# Sylvain Moineau

### List of Publications by Citations

Source: https://exaly.com/author-pdf/3751674/sylvain-moineau-publications-by-citations.pdf

Version: 2024-04-19

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.

20,923 141 243 57 h-index g-index citations papers 6.89 263 25,351 7.3 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
243	CRISPR provides acquired resistance against viruses in prokaryotes. <i>Science</i> , <b>2007</b> , 315, 1709-12	33.3	3735
242	Evolution and classification of the CRISPR-Cas systems. <i>Nature Reviews Microbiology</i> , <b>2011</b> , 9, 467-77	22.2	1604
241	The CRISPR/Cas bacterial immune system cleaves bacteriophage and plasmid DNA. <i>Nature</i> , <b>2010</b> , 468, 67-71	50.4	1462
240	An updated evolutionary classification of CRISPR-Cas systems. <i>Nature Reviews Microbiology</i> , <b>2015</b> , 13, 722-36	22.2	1434
239	Bacteriophage resistance mechanisms. <i>Nature Reviews Microbiology</i> , <b>2010</b> , 8, 317-27	22.2	1382
238	Phage response to CRISPR-encoded resistance in Streptococcus thermophilus. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 1390-400	3.5	897
237	Diversity, activity, and evolution of CRISPR loci in Streptococcus thermophilus. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 1401-12	3.5	586
236	Evolutionary classification of CRISPR-Cas systems: a burst of class 2 and derived variants. <i>Nature Reviews Microbiology</i> , <b>2020</b> , 18, 67-83	22.2	545
235	Revenge of the phages: defeating bacterial defences. <i>Nature Reviews Microbiology</i> , <b>2013</b> , 11, 675-87	22.2	421
234	CRISPR/Cas system and its role in phage-bacteria interactions. <i>Annual Review of Microbiology</i> , <b>2010</b> , 64, 475-93	17.5	405
233	Methods for sampling of airborne viruses. <i>Microbiology and Molecular Biology Reviews</i> , <b>2008</b> , 72, 413-44	13.2	255
232	Biodiversity and classification of lactococcal phages. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 4338-46	4.8	200
231	Biochemistry, genetics, and applications of exopolysaccharide production in Streptococcus thermophilus: a review. <i>Journal of Dairy Science</i> , <b>2003</b> , 86, 407-23	4	187
230	Phage diversity, genomics and phylogeny. <i>Nature Reviews Microbiology</i> , <b>2020</b> , 18, 125-138	22.2	160
229	Bacteriophages of lactic acid bacteria and their impact on milk fermentations. <i>Microbial Cell Factories</i> , <b>2011</b> , 10 Suppl 1, S20	6.4	153
228	Identification of a genetic determinant responsible for host specificity in Streptococcus thermophilus bacteriophages. <i>Molecular Microbiology</i> , <b>2001</b> , 41, 325-36	4.1	151
227	Evolution of a Lytic Bacteriophage via DNA Acquisition from the Lactococcus lactis Chromosome. <i>Applied and Environmental Microbiology</i> , <b>1994</b> , 60, 1832-41	4.8	145

#### (1996-2013)

226	CRISPR-Cas and restriction-modification systems are compatible and increase phage resistance. <i>Nature Communications</i> , <b>2013</b> , 4, 2087	17.4	137
225	Bacteriophages and dairy fermentations. <i>Bacteriophage</i> , <b>2012</b> , 2, 149-158		136
224	The population and evolutionary dynamics of phage and bacteria with CRISPR-mediated immunity. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003312	6	126
223	Structure of lactococcal phage p2 baseplate and its mechanism of activation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2010</b> , 107, 6852-7	11.5	124
222	An anti-CRISPR from a virulent streptococcal phage inhibits Streptococcus pyogenes Cas9. <i>Nature Microbiology</i> , <b>2017</b> , 2, 1374-1380	26.6	117
221	Lactococcal bacteriophage p2 receptor-binding protein structure suggests a common ancestor gene with bacterial and mammalian viruses. <i>Nature Structural and Molecular Biology</i> , <b>2006</b> , 13, 85-9	17.6	109
220	Widespread anti-CRISPR proteins in virulent bacteriophages inhibit a range of Cas9 proteins. <i>Nature Communications</i> , <b>2018</b> , 9, 2919	17.4	108
219	Comparison of five bacteriophages as models for viral aerosol studies. <i>Applied and Environmental Microbiology</i> , <b>2014</b> , 80, 4242-50	4.8	106
218	Multiplex PCR for detection and identification of lactococcal bacteriophages. <i>Applied and Environmental Microbiology</i> , <b>2000</b> , 66, 987-94	4.8	104
217	CRISPR-Cas: an efficient tool for genome engineering of virulent bacteriophages. <i>Nucleic Acids Research</i> , <b>2014</b> , 42, 9504-13	20.1	98
216	Characterization of lactococcal bacteriophages from Quebec cheese plants. <i>Canadian Journal of Microbiology</i> , <b>1992</b> , 38, 875-882	3.2	97
215	Receptor-binding protein of Lactococcus lactis phages: identification and characterization of the saccharide receptor-binding site. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 2400-10	3.5	95
214	Peptidoglycan hydrolase fusions maintain their parental specificities. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 2988-96	4.8	94
213	Complete genomic sequence of the lytic bacteriophage DT1 of Streptococcus thermophilus. <i>Virology</i> , <b>1999</b> , 255, 63-76	3.6	94
212	Adaptation in bacterial CRISPR-Cas immunity can be driven by defective phages. <i>Nature Communications</i> , <b>2014</b> , 5, 4399	17.4	93
211	Modular structure of the receptor binding proteins of Lactococcus lactis phages. The RBP structure of the temperate phage TP901-1. <i>Journal of Biological Chemistry</i> , <b>2006</b> , 281, 14256-62	5.4	93
210	Homologous recombination between a lactococcal bacteriophage and the chromosome of its host strain. <i>Virology</i> , <b>2000</b> , 270, 65-75	3.6	91
209	Isolation and Characterization of Lactococcal Bacteriophages from Cultured Buttermilk Plants in the United States. <i>Journal of Dairy Science</i> , <b>1996</b> , 79, 2104-2111	4	88

