

# Benjamin Raymond

## List of Publications by Year in descending order

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Version: 2024-02-01

52  
papers

1,956  
citations

236833

25  
h-index

265120

42  
g-index

56  
all docs

56  
docs citations

56  
times ranked

1987  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bacillus thuringiensis: an impotent pathogen?. Trends in Microbiology, 2010, 18, 189-194.	3.5	297
2	The Dynamics of Cooperative Bacterial Virulence in the Field. Science, 2012, 337, 85-88.	6.0	112
3	Environmental Factors Determining the Epidemiology and Population Genetic Structure of the Bacillus cereus Group in the Field. PLoS Pathogens, 2010, 6, e1000905.	2.1	94
4	A mid-gut microbiota is not required for the pathogenicity of Bacillus thuringiensis to diamondback moth larvae. Environmental Microbiology, 2009, 11, 2556-2563.	1.8	82
5	Live to cheat another day: bacterial dormancy facilitates the social exploitation of $\beta$ -lactamases. ISME Journal, 2016, 10, 778-787.	4.4	79
6	Genes and environment interact to determine the fitness costs of resistance to Bacillus thuringiensis. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 1519-1524.	1.2	74
7	Five rules for resistance management in the antibiotic apocalypse, a road map for integrated microbial management. Evolutionary Applications, 2019, 12, 1079-1091.	1.5	74
8	In defence of Bacillus thuringiensis, the safest and most successful microbial insecticide available to humanity—a response to EFSA. FEMS Microbiology Ecology, 2017, 93, .	1.3	73
9	Cooperation and the evolutionary ecology of bacterial virulence: The Bacillus cereus group as a novel study system. BioEssays, 2013, 35, 706-716.	1.2	60
10	Exploiting pathogens and their impact on fitness costs to manage the evolution of resistance to Bacillus thuringiensis. Journal of Applied Ecology, 2007, 44, 768-780.	1.9	59
11	Genetic and Biochemical Characterization of Field-Evolved Resistance to Bacillus thuringiensis Toxin Cry1Ac in the Diamondback Moth, Plutella xylostella. Applied and Environmental Microbiology, 2004, 70, 7010-7017.	1.4	56
12	Antagonistic competition moderates virulence in Bacillus thuringiensis. Ecology Letters, 2011, 14, 765-772.	3.0	55
13	The Social Biology of Quorum Sensing in a Naturalistic Host Pathogen System. Current Biology, 2014, 24, 2417-2422.	1.8	54
14	Host plant and population determine the fitness costs of resistance to Bacillus thuringiensis. Biology Letters, 2007, 3, 83-86.	1.0	45
15	Bacteria from natural populations transfer plasmids mostly towards their kin. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20191110.	1.2	45
16	Comparative Genomics of Bacillus thuringiensis Reveals a Path to Specialized Exploitation of Multiple Invertebrate Hosts. MBio, 2017, 8, .	1.8	43
17	Lineage-specific plasmid acquisition and the evolution of specialized pathogens in Bacillus thuringiensis and the Bacillus cereus group. Molecular Ecology, 2018, 27, 1524-1540.	2.0	43
18	Host shifting by Operophtera brumata into novel environments leads to population differentiation in life-history traits. Ecological Entomology, 2003, 28, 604-612.	1.1	42

