## **Roy Gross**

## List of Publications by Year in descending order

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ROV GROSS

#	Article	IF	CITATIONS
1	Regulation of bacterial virulence by two-component systems. Current Opinion in Microbiology, 2006, 9, 143-152.	2.3	371
2	The genome sequence of Blochmannia floridanus: Comparative analysis of reduced genomes. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 9388-9393.	3.3	338
3	Sequences required for expression of Bordetella pertussis virulence factors share homology with prokaryotic signal transduction proteins. Proceedings of the National Academy of Sciences of the United States of America, 1989, 86, 6671-6675.	3.3	306
4	Metabolic Interdependence of Obligate Intracellular Bacteria and Their Insect Hosts. Microbiology and Molecular Biology Reviews, 2004, 68, 745-770.	2.9	259
5	Nutritional upgrading for omnivorous carpenter ants by the endosymbiont Blochmannia. BMC Biology, 2007, 5, 48.	1.7	244
6	Families of bacterial signal-transducing proteins. Molecular Microbiology, 1989, 3, 1661-1667.	1.2	187
7	Intracellular endosymbiotic bacteria of Camponotus species (carpenter ants): systematics, evolution and ultrastructural characterization. Molecular Microbiology, 1996, 21, 479-489.	1.2	142
8	Signalling pathways in two-component phosphorelay systems. Trends in Microbiology, 1999, 7, 115-120.	3.5	136
9	Identification of Immunodominant Antigens from Helicobacter pylori and Evaluation of Their Reactivities with Sera from Patients with Different Gastroduodenal Pathologies. Infection and Immunity, 2000, 68, 915-920.	1.0	119
10	Positive regulation of pertussis toxin expression Proceedings of the National Academy of Sciences of the United States of America, 1988, 85, 3913-3917.	3.3	111
11	The virulence regulator protein of Listeria ivanovii is highly homologous to PrfA from Listeria monocytogenes and both belong to the Crp-Fnr family of transcription regulators. Molecular Microbiology, 1994, 13, 141-151.	1.2	98
12	Evolutionary relationships in the genus Bordetella. Molecular Microbiology, 1987, 1, 301-308.	1.2	92
13	Plasmid and chromosomal mutants in the iron(III)-aerobactin transport system of Escherichia coli. Use of streptonigrin for selection. Molecular Genetics and Genomics, 1983, 192, 131-139.	2.4	90
14	Analysis of and function predictions for previously conserved hypothetical or putative proteins in Blochmannia floridanus. BMC Microbiology, 2006, 6, 1.	1.3	87
15	Evolutionary trends in the genus Bordetella. Microbes and Infection, 2001, 3, 61-72.	1.0	86
16	The Lipopolysaccharide of <i>Bordetella bronchiseptica</i> Acts as a Protective Shield against Antimicrobial Peptides. Infection and Immunity, 1998, 66, 5607-5612.	1.0	86
17	The missing link: Bordetella petrii is endowed with both the metabolic versatility of environmental bacteria and virulence traits of pathogenic Bordetellae. BMC Genomics, 2008, 9, 449.	1.2	85
18	Bacterial microbiota associated with ants of the genus Tetraponera. Biological Journal of the Linnean Society, 2007, 90, 399-412.	0.7	82