208	Abortive infection mechanisms and prophage sequences significantly influence the genetic makeup of emerging lytic lactococcal phages. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 1482-7	3.5	87
207	Effect of exopolysaccharides on phage-host interactions in Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2002</b> , 68, 4364-9	4.8	85
206	Cleavage of phage DNA by the Streptococcus thermophilus CRISPR3-Cas system. <i>PLoS ONE</i> , <b>2012</b> , 7, e40913	3.7	82
205	Applications of phage resistance in lactic acid bacteria <b>1999</b> , 76, 377-382		82
204	Effect of Exopolysaccharides on Phage-Host Interactions in Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2003</b> , 69, 723-723	4.8	78
203	Bacteriophages in food fermentations: new frontiers in a continuous arms race. <i>Annual Review of Food Science and Technology</i> , <b>2013</b> , 4, 347-68	14.7	73
202	Detection of airborne lactococcal bacteriophages in cheese manufacturing plants. <i>Applied and Environmental Microbiology</i> , <b>2011</b> , 77, 491-7	4.8	72
201	Genomic organization and molecular analysis of virulent bacteriophage 2972 infecting an exopolysaccharide-producing Streptococcus thermophilus strain. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 4057-68	4.8	72
200	Characterization of two polyvalent phages infecting Enterobacteriaceae. Scientific Reports, 2017, 7, 40	34 <u>9</u> 9	71
199	Complete genomic sequence of bacteriophage ul36: demonstration of phage heterogeneity within the P335 quasi-species of lactococcal phages. <i>Virology</i> , <b>2002</b> , 296, 308-20	3.6	71
198	Structure, adsorption to host, and infection mechanism of virulent lactococcal phage p2. <i>Journal of Virology</i> , <b>2013</b> , 87, 12302-12	6.6	70
197	Morphological and genetic diversity of temperate phages in Clostridium difficile. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 7358-66	4.8	70
196	Costs of CRISPR-Cas-mediated resistance in Streptococcus thermophilus. <i>Proceedings of the Royal Society B: Biological Sciences</i> , <b>2015</b> , 282, 20151270	4.4	68
195	Evolution of Lactococcus lactis phages within a cheese factory. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 5336-44	4.8	68
194	Llama antibodies against a lactococcal protein located at the tip of the phage tail prevent phage infection. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 4531-41	3.5	65
193	Characterization of mesophilic mixed starter cultures used for the manufacture of aged cheddar cheese. <i>Journal of Dairy Science</i> , <b>2000</b> , 83, 620-7	4	64
192	Sequence and comparative genomic analysis of lactococcal bacteriophages jj50, 712 and P008: evolutionary insights into the 936 phage species. <i>FEMS Microbiology Letters</i> , <b>2006</b> , 261, 253-61	2.9	61
191	Molecular characterization of a theta replication plasmid and its use for development of a two-component food-grade cloning system for Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2001</b> , 67, 1700-9	4.8	61

## (2018-2014)

190	The three major types of CRISPR-Cas systems function independently in CRISPR RNA biogenesis in Streptococcus thermophilus. <i>Molecular Microbiology</i> , <b>2014</b> , 93, 98-112	4.1	60
189	Genome Engineering of Virulent Lactococcal Phages Using CRISPR-Cas9. <i>ACS Synthetic Biology</i> , <b>2017</b> , 6, 1351-1358	5.7	58
188	AbiQ, an abortive infection mechanism from Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>1998</b> , 64, 4748-56	4.8	58
187	Differentiation of Two Abortive Mechanisms by Using Monoclonal Antibodies Directed toward Lactococcal Bacteriophage Capsid Proteins. <i>Applied and Environmental Microbiology</i> , <b>1993</b> , 59, 208-12	4.8	57
186	Bacteriophages of lactobacillus. Frontiers in Bioscience - Landmark, 2009, 14, 1661-83	2.8	56
185	Crystal structure of the receptor-binding protein head domain from Lactococcus lactis phage bIL170. <i>Journal of Virology</i> , <b>2006</b> , 80, 9331-5	6.6	56
184	Galactose and lactose genes from the galactose-positive bacterium Streptococcus salivarius and the phylogenetically related galactose-negative bacterium Streptococcus thermophilus: organization, sequence, transcription, and activity of the gal gene products. <i>Journal of Bacteriology</i> ,	3.5	56
183	<b>2002</b> , 184, 785-93 Genome sequence and global gene expression of Q54, a new phage species linking the 936 and c2 phage species of Lactococcus lactis. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 6101-14	3.5	55
182	Characterization of coliphage PR772 and evaluation of its use for virus filter performance testing. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 4864-71	4.8	55
181	Characterization of 1706, a virulent phage from Lactococcus lactis with similarities to prophages from other Firmicutes. <i>Virology</i> , <b>2008</b> , 373, 298-309	3.6	54
180	Crystal structure and function of a DARPin neutralizing inhibitor of lactococcal phage TP901-1: comparison of DARPin and camelid VHH binding mode. <i>Journal of Biological Chemistry</i> , <b>2009</b> , 284, 3071	8 <sup>5</sup> 245	51
179	Functional and structural basis for a bacteriophage homolog of human RAD52. <i>Current Biology</i> , <b>2008</b> , 18, 1142-6	6.3	51
178	Phage production and maintenance of stocks, including expected stock lifetimes. <i>Methods in Molecular Biology</i> , <b>2009</b> , 501, 203-19	1.4	50
177	Characterization of a new virulent phage (MLC-A) of Lactobacillus paracasei. <i>Journal of Dairy Science</i> , <b>2006</b> , 89, 2414-23	4	50
176	A Unified Resource for Tracking Anti-CRISPR Names. CRISPR Journal, 2018, 1, 304-305	2.5	50
175	Prophages of the genus Bifidobacterium as modulating agents of the infant gut microbiota. <i>Environmental Microbiology</i> , <b>2016</b> , 18, 2196-213	5.2	49
174	The cell lysis activity of the Streptococcus agalactiae bacteriophage B30 endolysin relies on the cysteine, histidine-dependent amidohydrolase/peptidase domain. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 5108-12	4.8	49
173	Evolutionary emergence of infectious diseases in heterogeneous host populations. <i>PLoS Biology</i> , <b>2018</b> , 16, e2006738	9.7	49