#	ARTICLE	IF	CITATIONS
19	Rhizobacterial Community Assembly Patterns Vary Between Crop Species. <i>Frontiers in Microbiology</i> , 2019, 10, 581.	1.5	42
20	The role of food plant and pathogen-induced behaviour in the persistence of a nucleopolyhedrovirus. <i>Journal of Invertebrate Pathology</i> , 2005, 88, 49-57.	1.5	38
21	Competition and reproduction in mixed infections of pathogenic and non-pathogenic <i>Bacillus</i> spp.. <i>Journal of Invertebrate Pathology</i> , 2007, 96, 151-155.	1.5	38
22	Strong oviposition preference for Bt over non-Bt maize in <i>Spodoptera frugiperda</i> and its implications for the evolution of resistance. <i>BMC Biology</i> , 2014, 12, 48.	1.7	36
23	Quantifying the reproduction of <i>Bacillus thuringiensis</i> HD1 in cadavers and live larvae of <i>Plutella xylostella</i> . <i>Journal of Invertebrate Pathology</i> , 2008, 98, 307-313.	1.5	35
24	Ecological consequences of ingestion of <i>Bacillus cereus</i> on <i>Bacillus thuringiensis</i> infections and on the gut flora of a lepidopteran host. <i>Journal of Invertebrate Pathology</i> , 2008, 99, 103-111.	1.5	31
25	Impact of intraguild predation on parasitoid foraging behaviour. <i>Ecological Entomology</i> , 2010, 35, 183-189.	1.1	31
26	Biofilms facilitate cheating and social exploitation of $\beta$ -lactam resistance in <i>Escherichia coli</i> . <i>Npj Biofilms and Microbiomes</i> , 2019, 5, 36.	2.9	27
27	Bacterial Cooperation Causes Systematic Errors in Pathogen Risk Assessment due to the Failure of the Independent Action Hypothesis. <i>PLoS Pathogens</i> , 2015, 11, e1004775.	2.1	26
28	Division of labour and terminal differentiation in a novel <i>Bacillus thuringiensis</i> strain. <i>ISME Journal</i> , 2015, 9, 286-296.	4.4	26
29	Negative frequency dependent selection on plasmid carriage and low fitness costs maintain extended spectrum $\beta$ -lactamases in <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2019, 9, 17211.	1.6	25
30	Making pathogens sociable: The emergence of high relatedness through limited host invasibility. <i>ISME Journal</i> , 2015, 9, 2315-2323.	4.4	20
31	Ecological and genetic determinants of plasmid distribution in <i>Escherichia coli</i> . <i>Environmental Microbiology</i> , 2016, 18, 4230-4239.	1.8	16
32	Aseptic Rearing and Infection with Gut Bacteria Improve the Fitness of Transgenic Diamondback Moth, <i>Plutella xylostella</i> . <i>Insects</i> , 2019, 10, 89.	1.0	16
33	Limiting opportunities for cheating stabilizes virulence in insect parasitic nematodes. <i>Evolutionary Applications</i> , 2016, 9, 462-470.	1.5	15
34	Shifts along the parasite-mutualist continuum are opposed by fundamental trade-offs. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20190236.	1.2	13
35	Moderation of pathogen-induced mortality: the role of density in <i>Bacillus thuringiensis</i> virulence. <i>Biology Letters</i> , 2009, 5, 218-220.	1.0	12
36	Combining the high-dose/refuge strategy and self-limiting transgenic insects in resistance management—A test in experimental mesocosms. <i>Evolutionary Applications</i> , 2018, 11, 727-738.	1.5	12

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37	The impact of strain diversity and mixed infections on the evolution of resistance to <i>Bacillus thuringiensis</i> . Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20131497.	1.2	11
38	Crystal toxins and the volunteer's dilemma in bacteria. Journal of Evolutionary Biology, 2019, 32, 310-319.	0.8	11
39	Passage and the evolution of virulence in invertebrate pathogens: Fundamental and applied perspectives. Journal of Invertebrate Pathology, 2022, 187, 107692.	1.5	10
40	The Biology, Ecology and Taxonomy of <i>Bacillus thuringiensis</i> and Related Bacteria. , 2017, , 19-39.		9
41	The application of self-limiting transgenic insects in managing resistance in experimental metapopulations. Journal of Applied Ecology, 2019, 56, 688-698.	1.9	8
42	Controlling insecticide resistant clones of the aphid, <i>Myzus persicae</i> , using the entomopathogenic fungus <i>Akanthomyces muscarius</i> : Fitness cost of resistance under pathogen challenge. Pest Management Science, 2021, 77, 5286-5293.	1.7	8
43	Targeting antibiotic resistant bacteria with phage reduces bacterial density in an insect host. Biology Letters, 2019, 15, 20180895.	1.0	7
44	Divergence in environmental adaptation between terrestrial clades of the <i>Bacillus cereus</i> group. FEMS Microbiology Ecology, 2020, 97, .	1.3	7
45	An appeal for a more evidence based approach to biopesticide safety in the EU. FEMS Microbiology Ecology, 2018, 94, .	1.3	6
46	HTA-SuperSAGE of the gut tissue of a Vip3A-resistant <i>Heliothis virescens</i> (Lepidoptera: Noctuidae) strain provides insights into the basis of resistance. Insect Science, 2019, 26, 479-498.	1.5	5
47	Optimal Response to Quorum-Sensing Signals Varies in Different Host Environments with Different Pathogen Group Size. MBio, 2020, 11, .	1.8	5
48	Relative efficacy of biological control and cultural management for control of mollusc pests in cool climate vineyards. Biocontrol Science and Technology, 2021, 31, 725-738.	0.5	4
49	Signatures of selection in core and accessory genomes indicate different ecological drivers of diversification among <i>Bacillus cereus</i> clades. Molecular Ecology, 2022, 31, 3584-3597.	2.0	4
50	Lethal pathogens, non-lethal synergists and the evolutionary ecology of resistance. Journal of Theoretical Biology, 2008, 254, 339-349.	0.8	3
51	Strong Environment-Genotype Interactions Determine the Fitness Costs of Antibiotic Resistance <i>In Vitro</i> and in an Insect Model of Infection. Antimicrobial Agents and Chemotherapy, 2020, 64, .	1.4	3
52	Function is a better predictor of plant rhizosphere community membership than <i>16S</i> phylogeny. Environmental Microbiology, 2021, 23, 6089-6103.	1.8	3