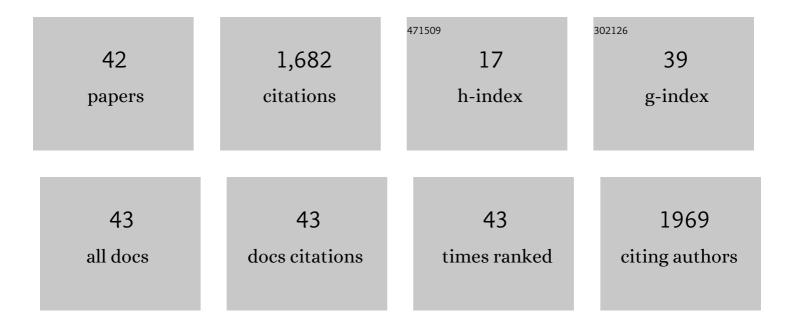
## Annika Gillis

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Bacilysin within the Bacillus subtilis group: gene prevalence versus antagonistic activity against Gram-negative foodborne pathogens. Journal of Biotechnology, 2021, 327, 28-35.	3.8	28
2	Modifications of cell wall polymers in Gram-positive bacteria by multi-component transmembrane glycosylation systems. Current Opinion in Microbiology, 2021, 60, 24-33.	5.1	19
3	Comparative Genomics of Prophages Sato and Sole Expands the Genetic Diversity Found in the Genus Betatectivirus. Microorganisms, 2021, 9, 1335.	3.6	1
4	Bacterial Viruses Subcommittee and Archaeal Viruses Subcommittee of the ICTV: update of taxonomy changes in 2021. Archives of Virology, 2021, 166, 3239-3244.	2.1	24
5	Bacillus cytotoxicus Genomics: Chromosomal Diversity and Plasmidome Versatility. Frontiers in Microbiology, 2021, 12, 789929.	3.5	5
6	Analysis of Spounaviruses as a Case Study for the Overdue Reclassification of Tailed Phages. Systematic Biology, 2020, 69, 110-123.	5.6	89
7	Characterization of PlyB221 and PlyP32, Two Novel Endolysins Encoded by Phages Preying on the Bacillus cereus Group. Viruses, 2020, 12, 1052.	3.3	12
8	Taxonomy of prokaryotic viruses: 2018-2019 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. Archives of Virology, 2020, 165, 1253-1260.	2.1	144
9	Tomato Twisted Leaf Virus: A Novel Indigenous New World Monopartite Begomovirus Infecting Tomato in Venezuela. Viruses, 2019, 11, 327.	3.3	18
10	Overview of the Antimicrobial Compounds Produced by Members of the Bacillus subtilis Group. Frontiers in Microbiology, 2019, 10, 302.	3.5	425
11	Biocontrol potential of phage Deep-Blue against psychrotolerant Bacillus weihenstephanensis. Food Control, 2019, 102, 94-103.	5.5	7
12	A novel T4SSâ€mediated DNA transfer used by pXO16, a conjugative plasmid from <i>Bacillus thuringiensis</i> serovar <i>israelensis</i> . Environmental Microbiology, 2018, 20, 1550-1561.	3.8	15
13	Complete Genome Sequence of Bacillus velezensis CN026 Exhibiting Antagonistic Activity against Gram-Negative Foodborne Pathogens. Genome Announcements, 2018, 6, .	0.8	7
14	Diversity and enzymatic potentialities of Bacillus sp. strains isolated from a polluted freshwater ecosystem in Cuba. World Journal of Microbiology and Biotechnology, 2018, 34, 28.	3.6	5
15	Taxonomy of prokaryotic viruses: 2017 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. Archives of Virology, 2018, 163, 1125-1129.	2.1	172
16	Complete genome sequence of two tomato-infecting begomoviruses in Venezuela: evidence of a putative novel species and a novel recombinant strain. Archives of Virology, 2018, 163, 555-558.	2.1	5
17	Role of plasmid plasticity and mobile genetic elements in the entomopathogen Bacillus thuringiensis serovar israelensis. FEMS Microbiology Reviews, 2018, 42, 829-856.	8.6	33
18	Bacterial Sexuality at the Nanoscale. Nano Letters, 2018, 18, 5821-5826.	9.1	11