172	Streptococcus thermophilus bacteriophages. International Dairy Journal, 2010, 20, 657-664	3.5	48
171	Restriction/Modification systems and restriction endonucleases are more effective on lactococcal bacteriophages that have emerged recently in the dairy industry. <i>Applied and Environmental Microbiology</i> , <b>1993</b> , 59, 197-202	4.8	48
170	Molecular insights on the recognition of a Lactococcus lactis cell wall pellicle by the phage 1358 receptor binding protein. <i>Journal of Virology</i> , <b>2014</b> , 88, 7005-15	6.6	47
169	Morphology, genome sequence, and structural proteome of type phage P335 from Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2008</b> , 74, 4636-44	4.8	47
168	Characterization of the two-component abortive phage infection mechanism AbiT from Lactococcus lactis. <i>Journal of Bacteriology</i> , <b>2002</b> , 184, 6325-32	3.5	47
167	Structure and activity of AbiQ, a lactococcal endoribonuclease belonging to the type III toxin-antitoxin system. <i>Molecular Microbiology</i> , <b>2013</b> , 87, 756-68	4.1	46
166	Phages as friends and enemies in food processing. <i>Current Opinion in Biotechnology</i> , <b>2018</b> , 49, 185-190	11.4	45
165	Characterization of Lactococcus lactis phage 949 and comparison with other lactococcal phages. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 6843-52	4.8	45
164	Lactococcal phage genes involved in sensitivity to AbiK and their relation to single-strand annealing proteins. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 3649-52	3.5	45
163	Phage morphology recapitulates phylogeny: the comparative genomics of a new group of myoviruses. <i>PLoS ONE</i> , <b>2012</b> , 7, e40102	3.7	43
162	Identification of a new P335 subgroup through molecular analysis of lactococcal phages Q33 and BM13. <i>Applied and Environmental Microbiology</i> , <b>2013</b> , 79, 4401-9	4.8	43
161	Diversity of Streptococcus thermophilus phages in a large-production cheese factory in Argentina. <i>Journal of Dairy Science</i> , <b>2006</b> , 89, 3791-9	4	43
160	Genome annotation and intraviral interactome for the Streptococcus pneumoniae virulent phage Dp-1. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 551-62	3.5	39
159	Expression and site-directed mutagenesis of the lactococcal abortive phage infection protein AbiK. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 3721-30	3.5	39
158	Novel food-grade plasmid vector based on melibiose fermentation for the genetic engineering of Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2002</b> , 68, 6152-61	4.8	39
157	Preventing phage lysis of Lactococcus lactis in cheese production using a neutralizing heavy-chain antibody fragment from llama. <i>Journal of Dairy Science</i> , <b>2002</b> , 85, 1376-82	4	39
156	Improving the safety of Staphylococcus aureus polyvalent phages by their production on a Staphylococcus xylosus strain. <i>PLoS ONE</i> , <b>2014</b> , 9, e102600	3.7	37
155	Fat-free yogurt made using a galactose-positive exopolysaccharide-producing recombinant strain of Streptococcus thermophilus. <i>Journal of Dairy Science</i> , <b>2009</b> , 92, 477-82	4	37

