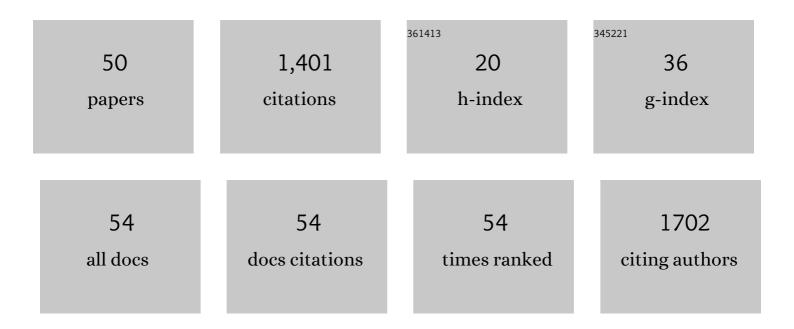
Laura C Miller

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptomic Analysis of Liver Indicates Novel Vaccine to Porcine Reproductive and Respiratory Virus Promotes Homeostasis in T-Cell and Inflammatory Immune Responses Compared to a Commercial Vaccine in Pigs. Frontiers in Veterinary Science, 2022, 9, 791034.	2.2	0
2	Epigenetic Evolution of ACE2 and IL-6 Genes: Non-Canonical Interferon-Stimulated Genes Correlate to COVID-19 Susceptibility in Vertebrates. Genes, 2021, 12, 154.	2.4	31
3	Inhibition of Antiviral Innate Immunity by Foot-and-Mouth Disease Virus L ^{pro} through Interaction with the N-Terminal Domain of Swine RNase L. Journal of Virology, 2021, 95, e0036121.	3.4	6
4	Harness Organoid Models for Virological Studies in Animals: A Cross-Species Perspective. Frontiers in Microbiology, 2021, 12, 725074.	3.5	5
5	Immunometabolic Dysregulation at the Intersection of Obesity and COVID-19. Frontiers in Immunology, 2021, 12, 732913.	4.8	16
6	Impact of Porcine Arterivirus, Influenza B, and Their Coinfection on Antiviral Response in the Porcine Lung. Pathogens, 2020, 9, 934.	2.8	8
7	Integrate structural analysis, isoform diversity, and interferon-inductive propensity of ACE2 to predict SARS-CoV2 susceptibility in vertebrates. Heliyon, 2020, 6, e04818.	3.2	13
8	Comparison of the Transcriptome Response within the Swine Tracheobronchial Lymphnode Following Infection with PRRSV, PCV-2 or IAV-S Pathogens, 2020, 9, 99.	2.8	7
9	Differentially Expressed MiRNAs and tRNA Genes Affect Host Homeostasis During Highly Pathogenic Porcine Reproductive and Respiratory Syndrome Virus Infections in Young Pigs. Frontiers in Genetics, 2019, 10, 691.	2.3	15
10	Cross-Species Genome-Wide Analysis Reveals Molecular and Functional Diversity of the Unconventional Interferon-ω Subtype. Frontiers in Immunology, 2019, 10, 1431.	4.8	28
11	Identification of small non-coding RNA classes expressed in swine whole blood during HP-PRRSV infection. Virology, 2018, 517, 56-61.	2.4	13
12	Porcine reproductive and respiratory disease virus: Evolution and recombination yields distinct ORF5 RFLP 1-7-4 viruses with individual pathogenicity. Virology, 2018, 513, 168-179.	2.4	75
13	Identification of Coding and Non-coding RNA Classes Expressed in Swine Whole Blood. Journal of Visualized Experiments, 2018, , .	0.3	Ο
14	Against the Odds: Syringe Exchange Policy Implementation in Indiana. AIDS and Behavior, 2017, 21, 973-981.	2.7	14
15	Reduction of infection by inhibiting mTOR pathway is associated with reversed repression of type I interferon by porcine reproductive and respiratory syndrome virus. Journal of General Virology, 2017, 98, 1316-1328.	2.9	20
16	Comparative analysis of signature genes in PRRSV-infected porcine monocyte-derived cells to different stimuli. PLoS ONE, 2017, 12, e0181256.	2.5	13
17	It's Not Just a Yes or No Answer: Expressions of Local Health Department Accreditation. Frontiers in Public Health, 2016, 4, 21.	2.7	6
18	Enhancement of innate immunity with granulocyte colony-stimulating factor did not mitigate disease in pigs infected with a highly pathogenic Chinese PRRSV strain. Veterinary Immunology and Immunopathology, 2016, 179, 70-76.	1.2	3

