## Helen Billman-Jacobe

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/341414/publications.pdf

Version: 2024-02-01

70 papers 3,017 citations

30 h-index 52 g-index

72 all docs

72 docs citations

times ranked

72

3267 citing authors

#	Article	IF	CITATIONS
1	Antimicrobial stewardship in companion animal practice: an implementation trial in 135 general practice veterinary clinics. JAC-Antimicrobial Resistance, 2022, 4, dlac015.	2.1	10
2	Faecal microbiota and antimicrobial resistance gene profiles of healthy foals. Equine Veterinary Journal, 2021, 53, 806-816.	1.7	6
3	In-Water Antibiotic Dosing Practices on Pig Farms. Antibiotics, 2021, 10, 169.	3.7	12
4	Effect of Drinking Water Distribution System Design on Antimicrobial Delivery to Pigs. Animals, 2021, 11, 2362.	2.3	5
5	Infection of Slugs with Theronts of the Ciliate Protozoan, Tetrahymena rostrata. Microorganisms, 2021, 9, 1970.	3.6	4
6	Antimicrobial stewardship in Australia: the role of qualitative research in programme development. JAC-Antimicrobial Resistance, 2021, 3, dlab166.	2.1	8
7	Intraspecies Variation in Tetrahymena rostrata. Microorganisms, 2021, 9, 2100.	3.6	2
8	Water Distribution Systems in Pig Farm Buildings: Critical Elements of Design and Management. Animals, 2021, 11, 3268.	2.3	6
9	Colonization of a hand washing sink in a veterinary hospital by an Enterobacter hormaechei strain carrying multiple resistances to high importance antimicrobials. Antimicrobial Resistance and Infection Control, 2020, 9, 163.	4.1	13
10	Use of cefovecin in dogs and cats attending firstâ€opinion veterinary practices in Australia. Veterinary Record, 2020, 187, e95.	0.3	14
11	A practical guide for managing a self-sustaining colony of <i>Deroceras reticulatum</i> (Müller) (Mollusca: Pulmonata). Biocontrol Science and Technology, 2020, 30, 920-928.	1.3	1
12	Antibiotic Resistance Genes in Antibiotic-Free Chicken Farms. Antibiotics, 2020, 9, 120.	3.7	14
13	Impact of insertion sequences on convergent evolution of Shigella species. PLoS Genetics, 2020, 16, e1008931.	3.5	43
14	The mitochondrial genome of <i>Tetrahymena rostrata</i> . Mitochondrial DNA Part B: Resources, 2020, 5, 53-54.	0.4	4
15	Survey of veterinary prescribing for poultry disease. Australian Veterinary Journal, 2019, 97, 288-288.	1.1	3
16	Appraisal of the Australian Veterinary Prescribing Guidelines for antimicrobial prophylaxis for surgery in dogs and cats. Australian Veterinary Journal, 2019, 97, 316-322.	1.1	8
17	Review: Water medication of growing pigs: sources of between-animal variability in systemic exposure to antimicrobials. Animal, 2019, 13, 3031-3040.	3.3	18
18	Z/I1 Hybrid Virulence Plasmids Carrying Antimicrobial Resistance genes in S. Typhimurium from Australian Food Animal Production. Microorganisms, 2019, 7, 299.	3.6	7