154	KSY1, a lactococcal phage with a T7-like transcription. <i>Virology</i> , <b>2007</b> , 365, 1-9	3.6	37
153	Characterization of Streptococcus thermophilus host range phage mutants. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 3036-41	4.8	37
152	Global gene expression analysis of two Streptococcus thermophilus bacteriophages using DNA microarray. <i>Virology</i> , <b>2005</b> , 340, 192-208	3.6	37
151	Isolation and characterization of a Streptococcus thermophilus plasmid closely related to the pMV158 family. <i>Plasmid</i> , <b>2001</b> , 45, 171-83	3.3	37
150	Evaluation of Filters for the Sampling and Quantification of RNA Phage Aerosols. <i>Aerosol Science and Technology</i> , <b>2010</b> , 44, 893-901	3.4	36
149	Monoclonal antibodies raised against native major capsid proteins of lactococcal c2-like bacteriophages. <i>Applied and Environmental Microbiology</i> , <b>1998</b> , 64, 4255-9	4.8	36
148	A reverse transcriptase-related protein mediates phage resistance and polymerizes untemplated DNA in vitro. <i>Nucleic Acids Research</i> , <b>2011</b> , 39, 7620-9	20.1	35
147	Characterization of genes involved in the metabolism of alpha-galactosides by Lactococcus raffinolactis. <i>Applied and Environmental Microbiology</i> , <b>2003</b> , 69, 4049-56	4.8	35
146	Immune loss as a driver of coexistence during host-phage coevolution. ISME Journal, 2018, 12, 585-597	11.9	34
145	7-Deazaguanine modifications protect phage DNA from host restriction systems. <i>Nature Communications</i> , <b>2019</b> , 10, 5442	17.4	34
145		17.4 4.9	34
	Communications, 2019, 10, 5442  Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in		
144	Communications, 2019, 10, 5442  Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in Uruguay and ectopic spacer acquisition in CRISPR array. Scientific Reports, 2017, 7, 43438  Effect of the abortive infection mechanism and type III toxin/antitoxin system AbiQ on the lytic	4.9	33
144	Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in Uruguay and ectopic spacer acquisition in CRISPR array. <i>Scientific Reports</i> , <b>2017</b> , 7, 43438  Effect of the abortive infection mechanism and type III toxin/antitoxin system AbiQ on the lytic cycle of Lactococcus lactis phages. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 3947-56  Genome organization and characterization of the virulent lactococcal phage 1358 and its	4·9 3·5	33
144 143 142	Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in Uruguay and ectopic spacer acquisition in CRISPR array. <i>Scientific Reports</i> , <b>2017</b> , 7, 43438  Effect of the abortive infection mechanism and type III toxin/antitoxin system AbiQ on the lytic cycle of Lactococcus lactis phages. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 3947-56  Genome organization and characterization of the virulent lactococcal phage 1358 and its similarities to Listeria phages. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 1623-32  Solution and electron microscopy characterization of lactococcal phage baseplates expressed in	4·9 3·5 4.8	<ul><li>33</li><li>33</li><li>33</li></ul>
144 143 142 141	Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in Uruguay and ectopic spacer acquisition in CRISPR array. <i>Scientific Reports</i> , <b>2017</b> , 7, 43438  Effect of the abortive infection mechanism and type III toxin/antitoxin system AbiQ on the lytic cycle of Lactococcus lactis phages. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 3947-56  Genome organization and characterization of the virulent lactococcal phage 1358 and its similarities to Listeria phages. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 1623-32  Solution and electron microscopy characterization of lactococcal phage baseplates expressed in Escherichia coli. <i>Journal of Structural Biology</i> , <b>2010</b> , 172, 75-84  Genome analysis of two virulent Streptococcus thermophilus phages isolated in Argentina.	4·9 3·5 4.8	<ul><li>33</li><li>33</li><li>33</li><li>33</li></ul>
144 143 142 141 140	Phage-host interactions in Streptococcus thermophilus: Genome analysis of phages isolated in Uruguay and ectopic spacer acquisition in CRISPR array. <i>Scientific Reports</i> , <b>2017</b> , 7, 43438  Effect of the abortive infection mechanism and type III toxin/antitoxin system AbiQ on the lytic cycle of Lactococcus lactis phages. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 3947-56  Genome organization and characterization of the virulent lactococcal phage 1358 and its similarities to Listeria phages. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 1623-32  Solution and electron microscopy characterization of lactococcal phage baseplates expressed in Escherichia coli. <i>Journal of Structural Biology</i> , <b>2010</b> , 172, 75-84  Genome analysis of two virulent Streptococcus thermophilus phages isolated in Argentina. <i>International Journal of Food Microbiology</i> , <b>2009</b> , 136, 101-9  Machine learning assisted design of highly active peptides for drug discovery. <i>PLoS Computational</i>	4.9 3.5 4.8 3.4 5.8	<ul><li>33</li><li>33</li><li>33</li><li>33</li><li>33</li></ul>

