

Scott Kenney

List of Publications by Year in descending order

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43
papers

2,142
citations

331670

21
h-index

276875

41
g-index

45
all docs

45
docs citations

45
times ranked

2883
citing authors

#	ARTICLE	IF	CITATIONS
1	Porcine Deltacoronaviruses: Origin, Evolution, Cross-Species Transmission and Zoonotic Potential. <i>Pathogens</i> , 2022, 11, 79.	2.8	23
2	Characterization of the Cross-Species Transmission Potential for Porcine Deltacoronaviruses Expressing Sparrow Coronavirus Spike Protein in Commercial Poultry. <i>Viruses</i> , 2022, 14, 1225.	3.3	2
3	Differing coronavirus genres alter shared host signaling pathways upon viral infection. <i>Scientific Reports</i> , 2022, 12, .	3.3	1
4	Ectopic Expression of Genotype 1 Hepatitis E Virus ORF4 Increases Genotype 3 HEV Viral Replication in Cell Culture. <i>Viruses</i> , 2021, 13, 75.	3.3	30
5	Comparative Transcriptome Profiling of Human and Pig Intestinal Epithelial Cells after Porcine Deltacoronavirus Infection. <i>Viruses</i> , 2021, 13, 292.	3.3	14
6	Hepatitis E Virus Immunopathogenesis. <i>Pathogens</i> , 2021, 10, 1180.	2.8	12
7	Naturally Occurring Animal Coronaviruses as Models for Studying Highly Pathogenic Human Coronaviral Disease. <i>Veterinary Pathology</i> , 2021, 58, 438-452.	1.7	30
8	Dissecting the potential role of hepatitis E virus ORF1 nonstructural gene in crossâ€species infection by using intergenotypic chimeric viruses. <i>Journal of Medical Virology</i> , 2020, 92, 3563-3571.	5.0	7
9	Porcine Deltacoronavirus Infection and Transmission in Poultry, United States1. <i>Emerging Infectious Diseases</i> , 2020, 26, 255-265.	4.3	99
10	The COVID-19 Pandemic: A Comprehensive Review of Taxonomy, Genetics, Epidemiology, Diagnosis, Treatment, and Control. <i>Journal of Clinical Medicine</i> , 2020, 9, 1225.	2.4	480
11	Deltacoronavirus Evolution and Transmission: Current Scenario and Evolutionary Perspectives. <i>Frontiers in Veterinary Science</i> , 2020, 7, 626785.	2.2	19
12	Isolation and Tissue Culture Adaptation of Porcine Deltacoronavirus: A Case Study. <i>Methods in Molecular Biology</i> , 2020, 2203, 77-88.	0.9	3
13	CD8⁺ lymphocytes but not B lymphocytes are required for protection against chronic hepatitis E virus infection in chickens. <i>Journal of Medical Virology</i> , 2019, 91, 1960-1969.	5.0	4
14	The Current Host Range of Hepatitis E Viruses. <i>Viruses</i> , 2019, 11, 452.	3.3	69
15	Emerging and re-emerging coronaviruses in pigs. <i>Current Opinion in Virology</i> , 2019, 34, 39-49.	5.4	276
16	Evidence for an unknown agent antigenically related to the hepatitis E virus in dairy cows in the United States. <i>Journal of Medical Virology</i> , 2019, 91, 677-686.	5.0	23
17	Hepatitis E Virus: Animal Models and Zoonosis. <i>Annual Review of Animal Biosciences</i> , 2019, 7, 427-448.	7.4	24
18	Hepatitis E Virus Genome Structure and Replication Strategy. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2019, 9, a031724.	6.2	101

