

# Maarten J Gilbert

## List of Publications by Year in descending order

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16  
papers

394  
citations

933447

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940533

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16  
docs citations

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times ranked

424  
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Campylobacter fetus</i> subsp. <i>testudinum</i> subsp. nov., isolated from humans and reptiles. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 2944-2948.	1.7	78
2	Human Infections with New Subspecies of <i>Campylobacter fetus</i> . <i>Emerging Infectious Diseases</i> , 2013, 19, 1678-80.	4.3	66
3	Occurrence, Diversity, and Host Association of Intestinal <i>Campylobacter</i> , <i>Arcobacter</i> , and <i>Helicobacter</i> in Reptiles. <i>PLoS ONE</i> , 2014, 9, e101599.	2.5	43
4	<i>Campylobacter iguaniorum</i> sp. nov., isolated from reptiles. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 975-982.	1.7	32
5	<i>Campylobacter pinnipediorum</i> sp. nov., isolated from pinnipeds, comprising <i>Campylobacter pinnipediorum</i> subsp. <i>pinnipediorum</i> subsp. nov. and <i>Campylobacter pinnipediorum</i> subsp. <i>caledonicus</i> subsp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1961-1968.	1.7	28
6	Mitigating <i>Batrachochytrium</i> salamandrivorans in Europe. <i>Amphibia - Reptilia</i> , 2019, 40, 265-290.	0.5	26
7	Comparative Genomics of <i>Campylobacter fetus</i> from Reptiles and Mammals Reveals Divergent Evolution in Host-Associated Lineages. <i>Genome Biology and Evolution</i> , 2016, 8, 2006-2019.	2.5	25
8	Whole genome-based phylogeny of reptile-associated <i>Helicobacter</i> indicates independent niche adaptation followed by diversification in a poikilothermic host. <i>Scientific Reports</i> , 2017, 7, 8387.	3.3	19
9	Living in Cold Blood: <i>Arcobacter</i> , <i>Campylobacter</i> , and <i>Helicobacter</i> in Reptiles. <i>Frontiers in Microbiology</i> , 2019, 10, 1086.	3.5	18
10	Homologous Recombination between Genetically Divergent <i>Campylobacter fetus</i> Lineages Supports Host-Associated Speciation. <i>Genome Biology and Evolution</i> , 2018, 10, 716-722.	2.5	16
11	Complete Genome Sequence of <i>Campylobacter fetus</i> subsp. <i>testudinum</i> Strain 03-427 <sup>T</sup> . <i>Genome Announcements</i> , 2013, 1, .	0.8	10
12	Complete Genome Sequence of <i>Campylobacter iguaniorum</i> Strain 1485E T, Isolated from a Bearded Dragon ( <i>Pogona vitticeps</i> ). <i>Genome Announcements</i> , 2014, 2, .	0.8	10
13	Forensic microbiology reveals that <i>Neisseria animaloris</i> infections in harbour porpoises follow traumatic injuries by grey seals. <i>Scientific Reports</i> , 2019, 9, 14338.	3.3	9
14	Comparative Genomics of <i>Campylobacter iguaniorum</i> to Unravel Genetic Regions Associated with Reptilian Hosts. <i>Genome Biology and Evolution</i> , 2016, 8, 3022-3029.	2.5	8
15	After the bite: bacterial transmission from grey seals ( <i>Halichoerus grypus</i> ) to harbour porpoises ( ) Tj ETQq1 1 0.784314 rgBT <sub>5</sub> /Overlook	2.4	5
16	<i>Campylobacter pinnipediorum</i> subsp. <i>caledonicus</i> and <i>C. pinnipediorum</i> subsp. <i>pinnipediorum</i> recovered from abscesses in pinnipeds. <i>Diseases of Aquatic Organisms</i> , 2020, 142, 41-46.	1.0	1