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19	Versatile Antagonistic Activities of Soil-Borne Bacillus spp. and Pseudomonas spp. against Phytophthora infestans and Other Potato Pathogens. Frontiers in Microbiology, 2018, 9, 143.	3.5	114
20	Complete genome sequence of bacteriophage Deep-Purple, a novel member of the family Siphoviridae infecting Bacillus cereus. Archives of Virology, 2018, 163, 2555-2559.	2.1	6
21	Horizontal transfer of chromosomal markers mediated by the large conjugative plasmid pXO16 from Bacillus thuringiensis serovar israelensis. Plasmid, 2017, 91, 76-81.	1.4	12
22	First Report of Bell Pepper Soft-Rot Caused by Pectobacterium carotovorum subsp. brasiliense in Venezuela. Plant Disease, 2017, 101, 1671-1671.	1.4	18
23	Molecular and biological characterization of a new Tomato mild yellow leaf curl Aragua virus strain producing severe symptoms in tomato. Virus Genes, 2017, 53, 939-942.	1.6	1
24	Taxonomy of prokaryotic viruses: 2016 update from the ICTV bacterial and archaeal viruses subcommittee. Archives of Virology, 2017, 162, 1153-1157.	2.1	57
25	Detection of the cryptic prophage-like molecule pBtic235 in Bacillus thuringiensis subsp. israelensis. Research in Microbiology, 2017, 168, 319-330.	2.1	19
26	Comparative genomics of extrachromosomal elements in Bacillus thuringiensis subsp. israelensis. Research in Microbiology, 2017, 168, 331-344.	2.1	28
27	Occurrence of Six Begomoviruses Infecting Tomato Fields in Venezuela and Genetic Characterization of <i>Potato Yellow Mosaic Virus</i> Isolates. Journal of Phytopathology, 2016, 164, 697-703.	1.0	14
28	Low genetic diversity of Banana bunchy top virus, with a sub-regional pattern of variation, in Democratic Republic of Congo. Virus Genes, 2016, 52, 900-905.	1.6	13
29	Complete Genome Sequence of Bacteriophage Deep-Blue Infecting Emetic Bacillus cereus. Genome Announcements, 2016, 4, .	0.8	4
30	Taxonomy of prokaryotic viruses: update from the ICTV bacterial and archaeal viruses subcommittee. Archives of Virology, 2016, 161, 1095-1099.	2.1	83
31	pXO16 from Bacillus thuringiensis serovar israelensis: Almost 350 kb of terra incognita. Plasmid, 2015, 80, 8-15.	1.4	15
32	Influence of Lysogeny of Tectiviruses GIL01 and GIL16 on Bacillus thuringiensis Growth, Biofilm Formation, and Swarming Motility. Applied and Environmental Microbiology, 2014, 80, 7620-7630.	3.1	28
33	Phages Preying on Bacillus anthracis, Bacillus cereus, and Bacillus thuringiensis: Past, Present and Future. Viruses, 2014, 6, 2623-2672.	3.3	89
34	Full-length genome sequencing of the mild strain of Tomato yellow leaf curl virus in Venezuela reveals a third introduction event of this virus in New World. Australasian Plant Disease Notes, 2014, 9, 1.	0.7	5
35	Prevalence, Genetic Diversity, and Host Range of Tectiviruses among Members of the Bacillus cereus Group. Applied and Environmental Microbiology, 2014, 80, 4138-4152.	3.1	28
36	Serratia marcescens associated with bell pepper (Capsicum annuum L.) soft-rot disease under greenhouse conditions. European Journal of Plant Pathology, 2014, 138, 1-8.	1.7	21

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37	An improved method for rapid generation and screening of Bacillus thuringiensis phage-resistant mutants. Journal of Microbiological Methods, 2014, 106, 101-103.	1.6	5
38	Identification of five novel tectiviruses in Bacillus strains: analysis ofÂa highly variable region generating genetic diversity. Research in Microbiology, 2013, 164, 118-126.	2.1	24
39	Nanoscale imaging of Bacillus thuringiensis flagella using atomic force microscopy. Nanoscale, 2012, 4, 1585-1591.	5.6	34
40	Atomic force microscopy: A powerful tool for studying bacterial swarming motility. Micron, 2012, 43, 1304-1311.	2.2	16
41	Bacillus thuringiensis improved isolation methodology from soil samples. Journal of Microbiological Methods, 2008, 75, 357-358.	1.6	17
42	Detección de genes cry1 y cry2 en cepas venezolanas de Bacillus thuringiensis y sus posibles aplicaciones en el campo de la BiotecnologÃa AgrÃcola. IFMBE Proceedings, 2007, , 975-978.	0.3	0