136	The targeted recognition of Lactococcus lactis phages to their polysaccharide receptors. <i>Molecular Microbiology</i> , <b>2015</b> , 96, 875-86	4.1	29
135	Inactivation of dairy bacteriophages by commercial sanitizers and disinfectants. <i>International Journal of Food Microbiology</i> , <b>2014</b> , 171, 41-7	5.8	29
134	AbiV, a novel antiphage abortive infection mechanism on the chromosome of Lactococcus lactis subsp. cremoris MG1363. <i>Applied and Environmental Microbiology</i> , <b>2008</b> , 74, 6528-37	4.8	29
133	Genetic and biochemical characterization of the phosphoenolpyruvate:glucose/mannose phosphotransferase system of Streptococcus thermophilus. <i>Applied and Environmental Microbiology</i> , <b>2003</b> , 69, 5423-32	4.8	29
132	DNA sequence analysis of three Lactococcus lactis plasmids encoding phage resistance mechanisms. <i>Journal of Dairy Science</i> , <b>2001</b> , 84, 1610-20	4	29
131	Resistance of Aerosolized Bacterial Viruses to Relative Humidity and Temperature. <i>Applied and Environmental Microbiology</i> , <b>2015</b> , 81, 7305-11	4.8	28
130	A Syst-OMICS Approach to Ensuring Food Safety and Reducing the Economic Burden of Salmonellosis. <i>Frontiers in Microbiology</i> , <b>2017</b> , 8, 996	5.7	28
129	Characterization of two virulent phages of Lactobacillus plantarum. <i>Applied and Environmental Microbiology</i> , <b>2012</b> , 78, 8719-34	4.8	28
128	Cryo-electron microscopy structure of lactococcal siphophage 1358 virion. <i>Journal of Virology</i> , <b>2014</b> , 88, 8900-10	6.6	27
127	P087, a lactococcal phage with a morphogenesis module similar to an Enterococcus faecalis prophage. <i>Virology</i> , <b>2009</b> , 388, 49-56	3.6	27
126	Microbiological and molecular impacts of AbiK on the lytic cycle of Lactococcus lactis phages of the 936 and P335 species. <i>Microbiology (United Kingdom)</i> , <b>2000</b> , 146 ( Pt 2), 445-453	2.9	27
125	Comparison of advanced whole genome sequence-based methods to distinguish strains of Salmonella enterica serovar Heidelberg involved in foodborne outbreaks in Qubec. <i>Food Microbiology</i> , <b>2018</b> , 73, 99-110	6	25
124	Efficacy of two Staphylococcus aureus phage cocktails in cheese production. <i>International Journal of Food Microbiology</i> , <b>2016</b> , 217, 7-13	5.8	25
123	Crystal structure of ORF12 from Lactococcus lactis phage p2 identifies a tape measure protein chaperone. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 728-34	3.5	25
122	Characterization of a galactokinase-positive recombinant strain of Streptococcus thermophilus. <i>Applied and Environmental Microbiology</i> , <b>2004</b> , 70, 4596-603	4.8	25
121	Versatile and robust genome editing with CRISPR1-Cas9. <i>Genome Research</i> , <b>2020</b> , 30, 107-117	9.7	25
120	Prophage Sequence Profiles Reflect Genome Diversity and Can Be Used for High Discrimination Subtyping. <i>Frontiers in Microbiology</i> , <b>2018</b> , 9, 836	5.7	24
119	Cas9 Allosteric Inhibition by the Anti-CRISPR Protein AcrIIA6. <i>Molecular Cell</i> , <b>2019</b> , 76, 922-937.e7	17.6	24

# (2016-2007)

118	Lactobacilli expressing llama VHH fragments neutralise Lactococcus phages. <i>BMC Biotechnology</i> , <b>2007</b> , 7, 58	3.5	24
117	Argentinean Lactococcus lactis bacteriophages: genetic characterization and adsorption studies. <i>Journal of Applied Microbiology</i> , <b>2008</b> , 104, 371-9	4.7	24
116	CRISPRStudio: A User-Friendly Software for Rapid CRISPR Array Visualization. Viruses, 2018, 10,	6.2	24
115	Characterization and diversity of phages infecting Aeromonas salmonicida subsp. salmonicida. <i>Scientific Reports</i> , <b>2017</b> , 7, 7054	4.9	23
114	Distribution and composition of the lysis cassette of Lactococcus lactis phages and functional analysis of bacteriophage ul36 holin. <i>FEMS Microbiology Letters</i> , <b>2004</b> , 233, 37-43	2.9	23
113	Detecting natural adaptation of the Streptococcus thermophilus CRISPR-Cas systems in research and classroom settings. <i>Nature Protocols</i> , <b>2017</b> , 12, 547-565	18.8	22
112	Lactococcal phage p2 ORF35-Sak3 is an ATPase involved in DNA recombination and AbiK mechanism. <i>Molecular Microbiology</i> , <b>2011</b> , 80, 102-16	4.1	22
111	Identification and characterization of the phage gene sav, involved in sensitivity to the lactococcal abortive infection mechanism AbiV. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 2484-94	4.8	22
110	Comparison of Polycarbonate and Polytetrafluoroethylene Filters for Sampling of Airborne Bacteriophages. <i>Aerosol Science and Technology</i> , <b>2010</b> , 44, 197-201	3.4	20
109	Crystal structure of a chimeric receptor binding protein constructed from two lactococcal phages. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 3220-5	3.5	20
108	Involvement of the major capsid protein and two early-expressed phage genes in the activity of the lactococcal abortive infection mechanism AbiT. <i>Applied and Environmental Microbiology</i> , <b>2012</b> , 78, 6890-	g4.8	20
107	Role of galK and galM in galactose metabolism by Streptococcus thermophilus. <i>Applied and Environmental Microbiology</i> , <b>2008</b> , 74, 1264-7	4.8	20
106	Characterization of Five Podoviridae Phages Infecting Citrobacter freundii. <i>Frontiers in Microbiology</i> , <b>2016</b> , 7, 1023	5.7	20
105	Evaluation of bacterial contaminants found on unused paper towels and possible postcontamination after handwashing: a pilot study. <i>American Journal of Infection Control</i> , <b>2012</b> , 40, e5-	<b>3</b> .8	19
104	Lactococcal abortive infection protein AbiV interacts directly with the phage protein SaV and prevents translation of phage proteins. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 7085-92	4.8	19
103	Deciphering the function of lactococcal phage ul36 Sak domains. <i>Journal of Structural Biology</i> , <b>2010</b> , 170, 462-9	3.4	19
102	Biology and genome sequence of Streptococcus mutans phage M102AD. <i>Applied and Environmental Microbiology</i> , <b>2012</b> , 78, 2264-71	4.8	19
101	Programming Native CRISPR Arrays for the Generation of Targeted Immunity. <i>MBio</i> , <b>2016</b> , 7,	7.8	18