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19	Exploration of antibiotic resistance risks in a veterinary teaching hospital with Oxford Nanopore long read sequencing. PLoS ONE, 2019, 14, e0217600.	2.5	15
20	Antimicrobial dosing for common equine drugs: a content review and practical advice for veterinarians in Australia. Australian Veterinary Journal, 2019, 97, 103-107.	1.1	10
21	Global phylogenomics of multidrug-resistant Salmonella enterica serotype Kentucky ST198. Microbial Genomics, 2019, 5, .	2.0	69
22	Barriers to and enablers of implementing antimicrobial stewardship programs in veterinary practices. Journal of Veterinary Internal Medicine, 2018, 32, 1092-1099.	1.6	77
23	Antimicrobial susceptibility testing by Australian veterinary diagnostic laboratories. Australian Veterinary Journal, 2018, 96, 142-146.	1.1	8
24	pSTM6-275, a Conjugative IncHI2 Plasmid of Salmonella enterica That Confers Antibiotic and Heavy-Metal Resistance under Changing Physiological Conditions. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	44
25	Antimicrobials used for surgical prophylaxis by equine veterinary practitioners in Australia. Equine Veterinary Journal, 2018, 50, 65-72.	1.7	23
26	Population wide assessment of antimicrobial use in dogs and cats using a novel data source – A cohort study using pet insurance data. Veterinary Microbiology, 2018, 225, 34-39.	1.9	40
27	Antimicrobial labelling in Australia: a threat to antimicrobial stewardship?. Australian Veterinary Journal, 2018, 96, 151-154.	1.1	23
28	Longitudinal study of Salmonella 1,4,[5],12:i:- shedding in five Australian pig herds. Preventive Veterinary Medicine, 2017, 136, 19-28.	1.9	16
29	Antimicrobials used for surgical prophylaxis by companion animal veterinarians in Australia. Veterinary Microbiology, 2017, 203, 301-307.	1.9	25
30	Crossâ€sectional study of antimicrobials used for surgical prophylaxis by bovine veterinary practitioners in Australia. Veterinary Record, 2017, 181, 426-426.	0.3	14
31	Genome Sequence of an Australian Monophasic Salmonella enterica subsp. enterica Typhimurium Isolate (TW-Stm6) Carrying a Large Plasmid with Multiple Antimicrobial Resistance Genes. Genome Announcements, 2017, 5, .	0.8	31
32	Meeting the Capstone Challenge in Postgraduate Food Science Education. Journal of Food Science Education, 2017, 16, 77-80.	1.0	2
33	ISMapper: identifying transposase insertion sites in bacterial genomes from short read sequence data. BMC Genomics, 2015, 16, 667.	2.8	119
34	Evidence of microevolution of Salmonella Typhimurium during a series of egg-associated outbreaks linked to a single chicken farm. BMC Genomics, 2013, 14, 800.	2.8	67
35	Identification of a Novel Gene Product That Promotes Survival of Mycobacterium smegmatis in Macrophages. PLoS ONE, 2012, 7, e31788.	2.5	14
36	Stress-induced Synthesis of Phosphatidylinositol 3-Phosphate in Mycobacteria. Journal of Biological Chemistry, 2010, 285, 16643-16650.	3.4	22

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37	Investigating the Function of the Putative Mycolic Acid Methyltransferase UmaA. Journal of Biological Chemistry, 2008, 283, 1419-1427.	3.4	33
38	Mutations in $\langle i \rangle$ pimE $\langle  i \rangle$ Restore Lipoarabinomannan Synthesis and Growth in a $\langle i \rangle$ Mycobacterium smegmatis lpqW $\langle  i \rangle$ Mutant. Journal of Bacteriology, 2008, 190, 3690-3699.	2.2	38
39	Population Genetics Study of Isoniazid Resistance Mutations and Evolution of Multidrug-Resistant Mycobacterium tuberculosis. Antimicrobial Agents and Chemotherapy, 2006, 50, 2640-2649.	3.2	364
40	PimE Is a Polyprenol-phosphate-mannose-dependent Mannosyltransferase That Transfers the Fifth Mannose of Phosphatidylinositol Mannoside in Mycobacteria. Journal of Biological Chemistry, 2006, 281, 25143-25155.	3.4	118
41	Identification of a Novel Protein with a Role in Lipoarabinomannan Biosynthesis in Mycobacteria. Journal of Biological Chemistry, 2006, 281, 9011-9017.	3.4	63
42	Function of Phosphatidylinositol in Mycobacteria. Journal of Biological Chemistry, 2005, 280, 10981-10987.	3.4	58
43	Role of <i>embB</i> Codon 306 Mutations in <i>Mycobacterium tuberculosis</i> Revisited: a Novel Association with Broad Drug Resistance and IS <i>6110</i> Clustering Rather than Ethambutol Resistance. Antimicrobial Agents and Chemotherapy, 2005, 49, 3794-3802.	3.2	103
44	Compartmentalization of Lipid Biosynthesis in Mycobacteria. Journal of Biological Chemistry, 2005, 280, 21645-21652.	3.4	92
45	Molecular Characterization of Isoniazid-Resistant Mycobacterium tuberculosis Isolates Collected in Australia. Antimicrobial Agents and Chemotherapy, 2005, 49, 4068-4074.	3.2	52
46	Methylation of GPLs in Mycobacterium smegmatis and Mycobacterium avium. Journal of Bacteriology, 2004, 186, 6792-6799.	2.2	33
47	A novel IS element, ISMpa1, in Mycobacterium avium subsp. paratuberculosis. Veterinary Microbiology, 2004, 98, 297-306.	1.9	13
48	Biosynthesis of mycobacterial phosphatidylinositol mannosides. Biochemical Journal, 2004, 378, 589-597.	3.7	93
49	Inactivation of mshB, a key gene in the mycothiol biosynthesis pathway in Mycobacterium smegmatis. Microbiology (United Kingdom), 2003, 149, 1341-1349.	1.8	61
50	Mannose metabolism is required for mycobacterial growth. Biochemical Journal, 2003, 372, 77-86.	3.7	59
51	Identification of a peptide synthetase involved in the biosynthesis of glycopeptidolipids of Mycobacterium smegmatis. Molecular Microbiology, 2002, 33, 1244-1253.	2.5	101
52	Modification of glycopeptidolipids by an O-methyltransferase of Mycobacterium smegmatis a aThe GenBank accession number for the sequence determined in this work is AY138899 Microbiology (United Kingdom), 2002, 148, 3079-3087.	1.8	44
53	The impact of the absence of glycopeptidolipids on the ultrastructure, cell surface and cell wall properties, and phagocytosis of Mycobacterium smegmatis. Microbiology (United Kingdom), 2002, 148, 3089-3100.	1.8	116
54	Antibody Responses to Infections with Strains of Plasmodium falciparum Expressing Diverse Forms of Merozoite Surface Protein 2. Infection and Immunity, 2001, 69, 959-967.	2.2	35

