

Lone BrÃndsted

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

941
citations

516710

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642732

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docs citations

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times ranked

922
citing authors

#	ARTICLE	IF	CITATIONS
1	Bacteriophage F336 Recognizes the Capsular Phosphoramidate Modification of <i>Campylobacter jejuni</i> NCTC11168. <i>Journal of Bacteriology</i> , 2011, 193, 6742-6749.	2.2	115
2	Phase Variable Expression of Capsular Polysaccharide Modifications Allows <i>Campylobacter jejuni</i> to Avoid Bacteriophage Infection in Chickens. <i>Frontiers in Cellular and Infection Microbiology</i> , 2012, 2, 11.	3.9	87
3	Energy Taxis Drives <i>Campylobacter jejuni</i> toward the Most Favorable Conditions for Growth. <i>Applied and Environmental Microbiology</i> , 2009, 75, 5308-5314.	3.1	84
4	Structure and Function of the Branched Receptor-Binding Complex of Bacteriophage CBA120. <i>Journal of Molecular Biology</i> , 2019, 431, 3718-3739.	4.2	67
5	<i>Campylobacter jejuni</i> Motility Is Required for Infection of the Flagellotropic Bacteriophage F341. <i>Applied and Environmental Microbiology</i> , 2014, 80, 7096-7106.	3.1	60
6	Primary Isolation Strain Determines Both Phage Type and Receptors Recognised by <i>Campylobacter jejuni</i> Bacteriophages. <i>PLoS ONE</i> , 2015, 10, e0116287.	2.5	58
7	Exploiting phage receptor binding proteins to enable endolysins to kill Gram-negative bacteria. <i>Scientific Reports</i> , 2020, 10, 12087.	3.3	57
8	Contribution of Conserved ATP-Dependent Proteases of <i>Campylobacter jejuni</i> to Stress Tolerance and Virulence. <i>Applied and Environmental Microbiology</i> , 2007, 73, 7803-7813.	3.1	53
9	The genera of bacteriophages and their receptors are the major determinants of host range. <i>Environmental Microbiology</i> , 2019, 21, 2095-2111.	3.8	45
10	Subtypes of tail spike proteins predicts the host range of Ackermannviridae phages. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 4854-4867.	4.1	36
11	Significance of phage-host interactions for biocontrol of <i>Campylobacter jejuni</i> in food. <i>Food Control</i> , 2017, 73, 1169-1175.	5.5	35
12	Phage exposure causes dynamic shifts in the expression states of specific phase-variable genes of <i>Campylobacter jejuni</i> . <i>Microbiology (United Kingdom)</i> , 2017, 163, 911-919.	1.8	35
13	Methods for Isolation, Purification, and Propagation of Bacteriophages of <i>Campylobacter jejuni</i> . <i>Methods in Molecular Biology</i> , 2017, 1512, 19-28.	0.9	33
14	Phase Variable Expression of a Single Phage Receptor in <i>Campylobacter jejuni</i> NCTC12662 Influences Sensitivity Toward Several Diverse CPS-Dependent Phages. <i>Frontiers in Microbiology</i> , 2018, 9, 82.	3.5	31
15	Natural Transformation of <i>Campylobacter jejuni</i> Occurs Beyond Limits of Growth. <i>PLoS ONE</i> , 2012, 7, e45467.	2.5	28
16	Developing Innolysins Against <i>Campylobacter jejuni</i> Using a Novel Prophage Receptor-Binding Protein. <i>Frontiers in Microbiology</i> , 2021, 12, 619028.	3.5	24
17	Phage S144, a New Polyvalent Phage Infecting <i>Salmonella</i> spp. and <i>Cronobacter sakazakii</i> . <i>International Journal of Molecular Sciences</i> , 2020, 21, 5196.	4.1	22
18	Looking into the future of phage-based control of zoonotic pathogens in food and animal production. <i>Current Opinion in Biotechnology</i> , 2021, 68, 96-103.	6.6	20

#	ARTICLE	IF	CITATIONS
19	Campylobacter phages use hypermutable polyG tracts to create phenotypic diversity and evade bacterial resistance. <i>Cell Reports</i> , 2021, 35, 109214.	6.4	15
20	Methods for Initial Characterization of Campylobacter jejuni Bacteriophages. <i>Methods in Molecular Biology</i> , 2017, 1512, 91-105.	0.9	8
21	Identification of Novel Phage Resistance Mechanisms in Campylobacter jejuni by Comparative Genomics. <i>Frontiers in Microbiology</i> , 2021, 12, 780559.	3.5	7
22	Two Distinct Modes of Lysis Regulation in Campylobacter Fletchervirus and Firehammervirus Phages. <i>Viruses</i> , 2020, 12, 1247.	3.3	6
23	Bacteriophages for Biological Control of Foodborne Pathogens. , 0, , 755-786.		4
24	Whole-Genome Sequence of the Bacteriophage-Sensitive Strain Campylobacter jejuni NCTC12662. <i>Genome Announcements</i> , 2017, 5, .	0.8	3