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19	Immune reactions of insects on bacterial pathogens and mutualists. Microbes and Infection, 2008, 10, 1082-1088.	1.0	82
20	Insights into the microbial world associated with ants. Archives of Microbiology, 2005, 184, 199-206.	1.0	80
21	Immunity and symbiosis. Molecular Microbiology, 2009, 73, 751-759.	1.2	80
22	Identification of the genes and their polypeptide products responsible for aerobactin synthesis by pCoIV plasmids. Molecular Genetics and Genomics, 1985, 201, 204-212.	2.4	79
23	Tissue Localization of the Endosymbiotic Bacterium " Candidatus Blochmannia floridanus―in Adults and Larvae of the Carpenter Ant Camponotus floridanus. Applied and Environmental Microbiology, 2002, 68, 4187-4193.	1.4	79
24	Bacteriocyte dynamics during development of a holometabolous insect, the carpenter ant Camponotus floridanus. BMC Microbiology, 2010, 10, 308.	1.3	72
25	Intracellular survival strategies of mutualistic and parasitic prokaryotes. Trends in Microbiology, 2001, 9, 267-273.	3.5	71
26	Insects as hosts for mutualistic bacteria. International Journal of Medical Microbiology, 2009, 299, 1-8.	1.5	70
27	Relevance of the Endosymbiosis of Blochmannia floridanus and Carpenter Ants at Different Stages of the Host. Applied and Environmental Microbiology, 2006, 72, 6027-6033.	1.4	69
28	Specificity of the BvgAS and EvgAS phosphorelay is mediated by the Câ€ŧerminal HPt domains of the sensor proteins. Molecular Microbiology, 1998, 27, 875-887.	1.2	65
29	The unorthodox histidine kinases BvgS and EvgS are responsive to the oxidation status of a quinone electron carrier. FEBS Journal, 2002, 269, 3479-3484.	0.2	63
30	Versatile roles of the chaperonin GroEL in microorganism-insect interactions. FEMS Microbiology Letters, 2014, 353, 1-10.	0.7	63
31	Replication of the Endosymbiotic Bacterium Blochmannia floridanus Is Correlated with the Developmental and Reproductive Stages of Its Ant Host. Applied and Environmental Microbiology, 2004, 70, 4096-4102.	1.4	62
32	In VivoCharacterization of the Unorthodox BvgS Two-component Sensor Protein ofBordetella pertussis. Journal of Molecular Biology, 1995, 248, 596-610.	2.0	61
33	Endosymbiont Tolerance and Control within Insect Hosts. Insects, 2012, 3, 553-572.	1.0	59
34	An engineered 3D human airway mucosa model based on an SIS scaffold. Biomaterials, 2014, 35, 7355-7362.	5.7	59
35	Pertussis toxin promoter sequences involved in modulation. Journal of Bacteriology, 1989, 171, 4026-4030.	1.0	57
36	Genetic and biochemical characterization of the aerobactin synthesis operon on pColV. Molecular Genetics and Genomics, 1984, 196, 74-80.	2.4	55

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37	Transcriptional profiling of the endosymbiont <i>Blochmannia floridanus</i> during different developmental stages of its holometabolous ant host. Environmental Microbiology, 2009, 11, 877-888.	1.8	47
38	Effect of mutations causing overexpression of RNA polymerase alpha subunit on regulation of virulence factors in Bordetella pertussis. Journal of Bacteriology, 1994, 176, 7267-7273.	1.0	45
39	A new gene locus of Bordetella pertussis defines a novel family of prokaryotic transcriptional accessory proteins. Journal of Bacteriology, 1996, 178, 4445-4452.	1.0	42
40	Transcriptional profiling of <i>Bordetella pertussis</i> reveals requirement of RNA chaperone Hfq for Type III secretion system functionality. RNA Biology, 2015, 12, 175-185.	1.5	42
41	Gene expression analysis of the endosymbiont-bearing midgut tissue during ontogeny of the carpenter ant Camponotus floridanus. Journal of Insect Physiology, 2013, 59, 611-623.	0.9	41
42	Phagosome Acidification Has Opposite Effects on Intracellular Survival of Bordetella pertussis and B. bronchiseptica. Infection and Immunity, 2000, 68, 7039-7048.	1.0	40
43	Phase variation affects long-term survival of Bordetella bronchiseptica in professional phagocytes. Infection and Immunity, 1997, 65, 3469-3473.	1.0	40
44	Genetics of pertussis toxin. Molecular Microbiology, 1989, 3, 119-124.	1.2	39
45	Cloning and characterization of an Mn-containing superoxide dismutase (SodA) of Bordetella pertussis. Journal of Bacteriology, 1997, 179, 2194-2201.	1.0	39
46	Genomic island excisions in Bordetella petrii. BMC Microbiology, 2009, 9, 141.	1.3	39
47	Microsatellites reveal clonal structure of populations of the thelytokous ant Platythyrea punctata (F. Smith) (Hymenoptera; Formicidae). Molecular Ecology, 1999, 8, 1497-1507.	2.0	37
48	Identification and characterization of two functional domains of the hemolysin translocator protein HlyD. Molecular Genetics and Genomics, 1994, 245, 203-211.	2.4	36
49	Resemblance and divergence: the "new―members of the genus Bordetella. Medical Microbiology and Immunology, 2010, 199, 155-163.	2.6	36
50	Immune response of the ant Camponotus floridanus against pathogens and its obligate mutualistic endosymbiont. Insect Biochemistry and Molecular Biology, 2011, 41, 529-536.	1.2	36
51	Scrutinizing the immune defence inventory of Camponotus floridanus applying total transcriptome sequencing. BMC Genomics, 2015, 16, 540.	1.2	33
52	A phase variant of Bordetella pertussis with a mutation in a new locus involved in the regulation of pertussis toxin and adenylate cyclase toxin expression. Journal of Bacteriology, 1993, 175, 6679-6688.	1.0	32
53	Characterization of a Bordetella pertussisDiaminopimelate (DAP) Biosynthesis Locus Identifies dapC, a Novel Gene Coding for anN-Succinyl-I,I-DAP Aminotransferase. Journal of Bacteriology, 2000, 182, 3626-3631.	1.0	30
54	New insights into symbiotic associations between ants and bacteria. Research in Microbiology, 2000, 151, 513-519.	1.0	30