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19	Isolation of Peripheral Blood CD8 T Cells Specific to Porcine Reproductive and Respiratory Syndrome Virus Utilizing Porcine CD137 Activation Marker. <i>Viral Immunology</i> , 2018, 31, 333-337.	1.3	0
20	Broad receptor engagement of an emerging global coronavirus may potentiate its diverse cross-species transmissibility. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E5135-E5143.	7.1	192
21	Infection Dynamics of Hepatitis E Virus in Wild-Type and Immunoglobulin Heavy Chain Knockout J _H ^{â”/â”} Gnotobiotic Piglets. <i>Journal of Virology</i> , 2018, 92, .	3.4	17
22	Pig model mimicking chronic hepatitis E virus infection in immunocompromised patients to assess immune correlates during chronicity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 6914-6923.	7.1	69
23	Efficient priming of CD4â”T cells by Langerin-expressing dendritic cells targeted with porcine epidemic diarrhea virus spike protein domains in pigs. <i>Virus Research</i> , 2017, 227, 212-219.	2.2	22
24	Modulation of Proinflammatory Cytokines in Monocyte-Derived Dendritic Cells by Porcine Reproductive and Respiratory Syndrome Virus Through Interaction with the Porcine Intercellular-Adhesion-Molecule-3-Grabbing Nonintegrin. <i>Viral Immunology</i> , 2016, 29, 546-556.	1.3	27
25	Characterization of Seven Outbreaks of Hemorrhagic Hepatopathy Syndrome in Commercial Pullets Following the Administration of a <i>Salmonella</i> Enteritidis Bacterin in California. <i>Avian Diseases</i> , 2016, 60, 33-42.	1.0	5
26	Evaluation of the use of non-pathogenic porcine circovirus type 1 as a vaccine delivery virus vector to express antigenic epitopes of porcine reproductive and respiratory syndrome virus. <i>Virus Research</i> , 2016, 213, 100-108.	2.2	9
27	RNA transcripts of full-length cDNA clones of rabbit hepatitis E virus are infectious in rabbits. <i>Journal of General Virology</i> , 2015, 96, 1190-1190.	2.9	0
28	The Lysine Residues within the Human Ribosomal Protein S17 Sequence Naturally Inserted into the Viral Nonstructural Protein of a Unique Strain of Hepatitis E Virus Are Important for Enhanced Virus Replication. <i>Journal of Virology</i> , 2015, 89, 3793-3803.	3.4	39
29	Therapeutic targets for the treatment of hepatitis E virus infection. <i>Expert Opinion on Therapeutic Targets</i> , 2015, 19, 1245-1260.	3.4	15
30	Expression of antigenic epitopes of porcine reproductive and respiratory syndrome virus (PRRSV) in a modified live-attenuated porcine circovirus type 2 (PCV2) vaccine virus (PCV1-2a) as a potential bivalent vaccine against both PCV2 and PRRSV. <i>Virus Research</i> , 2015, 210, 154-164.	2.2	14
31	An SH3 binding motif within the nucleocapsid protein of porcine reproductive and respiratory syndrome virus interacts with the host cellular signaling proteins STAM1, TXK, Fyn, Hck, and cortactin. <i>Virus Research</i> , 2015, 204, 31-39.	2.2	11
32	Identification and Fine Mapping of Nuclear and Nucleolar Localization Signals within the Human Ribosomal Protein S17. <i>PLoS ONE</i> , 2015, 10, e0124396.	2.5	16
33	Replacement of the hepatitis E virus ORF3 protein PxxP motif with heterologous late domain motifs affects virus release via interaction with TSG101. <i>Virology</i> , 2015, 486, 198-208.	2.4	32
34	In vivo targeting of porcine reproductive and respiratory syndrome virus antigen through porcine DC-SIGN to dendritic cells elicits antigen-specific CD4T cell immunity in pigs. <i>Vaccine</i> , 2014, 32, 6768-6775.	3.8	21
35	Cross-species infection of pigs with a novel rabbit, but not rat, strain of hepatitis E virus isolated in the United States. <i>Journal of General Virology</i> , 2012, 93, 1687-1695.	2.9	115
36	The PSAP Motif within the ORF3 Protein of an Avian Strain of the Hepatitis E Virus Is Not Critical for Viral Infectivity <i>In Vivo</i> but Plays a Role in Virus Release. <i>Journal of Virology</i> , 2012, 86, 5637-5646.	3.4	44

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37	Assessment of the cross-protective capability of recombinant capsid proteins derived from pig, rat, and avian hepatitis E viruses (HEV) against challenge with a genotype 3 HEV in pigs. <i>Vaccine</i> , 2012, 30, 6249-55.	3.8	16
38	Productive infection of human hepatocellular carcinoma cells by porcine circovirus type 1. <i>Vaccine</i> , 2011, 29, 7303-7306.	3.8	23
39	Expression of the putative ORF1 capsid protein of Torque teno sus virus 2 (TTSuV2) and development of Western blot and ELISA serodiagnostic assays: Correlation between TTSuV2 viral load and IgG antibody level in pigs. <i>Virus Research</i> , 2011, 158, 79-88.	2.2	41
40	Mutational Analysis of the Hypervariable Region of Hepatitis E Virus Reveals Its Involvement in the Efficiency of Viral RNA Replication. <i>Journal of Virology</i> , 2011, 85, 10031-10040.	3.4	66
41	Genetic Evidence for a Connection between Rous Sarcoma Virus Gag Nuclear Trafficking and Genomic RNA Packaging. <i>Journal of Virology</i> , 2009, 83, 6790-6797.	3.4	64
42	Intermolecular Interactions between Retroviral Gag Proteins in the Nucleus. <i>Journal of Virology</i> , 2008, 82, 683-691.	3.4	34
43	Overlapping Roles of the Rous Sarcoma Virus Gag p10 Domain in Nuclear Export and Virion Core Morphology. <i>Journal of Virology</i> , 2007, 81, 10718-10728.	3.4	33