100	Diversity and Host Specificity Revealed by Biological Characterization and Whole Genome Sequencing of Bacteriophages Infecting. <i>Viruses</i> , <b>2019</b> , 11,	6.2	17
99	Study of mesophilic Aeromonas salmonicida A527 strain sheds light on the speciesRifestyles and taxonomic dilemma. <i>FEMS Microbiology Letters</i> , <b>2017</b> , 364,	2.9	17
98	Multilocus sequence typing scheme for the characterization of 936-like phages infecting Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2012</b> , 78, 4646-53	4.8	17
97	Applications of phage resistance in lactic acid bacteria <b>1999</b> , 377-382		17
96	Streamlining CRISPR spacer-based bacterial host predictions to decipher the viral dark matter. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, 3127-3138	20.1	17
95	A proposed new bacteriophage subfamily: "Jerseyvirinae". <i>Archives of Virology</i> , <b>2015</b> , 160, 1021-33	2.6	16
94	The CRISPR-Cas Immune System and Genetic Transfers: Reaching an Equilibrium. <i>Microbiology Spectrum</i> , <b>2015</b> , 3, PLAS-0034-2014	8.9	16
93	The doubly phosphorylated form of HPr, HPr(Ser~P)(His-P), is abundant in exponentially growing cells of Streptococcus thermophilus and phosphorylates the lactose transporter LacS as efficiently as HPr(His~P). <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 1364-72	4.8	16
92	Production of Monoclonal Antibodies against the Major Capsid Protein of the Lactococcus Bacteriophage ul36 and Development of an Enzyme-Linked Immunosorbent Assay for Direct Phage Detection in Whey and Milk. <i>Applied and Environmental Microbiology</i> , <b>1993</b> , 59, 2034-40	4.8	16
91	The proteome and interactome of Streptococcus pneumoniae phage Cp-1. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 3135-8	3.5	15
90	Use of an alpha-galactosidase gene as a food-grade selection marker for Streptococcus thermophilus. <i>Journal of Dairy Science</i> , <b>2005</b> , 88, 2341-7	4	15
89	Characterization of a theta-replicating plasmid from Streptococcus thermophilus. <i>Plasmid</i> , <b>2004</b> , 51, 24-36	3.3	15
88	Resistance of Aerosolized Bacterial Viruses to Four Germicidal Products. <i>PLoS ONE</i> , <b>2016</b> , 11, e0168815	3.7	15
87	A mutation in the methionine aminopeptidase gene provides phage resistance in Streptococcus thermophilus. <i>Scientific Reports</i> , <b>2019</b> , 9, 13816	4.9	14
86	Phages of dairy Leuconostoc mesenteroides: genomics and factors influencing their adsorption. <i>International Journal of Food Microbiology</i> , <b>2015</b> , 201, 58-65	5.8	14
85	A genomic approach to understand interactions between Streptococcus pneumoniae and its bacteriophages. <i>BMC Genomics</i> , <b>2015</b> , 16, 972	4.5	14
84	Structure and function of phage p2 ORF34(p2), a new type of single-stranded DNA binding protein. <i>Molecular Microbiology</i> , <b>2009</b> , 73, 1156-70	4.1	14
83	Detection and quantification of capsular exopolysaccharides from Streptococcus thermophilus using lectin probes. <i>Journal of Dairy Science</i> , <b>2006</b> , 89, 4156-62	4	14

82	Investigating the requirement for calcium during lactococcal phage infection. <i>International Journal of Food Microbiology</i> , <b>2015</b> , 201, 47-51	5.8	13
81	Complete Genome Sequence of Streptococcus thermophilus SMQ-301, a Model Strain for Phage-Host Interactions. <i>Genome Announcements</i> , <b>2015</b> , 3,		13
80	Lactococcus lactis type III-A CRISPR-Cas system cleaves bacteriophage RNA. RNA Biology, <b>2019</b> , 16, 461	1-468	13
79	Characterization of a Novel Panton-Valentine leukocidin (PVL)-encoding staphylococcal phage and its naturally PVL-lacking variant. <i>Applied and Environmental Microbiology</i> , <b>2013</b> , 79, 2828-32	4.8	13
78	Control of Bacteriophages in Industrial Fermentations 2004,		13
77	Characterization of the cro-ori region of the Streptococcus thermophilus virulent bacteriophage DT1. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 1237-46	4.8	13
76	Variability in the durability of CRISPR-Cas immunity. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , <b>2019</b> , 374, 20180097	5.8	12
75	A Protocol for Extraction of Infective Viromes Suitable for Metagenomics Sequencing from Low Volume Fecal Samples. <i>Viruses</i> , <b>2019</b> , 11,	6.2	12
74	Phosphorylation of Streptococcus salivarius lactose permease (LacS) by HPr(His ~ P) and HPr(Ser-P)(His ~ P) and effects on growth. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 6764-72	3.5	12
73	Direct detection of lactococcal bacteriophages in cheese whey using DNA probes. <i>FEMS Microbiology Letters</i> , <b>1992</b> , 92, 169-174	2.9	12
72	A new Microviridae phage isolated from a failed biotechnological process driven by Escherichia coli. <i>Applied and Environmental Microbiology</i> , <b>2014</b> , 80, 6992-7000	4.8	11
71	Diverse virulent pneumophages infect Streptococcus mitis. <i>PLoS ONE</i> , <b>2015</b> , 10, e0118807	3.7	11
70	Technical note: Use of RFLP to characterize Lactococcus lactis strains producing exopolysaccharides. <i>Journal of Dairy Science</i> , <b>2003</b> , 86, 1472-5	4	11
69	Delivery of CRISPR-Cas systems using phage-based vectors. <i>Current Opinion in Biotechnology</i> , <b>2021</b> , 68, 174-180	11.4	11
68	Characterization of the Virulent Myophage ST32. Viruses, 2018, 10,	6.2	11
67	Phagebook: The Social Network. <i>Molecular Cell</i> , <b>2017</b> , 65, 963-964	17.6	10
66	Investigation of the protective effect of whey proteins on lactococcal phages during heat treatment at various pH. <i>International Journal of Food Microbiology</i> , <b>2015</b> , 210, 33-41	5.8	10
65	First Complete Genome Sequence of Staphylococcus xylosus, a Meat Starter Culture and a Host to Propagate Staphylococcus aureus Phages. <i>Genome Announcements</i> , <b>2014</b> , 2,		10