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19	Susceptibility of swine to H5 and H7 low pathogenic avian influenza viruses. Influenza and Other Respiratory Viruses, 2016, 10, 346-352.	3.4	17
20	Status of vaccines for porcine epidemic diarrhea virus in the United States and Canada. Virus Research, 2016, 226, 108-116.	2.2	65
21	Leading edge analysis of transcriptomic changes during pseudorabies virus infection. Genomics Data, 2016, 10, 104-106.	1.3	7
22	Evaluation of two real-time polymerase chain reaction assays for <i>Porcine epidemic diarrhea virus</i> (PEDV) to assess PEDV transmission in growing pigs. Journal of Veterinary Diagnostic Investigation, 2016, 28, 20-29.	1.1	20
23	Effects of Pseudorabies Virus Infection on the Tracheobronchial Lymph Node Transcriptome. Bioinformatics and Biology Insights, 2015, 9s2, BBI.S30522.	2.0	4
24	Macrophage Polarization in Virus-Host Interactions. Journal of Clinical & Cellular Immunology, 2015, 06, .	1.5	73
25	Evaluation of porcine epidemic diarrhea virus transmission and the immune response in growing pigs. Veterinary Research, 2015, 46, 49.	3.0	44
26	Efficacy of Type 2 PRRSV vaccine against Chinese and Vietnamese HP-PRRSV challenge in pigs. Vaccine, 2014, 32, 6457-6462.	3.8	33
27	Dual purpose with dual benefit research models in veterinary and biomedical research. Veterinary Immunology and Immunopathology, 2014, 159, 111-112.	1.2	1
28	Evolutionary characterization of pig interferon-inducible transmembrane gene family and member expression dynamics in tracheobronchial lymph nodes of pigs infected with swine respiratory disease viruses. Veterinary Immunology and Immunopathology, 2014, 159, 180-191.	1.2	19
29	Chinese and Vietnamese strains of HP-PRRSV cause different pathogenic outcomes in United States high health swine. Virology, 2013, 446, 238-250.	2.4	26
30	Experimental infection of United States swine with a Chinese highly pathogenic strain of porcine reproductive and respiratory syndrome virus. Virology, 2013, 435, 372-384.	2.4	98
31	Development and characterization of two porcine monocyte-derived macrophage cell lines. Results in Immunology, 2013, 3, 26-32.	2.2	15
32	Theme Issue on Asia Knowledge: Inside and Outside the Ivory Tower. Journal of American-East Asian Relations, 2013, 20, 311-323.	0.1	0
33	Reactomes of Porcine Alveolar Macrophages Infected with Porcine Reproductive and Respiratory Syndrome Virus. PLoS ONE, 2013, 8, e59229.	2.5	33
34	The Presence of Alpha Interferon at the Time of Infection Alters the Innate and Adaptive Immune Responses to Porcine Reproductive and Respiratory Syndrome Virus. Vaccine Journal, 2012, 19, 508-514.	3.1	34
35	Genomic sequence and virulence comparison of four Type 2 porcine reproductive and respiratory syndrome virus strains. Virus Research, 2012, 169, 212-221.	2.2	128
36	Evaluation of a real-time polymerase chain reaction assay for Pseudorabies virus surveillance purposes. Journal of Veterinary Diagnostic Investigation, 2012, 24, 739-745.	1.1	21

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37	Analysis of the swine tracheobronchial lymph node transcriptomic response to infection with a Chinese highly pathogenic strain of porcine reproductive and respiratory syndrome virus. BMC Veterinary Research, 2012, 8, 208.	1.9	30
38	Utility of a Panviral Microarray for Detection of Swine Respiratory Viruses in Clinical Samples. Journal of Clinical Microbiology, 2011, 49, 1542-1548.	3.9	21
39	In-Depth Global Analysis of Transcript Abundance Levels in Porcine Alveolar Macrophages Following Infection with Porcine Reproductive and Respiratory Syndrome Virus. Advances in Virology, 2010, 2010, 1-12.	1.1	12
40	Cytokine Protein Expression Levels in Tracheobronchial Lymph Node Homogenates of Pigs Infected with Pseudorabies Virus. Vaccine Journal, 2010, 17, 728-734.	3.1	15
41	Role of Toll-Like Receptors in Activation of Porcine Alveolar Macrophages by Porcine Reproductive and Respiratory Syndrome Virus. Vaccine Journal, 2009, 16, 360-365.	3.1	42
42	Development of Cell Lines from the Sheep Used to Construct the CHORI-243 Ovine BAC Library. Animal Biotechnology, 2008, 19, 84-88.	1.5	2
43	Effect of Porcine Reproductive and Respiratory Syndrome Virus on Porcine Alveolar Macrophage Function as Determined Using Serial Analysis of Gene Expression (SAGE). Developments in Biologicals, 2008, 132, 169-174.	0.5	7
44	Structure of Foot-and-mouth disease virus serotype A1061 alone and complexed with oligosaccharide receptor: receptor conservation in the face of antigenic variation. Journal of General Virology, 2005, 86, 1909-1920.	2.9	95
45	Evaluation of a Real-Time PCR Kit for Detecting Escherichia coli O157 in Bovine Fecal Samples. Applied and Environmental Microbiology, 2004, 70, 1855-1857.	3.1	25
46	Interferon type I response in porcine reproductive and respiratory syndrome virus-infected MARC-145 cells. Archives of Virology, 2004, 149, 2453-2463.	2.1	101
47	Gene expression profiling of bovine macrophages in response to O157:H7 lipopolysaccharide. Developmental and Comparative Immunology, 2004, 28, 635-645.	2.3	25
48	Apoptosis and porcine reproductive and respiratory syndrome virus. Veterinary Immunology and Immunopathology, 2004, 102, 131-142.	1.2	62
49	Role of the Cytoplasmic Domain of the β-Subunit of Integrin αvβ6 in Infection by Foot-and-Mouth Disease Virus. Journal of Virology, 2001, 75, 4158-4164.	3.4	43
50	Epigenetic Evolution of ACE2 and IL-6 Genes as Non-Canonical Interferon-Stimulated Genes Correlate to COVID-19 Susceptibility in Vertebrates. SSRN Electronic Journal, 0, , .	0.4	3