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55	The BvgAS two-component system of Bordetella spp.: a versatile modulator of virulence gene expression. International Journal of Medical Microbiology, 2001, 291, 119-130.	1.5	30
56	Dimerization of signalling modules of the EvgAS and BvgAS phosphorelay systems. BBA - Proteins and Proteomics, 2000, 1478, 341-354.	2.1	25
57	Systemic gene knockdown in Camponotus floridanus workers by feeding of dsRNA. Insectes Sociaux, 2013, 60, 475-484.	0.7	25
58	Global regulatory mechanisms affect virulence gene expression in Bordetella pertussis. Molecular Genetics and Genomics, 1995, 247, 86-94.	2.4	24
59	Distribution of the obligate endosymbiont Blochmannia floridanus and expression analysis of putative immune genes in ovaries of the carpenter ant Camponotus floridanus. Arthropod Structure and Development, 2016, 45, 475-487.	0.8	24
60	Conserved sequence motifs in the unorthodox BvgS two-component sensor protein ofBordetella pertussis. Molecular Genetics and Genomics, 1996, 252, 169-176.	2.4	22
61	Molecular Characterization of Antimicrobial Peptide Genes of the Carpenter Ant Camponotus floridanus. PLoS ONE, 2012, 7, e43036.	1.1	21
62	Expression of bacterial cytotoxin genes in mammalian target cells. Molecular Microbiology, 1992, 6, 2651-2659.	1.2	18
63	Representational difference analysis identifies a strain-specific LPS biosynthesis locus in Bordetella spp Molecular Genetics and Genomics, 1999, 262, 189-198.	2.4	18
64	An antibiotic target ranking and prioritization pipeline combining sequence, structure and network-based approaches exemplified for Serratia marcescens. Gene, 2016, 591, 268-278.	1.0	17
65	Identification and regulation of expression of a gene encoding a filamentous hemagglutinin-related protein in Bordetella holmesii. BMC Microbiology, 2007, 7, 100.	1.3	14
66	Functional analysis of the pertussis toxin promoter. Research in Microbiology, 1992, 143, 671-681.	1.0	13
67	Promoter Characterization in the AT-Rich Genome of the Obligate Endosymbiont " <i>Candidatus</i> Blochmannia floridanusâ€# Journal of Bacteriology, 2009, 191, 3747-3751.	1.0	13
68	Identification and genomic organization of gene loci negatively controlled by the virulence regulatory BvgAS two-component system in Bordetella bronchiseptica. Molecular Genetics and Genomics, 2002, 267, 526-535.	1.0	11
69	Domain structure of the outer membrane transporter protein CyaE of Bordetella pertussis. Molecular Microbiology, 1995, 17, 1219-1220.	1.2	10
70	Rational design and molecular characterization of a chimaeric response regulator protein. Journal of Molecular Biology, 2001, 310, 283-290.	2.0	10
71	Development and characterization of attenuated metabolic mutants of <i>Bordetella bronchiseptica</i> ) for applications in vaccinology. Environmental Microbiology, 2013, 15, 64-76.	1.8	9
72	Structure-function relationships in the Bvg and Evg two-component phosphorelay systems. International Journal of Medical Microbiology, 2000, 290, 317-323.	1.5	8

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73	Molecular characterization of the BvgA response regulator of Bordetella holmesii. Microbiological Research, 2009, 164, 243-252.	2.5	8
74	Phg, a novel member of the autotransporter family present in Bordetella species. Microbiological Research, 2005, 160, 329-336.	2.5	5
75	JANE: efficient mapping of prokaryotic ESTs and variable length sequence reads on related template genomes. BMC Bioinformatics, 2009, 10, 391.	1.2	5