning of multitemplate PCR products – A factor causing bias in microbial community structure analysis. <i>Journal of Microbiological Methods</i> , 2011, 85, 131-136.	0.7	16
20	Genomic Diversity, Virulence Gene, and Prophage Arrays of Bovine and Human Shiga Toxigenic and Enteropathogenic <i>Escherichia coli</i> Strains Isolated in Hungary. <i>Frontiers in Microbiology</i> , 0, 13, .	1.5	1