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55	Quantitative Determination of the Biodegradable Polymer Poly( $\hat{l}^2$ -hydroxybutyrate) in a Recombinant Escherichia coli Strain by Use of Mid-Infrared Spectroscopy and Multivariative Statistics. Applied and Environmental Microbiology, 2000, 66, 3415-3420.	3.1	128
56	Identification of a Methyltransferase from Mycobacterium smegmatis Involved in Glycopeptidolipid Synthesis. Journal of Biological Chemistry, 2000, 275, 24900-24906.	3.4	61
57	Characterization of a <i>Mycobacterium smegmatis</i> Mutant Lacking Penicillin Binding Protein 1. Antimicrobial Agents and Chemotherapy, 1999, 43, 3011-3013.	3.2	29
58	A Plasmodium falciparum apical membrane antigen-1 (AMA-1) gene apparently generated by intragenic recombination. Molecular and Biochemical Parasitology, 1999, 100, 243-246.	1.1	15
59	Temporal Variation of the Merozoite Surface Protein-2 Gene of <i>Plasmodium falciparum </i> Infection and Immunity, 1998, 66, 239-246.	2.2	45
60	Expression in bacteria other than Escherichia coli. Current Opinion in Biotechnology, 1996, 7, 500-504.	6.6	25
61	Analysis of isoniazid-resistant transposon mutants of Mycobacterium smegmatis. FEMS Microbiology Letters, 1996, 144, 47-52.	1.8	18
62	Expression and secretion of heterologous proteases by Corynebacterium glutamicum. Applied and Environmental Microbiology, 1995, 61, 1610-1613.	3.1	52
63	Differential T Cell Responses To Mycobacteria-Secreted Proteins Distinguish Vaccination With Bacille Calmette-Guerin From Infection With Mycobacterium Tuberculosis. Journal of Infectious Diseases, 1994, 170, 1326-1330.	4.0	78
64	Diagnosis and epidemiology of bovine tuberculosis using molecular biological approaches. Veterinary Microbiology, 1994, 40, 83-94.	1.9	40
65	Nucleotide sequence of arecAgene fromCorny bacterium glutamicum. DNA Sequence, 1994, 4, 403-404.	0.7	2
66	Expression of ovine gamma interferon in Escherichia coli and Corynebacterium glutamicum. Applied and Environmental Microbiology, 1994, 60, 1641-1645.	3.1	27
67	T-cell determinants and antibody binding sites on the major mycobacterial secretory protein MPB59 of Mycobacterium bovis. Infection and Immunity, 1994, 62, 5319-5326.	2.2	66
68	A comparison of the interferon gamma assay with the absorbed ELISA for the diagnosis of Johne's disease in cattle. Australian Veterinary Journal, 1992, 69, 25-28.	1.1	72
69	Mapping of the T and B cell epitopes of the <i>Mycobacterium bovis</i> protein, MPB70. Immunology and Cell Biology, 1990, 68, 359-365.	2.3	30
70	Epitope mapping of the Mycobacterium bovis secretory protein MPB70 using overlapping peptide analysis. Journal of General Microbiology, 1990, 136, 265-272.	2.3	54