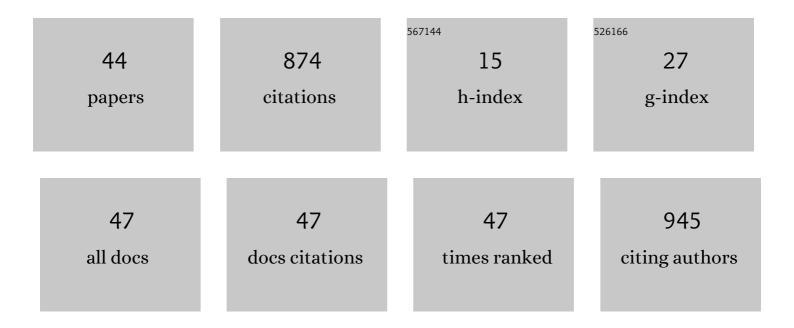
Conny Turni

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Nucleic acid purification from plants, animals and microbes in under 30 seconds. PLoS Biology, 2017, 15, e2003916.	2.6	190
2	Development of a Rapid Multiplex PCR Assay To Genotype Pasteurella multocida Strains by Use of the Lipopolysaccharide Outer Core Biosynthesis Locus. Journal of Clinical Microbiology, 2015, 53, 477-485.	1.8	89
3	Antimicrobial resistance in bacteria associated with porcine respiratory disease in Australia. Veterinary Microbiology, 2014, 171, 232-235.	0.8	62
4	Pasteurella multocida Heddleston Serovar 3 and 4 Strains Share a Common Lipopolysaccharide Biosynthesis Locus but Display both Inter- and Intrastrain Lipopolysaccharide Heterogeneity. Journal of Bacteriology, 2013, 195, 4854-4864.	1.0	37
5	Evaluation of a multiplex PCR to identify and serotype <i>Actinobacillus pleuropneumoniae</i> serovars 1, 5, 7, 12 and 15. Letters in Applied Microbiology, 2014, 59, 362-369.	1.0	34
6	Antimicrobial resistance genes in <i>Actinobacillus pleuropneumoniae</i> , <i>Haemophilus parasuis</i> and <i>Pasteurella multocida</i> isolated from Australian pigs. Australian Veterinary Journal, 2016, 94, 227-231.	0.5	33
7	Towards a Standardized Method for Broth Microdilution Susceptibility Testing of Haemophilus parasuis. Journal of Clinical Microbiology, 2017, 55, 264-273.	1.8	33
8	Epidemiology of Fowl Cholera in Free Range Broilers. Avian Diseases, 2014, 58, 124-128.	0.4	32
9	Use of a proposed antimicrobial susceptibility testing method for Haemophilus parasuis. Veterinary Microbiology, 2014, 172, 586-589.	0.8	28
10	Protective efficacy afforded by live Pasteurella multocida vaccines in chickens is independent of lipopolysaccharide outer core structure. Vaccine, 2016, 34, 1696-1703.	1.7	25
11	Studies on the presence and persistence of <i>Pasteurella multocida</i> serovars and genotypes in fowl cholera outbreaks. Avian Pathology, 2013, 42, 581-585.	0.8	24
12	Genotypic diversity of <scp><i>Pasteurella multocida</i></scp> isolates from pigs and poultry in Australia. Australian Veterinary Journal, 2018, 96, 390-394.	0.5	23
13	An evaluation of the apxIVA based PCR-REA method for differentiation of Actinobacillus pleuropneumoniae. Veterinary Microbiology, 2007, 121, 163-169.	0.8	22
14	Interrogating the bovine reproductive tract metagenomes using culture-independent approaches: a systematic review. Animal Microbiome, 2021, 3, 41.	1.5	22
15	Genomic analysis of phylogenetic group B2 extraintestinal pathogenic E. coli causing infections in dogs in Australia. Veterinary Microbiology, 2020, 248, 108783.	0.8	20
16	Glaesserella australis sp. nov., isolated from the lungs of pigs. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 3686-3692.	0.8	15
17	Virulenceâ€associated gene profiling, DNA fingerprinting and multilocus sequence typing of <i>Haemophilus parasuis</i> isolates in Australia. Australian Veterinary Journal, 2018, 96, 196-202.	0.5	14
18	Actinobacillus pleuropneumoniae: The molecular determinants of virulence and pathogenesis. Advances in Microbial Physiology, 2021, 78, 179-216.	1.0	14