64	Identification of an inducible bacteriophage in a virulent strain of Streptococcus suis serotype 2. <i>Infection and Immunity</i> , <b>2003</b> , 71, 6104-8	3.7	10
63	Effect of fermented milks on humoral immune response in mice. <i>International Dairy Journal</i> , <b>1991</b> , 1, 231-239	3.5	10
62	Cooperation between Different CRISPR-Cas Types Enables Adaptation in an RNA-Targeting System. <i>MBio</i> , <b>2021</b> , 12,	7.8	10
61	Novel Genus of Phages Infecting Streptococcus thermophilus: Genomic and Morphological Characterization. <i>Applied and Environmental Microbiology</i> , <b>2020</b> , 86,	4.8	10
60	Microencapsulation of a Staphylococcus phage for concentration and long-term storage. <i>Food Microbiology</i> , <b>2018</b> , 76, 304-309	6	9
59	Staphylococcus epidermidis bacteriophages from the anterior nares of humans. <i>Applied and Environmental Microbiology</i> , <b>2011</b> , 77, 7853-5	4.8	9
58	Analysis of two theta-replicating plasmids of Streptococcus thermophilus. <i>Plasmid</i> , <b>2007</b> , 58, 174-81	3.3	9
57	Galactose metabolism and capsule formation in a recombinant strain of Streptococcus thermophilus with a galactose-fermenting phenotype. <i>Journal of Dairy Science</i> , <b>2007</b> , 90, 4051-7	4	9
56	Beyond the A-layer: adsorption of lipopolysaccharides and characterization of bacteriophage-insensitive mutants of Aeromonas salmonicida subsp. salmonicida. <i>Molecular Microbiology</i> , <b>2019</b> , 112, 667-677	4.1	8
55	Mutational Analysis of the Antitoxin in the Lactococcal Type III Toxin-Antitoxin System AbiQ. <i>Applied and Environmental Microbiology</i> , <b>2015</b> , 81, 3848-55	4.8	8
54	Characterization of a Type II-A CRISPR-Cas System in. <i>MSphere</i> , <b>2020</b> , 5,	5	8
53	Complete Genome Sequence of Virulent Phage MS1. <i>Genome Announcements</i> , <b>2017</b> , 5,		8
52	Mobilome of Sheds Light on Its Genetic Diversity and Its Adaptation to Smear-Ripened Cheeses. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 1270	5.7	7
51	The double-edged sword of CRISPR-Cas systems. <i>Cell Research</i> , <b>2013</b> , 23, 15-7	24.7	7
50	Characterization of prophages of Lactococcus garvieae. Scientific Reports, 2017, 7, 1856	4.9	7
49	Activation and transfer of the chromosomal phage resistance mechanism AbiV in Lactococcus lactis. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 3358-61	4.8	7
48	A short overview of the CRISPR-Cas adaptation stage. Canadian Journal of Microbiology, 2021, 67, 1-12	3.2	7
47	The Tape Measure Protein Is Involved in the Heat Stability of Lactococcus lactis Phages. <i>Applied and Environmental Microbiology</i> , <b>2018</b> , 84,	4.8	7

46	The CRISPR-Cas app goes viral. Current Opinion in Microbiology, 2017, 37, 103-109	7.9	6
45	Investigating MG1363 Response to Phage p2 Infection at the Proteome Level. <i>Molecular and Cellular Proteomics</i> , <b>2019</b> , 18, 704-714	7.6	6
44	A virulent phage infecting Lactococcus garvieae, with homology to Lactococcus lactis phages. <i>Applied and Environmental Microbiology</i> , <b>2015</b> , 81, 8358-65	4.8	6
43	The DNA binding mechanism of a SSB protein from Lactococcus lactis siphophage p2. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2013</b> , 1834, 1070-6	4	6
42	Complete Genome Sequence of a Staphylococcus epidermidis Bacteriophage Isolated from the Anterior Nares of Humans. <i>Genome Announcements</i> , <b>2014</b> , 2,		6
41	Detection of preQ0 deazaguanine modifications in bacteriophage CAjan DNA using Nanopore sequencing reveals same hypermodification at two distinct DNA motifs. <i>Nucleic Acids Research</i> , <b>2020</b> , 48, 10383-10396	20.1	6
40	Molecular Structure of Lactoferrin Influences the Thermal Resistance of Lactococcal Phages. Journal of Agricultural and Food Chemistry, <b>2017</b> , 65, 2214-2221	5.7	5
39	Comparative genomic analysis of 142 bacteriophages infecting Salmonella enterica subsp. enterica. <i>BMC Genomics</i> , <b>2020</b> , 21, 374	4.5	5
38	Applications of CRISPR-Cas in its natural habitat. Current Opinion in Chemical Biology, 2016, 34, 30-36	9.7	5
37	Production of Bacteriophages by Listeria Cells Entrapped in Organic Polymers. Viruses, 2018, 10,	6.2	5
36	Characterization of CRISPR-Cas systems in the Ralstonia solanacearum species complex. <i>Molecular Plant Pathology</i> , <b>2019</b> , 20, 223-239	5.7	5
35	The effect of bacteriophages on the acidification of a vegetable juice medium by microencapsulated Lactobacillus plantarum. <i>Food Microbiology</i> , <b>2017</b> , 63, 28-34	6	4
34	The EcoChip: A Wireless Multi-Sensor Platform for Comprehensive Environmental Monitoring. <i>IEEE Transactions on Biomedical Circuits and Systems</i> , <b>2018</b> , 12, 1289-1300	5.1	4
33	CRISPR-Cas in the laboratory classroom. <i>Nature Microbiology</i> , <b>2017</b> , 2, 17018	26.6	3
32	Complete Genome Sequence of Escherichia coli Siphophage BRET. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	3
31	Source Tracking Based on Core Genome SNV and CRISPR Typing of Serovar Heidelberg Isolates Involved in Foodborne Outbreaks in Qubec, 2012. <i>Frontiers in Microbiology</i> , <b>2020</b> , 11, 1317	5.7	3
30	How are genes modified? Crossbreeding, mutagenesis, and CRISPR-Cas9 2020, 39-54		3
29	Immune Loss as a Driver of Coexistence During Host-Phage Coevolution		3

