Philippe Fravalo

List of Publications by Year in descending order

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25 papers	586 citations	686830 13 h-index	610482 24 g-index
25 all docs	25 docs citations	25 times ranked	783 citing authors

#	Article	IF	Citations
1	Chicken Caecal Microbiome Modifications Induced by Campylobacter jejuni Colonization and by a Non-Antibiotic Feed Additive. PLoS ONE, 2015, 10, e0131978.	1.1	123
2	Risk factors for Salmonella enterica subsp. enterica shedding by market-age pigs in French farrow-to-finish herds. Preventive Veterinary Medicine, 2004, 63, 103-120.	0.7	118
3	Prevalence of colistin resistance and mcr-1/mcr-2 genes in extended-spectrum Î²-lactamase/AmpC-producing Escherichia coli isolated from chickens in Canada, Senegal and Vietnam. Journal of Global Antimicrobial Resistance, 2019, 19, 222-227.	0.9	35
4	Source Attribution of Foodborne Diseases: Potentialities, Hurdles, and Future Expectations. Frontiers in Microbiology, 2018, 9, 1983.	1.5	32
5	Analysis of Listeria monocytogenes Strain Distribution in a Pork Slaughter and Cutting Plant in the Province of Quebec. Journal of Food Protection, 2014, 77, 2121-2128.	0.8	29
6	Critical Orientation in the Jungle of Currently Available Methods and Types of Data for Source Attribution of Foodborne Diseases. Frontiers in Microbiology, 2019, 10, 2578.	1.5	26
7	Lack of Evidence That Selenium-Yeast Improves Chicken Health and Modulates the Caecal Microbiota in the Context of Colonization by Campylobacter jejuni. Frontiers in Microbiology, 2017, 8, 451.	1.5	24
8	Impact of medicated feed along with clay mineral supplementation on Escherichia coli resistance to antimicrobial agents in pigs after weaning in field conditions. Research in Veterinary Science, 2015, 102, 72-79.	0.9	22
9	Detection and Phylogenetic Analysis of the Hepatitis E Virus in a Canadian Swine Production Network. Food and Environmental Virology, 2016, 8, 296-304.	1.5	22
10	Extensive characterization of Campylobacter jejuni chicken isolates to uncover genes involved in the ability to compete for gut colonization. BMC Microbiology, 2015, 15, 97.	1.3	21
11	Reduction of Salmonella Shedding by Sows during Gestation in Relation to Its Fecal Microbiome. Frontiers in Microbiology, 2017, 8, 2219.	1.5	17
12	Salmonella shedding status of the sow affects the microbiota of their piglets at weaning. Journal of Applied Microbiology, 2019, 126, 411-423.	1.4	16
13	Screening for fecal presence of colistin-resistant Escherichia coli and mcr-1 and mcr-2 genes in camel-calves in southern Tunisia. Acta Veterinaria Scandinavica, 2018, 60, 35.	0.5	15
14	Evolution of Pig Fecal Microbiota Composition and Diversity in Response to Enterotoxigenic Escherichia coli Infection and Colistin Treatment in Weaned Piglets. Microorganisms, 2021, 9, 1459.	1.6	14
15	First identification of mcr-1/mcr-2 genes in the fecal microbiota of Canadian commercial pigs during the growing and finishing period . Veterinary Medicine: Research and Reports, 2019, Volume 10, 65-67.	0.4	13
16	Characterisation of InlA truncation in Listeria monocytogenes isolates from farm animals and human cases in the province of Quebec. Veterinary Record Open, 2017, 4, e000199.	0.3	10
17	Persistence of Indicator and Pathogenic Microorganisms in Broccoli following Manure Spreading and Irrigation with Fecally Contaminated Water: Field Experiment. Journal of Food Protection, 2015, 78, 1776-1784.	0.8	9
18	Dynamics of Virus Distribution in a Defined Swine Production Network Using Enteric Viruses as Molecular Markers. Applied and Environmental Microbiology, 2017, 83, .	1.4	9

#	Article	IF	Citations
19	Comparison of microbiota of recycled manure solids and straw bedding used in dairy farms in eastern Canada. Journal of Dairy Science, 2022, 105, 389-408.	1.4	9
20	The swine enteric virome in a commercial production system and its association with neonatal diarrhea. Veterinary Microbiology, 2022, 266, 109366.	0.8	9
21	Feed presentation options in Swine early fattening mitigatesSalmonellashedding and specifically modulates the faecal microbiota. Journal of Applied Microbiology, 2017, 122, 30-39.	1.4	7
22	Treatments of porcine fecal samples affect high-throughput virome sequencing results. Journal of Virological Methods, 2021, 289, 114045.	1.0	4
23	Sows affect their pigletsâ€™ faecal microbiota until fattening but not their Salmonella enterica shedding status. Letters in Applied Microbiology, 2021, 72, 113-120.	1.0	1
24	Different types of stainless steel used in equipment in meat plants do not affect the initial microbial transfer, including pathogens, from pork skin. Canadian Journal of Veterinary Research, 2015, 79, 255-9.	0.2	1
25	Salmonella contamination in a network of 10 pig farms interconnected within the same cooperative. Veterinary Record Open, 2019, 6, e000269.	0.3	0