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19	Genetic diversity and toxin gene distribution among serovars of <i>Actinobacillus pleuropneumoniae</i> from Australian pigs. Australian Veterinary Journal, 2018, 96, 17-23.	0.5	13
20	Combining conventional and participatory approaches to identify and prioritise management and health-related constraints to smallholder pig production in San Simon, Pampanga, Philippines. Preventive Veterinary Medicine, 2020, 178, 104987.	0.7	13
21	Genomic analysis of fluoroquinolone-susceptible phylogenetic group B2 extraintestinal pathogenic Escherichia coli causing infections in cats. Veterinary Microbiology, 2020, 245, 108685.	0.8	12
22	Whole Genome Sequence Analysis of Pig Respiratory Bacterial Pathogens with Elevated Minimum Inhibitory Concentrations for Macrolides. Microbial Drug Resistance, 2016, 22, 531-537.	0.9	11
23	First Emergence of Resistance to Macrolides and Tetracycline Identified in Mannheimia haemolytica and Pasteurella multocida Isolates from Beef Feedlots in Australia. Microorganisms, 2021, 9, 1322.	1.6	11
24	Technical note: overcoming host contamination in bovine vaginal metagenomic samples with nanopore adaptive sequencing. Journal of Animal Science, 2022, 100, .	0.2	10
25	Human Wound Infection with Mannheimia glucosida following Lamb Bite. Journal of Clinical Microbiology, 2015, 53, 3374-3376.	1.8	8
26	Application of an enzyme-linked immunosorbent assay for detection of antibodies to <i>Actinobacillus pleuropneumoniae</i> serovar 15 in pig sera. Journal of Veterinary Medical Science, 2017, 79, 1968-1972.	0.3	8
27	Molecular and serological characterization of <i>Riemerella</i> isolates associated with poultry in Australia. Avian Pathology, 2021, 50, 31-40.	0.8	8
28	Genetic analysis of porcine circovirus type 2 (PCV2) in Queensland, Australia. Australian Veterinary Journal, 2020, 98, 388-395.	0.5	7
29	Pathogens associated with pleuritic pig lungs at an abattoir in Queensland Australia. Australian Veterinary Journal, 2021, 99, 163-171.	0.5	7
30	An Unusual Strain of Haemophilus Parasuis that Fails to React in a Species-Specific Polymerase Chain Reaction Assay. Journal of Veterinary Diagnostic Investigation, 2011, 23, 355-358.	0.5	6
31	Novel insights into pasteurellosis in captive pinnipeds. Veterinary Microbiology, 2019, 231, 232-237.	0.8	6
32	Phase variation in latB associated with a fatal Pasteurella multocida outbreak in captive squirrel gliders. Veterinary Microbiology, 2020, 243, 108612.	0.8	6
33	Using genomics to understand inter- and intra- outbreak diversity of Pasteurella multocida isolates associated with fowl cholera in meat chickens. Microbial Genomics, 2020, 6, .	1.0	6
34	Variation in the Antimicrobial Susceptibility ofActinobacillus pleuropneumoniaelsolates in a Pig, Within a Batch of Pigs, and Among Batches of Pigs from One Farm. Microbial Drug Resistance, 2015, 21, 491-496.	0.9	4
35	Diverse strains of Actinobacillus lignieresii isolated from clinically affected cattle in a geographically restricted area. Australian Veterinary Journal, 2019, 97, 440-446.	0.5	4
36	Detection of porcine circovirus type 2 (PCV2) in the Philippines and the complexity of PCV2-associated disease diagnosis. Tropical Animal Health and Production, 2021, 53, 371.	0.5	3

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#	Article	IF	CITATIONS
37	Latent class analysis identifies multimorbidity patterns in pigs with respiratory disease. Preventive Veterinary Medicine, 2021, 186, 105209.	0.7	2
38	Antimicrobial susceptibility, plasmid replicon typing, phylogenetic grouping, and virulence potential of avian pathogenic and faecal <i>Escherichia coli</i> isolated from meat chickens in Australia. Avian Pathology, 2022, 51, 349-360.	0.8	2
39	Risk Factors Associated with the Carriage of Pathogenic Escherichia coli in Healthy Commercial Meat Chickens in Queensland, Australia â€. Poultry, 2022, 1, 94-110.	0.5	2
40	Role of Staphylococcus agnetis and Staphylococcus hyicus in the Pathogenesis of Buffalo Fly Skin Lesions in Cattle. Microbiology Spectrum, 2022, 10, .	1.2	2
41	MOLECULAR IDENTIFICATION OF MEMBERS OF THE FAMILY PASTEURELLACEAE FROM THE ORAL CAVITY OF KOALAS (PHASCOLARCTOS CINEREUS) AND THEIR RELATIONSHIP WITH ISOLATES FROM KOALA BITE WOUNDS IN HUMANS. Journal of Zoo and Wildlife Medicine, 2021, 51, 771-779.	0.3	1
42	Development of a Luminex microbead-based serotyping assay for Glaesserella parasuis. Journal of Microbiological Methods, 2021, 182, 106159.	0.7	1
43	Phase variation in the glycosyltransferase genes of Pasteurella multocida associated with outbreaks of fowl cholera on free-range layer farms. Microbial Genomics, 2022, 8, .	1.0	1
44	An improved multiplex PCR for Actinobacillus pleuropneumoniae, Glaesserella australis and Pasteurella multocida. Journal of Microbiological Methods, 2021, 191, 106360.	0.7	0