28	Structural Insights into Lactococcal Siphophage p2 Baseplate Activation Mechanism. <i>Viruses</i> , <b>2020</b> , 12,	6.2	3
27	Ectopic Spacer Acquisition in CRISPR3 Array. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	3
26	Primed CRISPR-Cas Adaptation and Impaired Phage Adsorption in Streptococcus mutans. <i>MSphere</i> , <b>2021</b> , 6,	5	3
25	Hundreds of viral families in the healthy infant gut		3
24	DNA tandem repeats contribute to the genetic diversity of Brevibacterium aurantiacum phages. <i>Environmental Microbiology</i> , <b>2020</b> , 22, 3413-3428	5.2	2
23	A Jumbo Formation in the Viral Game Plan. CRISPR Journal, 2020, 3, 14-17	2.5	2
22	A Protocol for Extraction of Infective Viromes Suitable for Metagenomics Sequencing from Low Volume Fecal Samples		2
21	Targeted Genome Editing of Virulent Phages Using CRISPR-Cas9. <i>Bio-protocol</i> , <b>2018</b> , 8, e2674	0.9	2
20	Functional Study of the Type II-A CRISPR-Cas System of Hypervirulent Strains. <i>CRISPR Journal</i> , <b>2021</b> , 4, 233-242	2.5	2
19	Complete Genome Sequence of Brevibacterium linens SMQ-1335. <i>Genome Announcements</i> , <b>2016</b> , 4,		2
18	Complete Genome Sequence of Ebrios, a Novel T7virus Isolated from the Ebrie Lagoon in Abidjan, CEe dRvoire. <i>Genome Announcements</i> , <b>2018</b> , 6,		2
17	Procedures for Generating CRISPR Mutants with Novel Spacers Acquired from Viruses or Plasmids. <i>Methods in Molecular Biology</i> , <b>2015</b> , 1311, 195-222	1.4	1
16	Bacteriophages in Industrial Food Processing: Incidence and Control in Industrial Fermentation <b>2014</b> , 199-216		1
15	Type II: Streptococcus thermophilus <b>2013</b> , 171-200		1
14	Abortive Infection Mechanisms and Prophage Sequences Significantly Influence the Genetic Makeup of Emerging Lytic Lactococcal Phages. <i>Journal of Bacteriology</i> , <b>2007</b> , 189, 5787-5787	3.5	1
13	Type II: Streptococcus thermophilus <b>2013</b> , 171		1
12	Evolutionary emergence of infectious diseases in heterogeneous host populations		1
11	Versatile and robust genome editing with Streptococcus thermophilus CRISPR1-Cas9		1

#### LIST OF PUBLICATIONS

10	A Lactococcal Phage Protein Promotes Viral Propagation and Alters the Host Proteomic Response During Infection. <i>Viruses</i> , <b>2020</b> , 12,	6.2	1
9	Genomic diversity and CRISPR-Cas systems in the cyanobacterium Nostoc in the High Arctic. <i>Environmental Microbiology</i> , <b>2021</b> , 23, 2955-2968	5.2	1
8	The endless battle between phages and CRISPR-Cas systems in. <i>Biochemistry and Cell Biology</i> , <b>2021</b> , 99, 397-402	3.6	1
7	Would Bacteriophages Be a New Old Complement to Antibiotics in Aquaculture? <b>2019</b> , 51-68		O
6	Zebrafish: a big fish in the study of the gut microbiota. Current Opinion in Biotechnology, 2021, 73, 308-3	3 <b>1:3</b> .4	O
5	Induction and Elimination of Prophages Using CRISPR Interference. CRISPR Journal, 2021, 4, 549-557	2.5	O
4	A truncated anti-CRISPR protein prevents spacer acquisition but not interference <i>Nature Communications</i> , <b>2022</b> , 13, 2802	17.4	O
3	The CRISPR-Cas Immune System and Genetic Transfers: Reaching an Equilibrium209-218		
2	Phosphorylation, an Altruistic Bacterial Trick to Halt Phages. Cell Host and Microbe, 2016, 20, 409-410	23.4	
1	CRISPR-Cas Systems in Starter Cultures <b>2022</b> , 103-112		