Kristine M Wylie

List of Publications by Year in descending order

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279798 289244 2,319 45 23 40 citations g-index h-index papers 45 45 45 3803 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Fecal microbiome and bile acid metabolome in adult short bowel syndrome. American Journal of Physiology - Renal Physiology, 2022, 322, G154-G168.	3.4	13
2	Longitudinal Dynamics of Skin Bacterial Communities in the Context of Staphylococcus aureus Decolonization. Microbiology Spectrum, 2022, 10, e0267221.	3.0	3
3	Helping others enhances graduate student wellness and mental health. Nature Biotechnology, 2022, 40, 618-619.	17.5	3
4	Association of vaginal bacterial communities and reproductive outcomes with prophylactic antibiotic exposure in a subfertile population undergoing inÂvitro fertilization: a prospective exploratory study. F&S Science, 2021, 2, 71-79.	0.9	3
5	ViroMatch: A Computational Pipeline for the Detection of Viral Sequences from Complex Metagenomic Data. Microbiology Resource Announcements, 2021, 10, .	0.6	11
6	Chronic villitis of unknown etiology: Investigations into viral pathogenesis. Placenta, 2021, 107, 24-30.	1.5	11
7	Characterization of the Eukaryotic Virome of Mice from Different Sources. Microorganisms, 2021, 9, 2064.	3.6	5
8	Association of the Human Semen DNA Virome with Successful In Vitro Fertilization. F&S Science, 2021, 3, 2-9.	0.9	1
9	A virome sequencing approach to feline oral squamous cell carcinoma to evaluate viral causative factors. Veterinary Microbiology, 2020, 240, 108491.	1.9	27
10	The microbiome of the human female reproductive tract. Current Opinion in Physiology, 2020, 13, 87-93.	1.8	9
11	A Protein Epitope Targeted by the Antibody Response to Kawasaki Disease. Journal of Infectious Diseases, 2020, 222, 158-168.	4.0	31
12	Metagenomic analysis of <scp>DNA</scp> viruses from posttransplant lymphoproliferative disorders. Cancer Medicine, 2019, 8, 1013-1023.	2.8	19
13	Impaired Chylomicron Assembly Modifies Hepatic Metabolism Through Bile Acid–Dependent and Transmissible Microbial Adaptations. Hepatology, 2019, 70, 1168-1184.	7.3	12
14	Genome Organization of Canada Goose Coronavirus, A Novel Species Identified in a Mass Die-off of Canada Geese. Scientific Reports, 2019, 9, 5954.	3. 3	36
15	Whole-Genome Sequencing of Klebsiella pneumoniae Isolates to Track Strain Progression in a Single Patient With Recurrent Urinary Tract Infection. Frontiers in Cellular and Infection Microbiology, 2019, 9, 14.	3.9	12
16	The upper-airway microbiota and loss of asthma control among asthmatic children. Nature Communications, 2019, 10, 5714.	12.8	100
17	Differences in the lower airway microbiota of infants with and without cystic fibrosis. Journal of Cystic Fibrosis, 2019, 18, 646-652.	0.7	16
18	Carriage of the Toxic Shock Syndrome Toxin Gene by Contemporary Community-Associated Staphylococcus aureus Isolates. Journal of the Pediatric Infectious Diseases Society, 2019, 8, 470-473.	1.3	4

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19	Detection of Viruses in Clinical Samples by Use of Metagenomic Sequencing and Targeted Sequence Capture. Journal of Clinical Microbiology, 2018, 56, .	3.9	40
20	The vaginal eukaryotic DNA virome and preterm birth. American Journal of Obstetrics and Gynecology, 2018, 219, 189.e1-189.e12.	1.3	90
21	Remnant Small Bowel Length in Pediatric Short Bowel Syndrome and the Correlation with Intestinal Dysbiosis and Linear Growth. Journal of the American College of Surgeons, 2018, 227, 439-449.	0.5	28
22	Early pregnancy vaginal microbiome trends and pretermÂbirth. American Journal of Obstetrics and Gynecology, 2017, 217, 356.e1-356.e18.	1.3	203
23	Genome Sequence of Enterovirus D68 from St. Louis, Missouri, USA, 2016. Genome Announcements, 2017, 5, .	0.8	22
24	The lower airway microbiota in early cystic fibrosis lung disease: a longitudinal analysis. Thorax, 2017, 72, 1104-1112.	5.6	90
25	The Virome of the Human Respiratory Tract. Clinics in Chest Medicine, 2017, 38, 11-19.	2.1	78
26	Environmental Enteric Dysfunction and the Fecal Microbiota in Malawian Children. American Journal of Tropical Medicine and Hygiene, 2017, 96, 473-476.	1.4	41
27	Tissue polymerase chain reaction for the diagnosis of cytomegalovirus disease after allogeneic hematopoietic cell transplantation. American Journal of Hematology, 2017, 92, E19-E20.	4.1	2
28	The Gut Microbiome in Pediatric Allogeneic Hematopoietic Stem Cell Transplant Patients. Open Forum Infectious Diseases, 2017, 4, S231-S231.	0.9	0
29	Metagenomic Approach for Identification of the Pathogens Associated with Diarrhea in Stool Specimens. Journal of Clinical Microbiology, 2016, 54, 368-375.	3.9	98
30	The Value of Tissue Polymerase Chain Reaction Testing for the Diagnosis of Tissue-Invasive Cytomegalovirus Disease Following Allogeneic Hematopoietic Cell Transplantation. Blood, 2016, 128, 3420-3420.	1.4	0
31	The transcriptional profile of coronary arteritis in Kawasaki disease. BMC Genomics, 2015, 16, 1076.	2.8	63
32	Enhanced virome sequencing using targeted sequence capture. Genome Research, 2015, 25, 1910-1920.	5.5	209
33	Emergence of Rotavirus G12P[8] in St. Louis During the 2012â€"2013 Rotavirus Season. Journal of the Pediatric Infectious Diseases Society, 2015, 4, e84-e89.	1.3	6
34	Genome Sequence of Enterovirus D68 from St. Louis, Missouri, USA. Emerging Infectious Diseases, 2015, 21, 184-186.	4.3	35
35	Development and Evaluation of an Enterovirus D68 Real-Time Reverse Transcriptase PCR Assay. Journal of Clinical Microbiology, 2015, 53, 2641-2647.	3.9	55
36	Abstract O.20: Deep RNA Sequencing Reveals a Transcriptional Profile of Cytotoxic T Lymphocyte Activation, Antigen Presentation, Immunoglobulin Production, and Type I Interferon Response in Kawasaki Disease Arteritis. Circulation, 2015, 131, .	1.6	0

#	Article	IF	CITATIONS
37	Metagenomic analysis of double-stranded DNA viruses in healthy adults. BMC Biology, 2014, 12, 71.	3.8	181
38	Virome genomics: a tool for defining the human virome. Current Opinion in Microbiology, 2013, 16, 479-484.	5.1	38
39	Emerging view of the human virome. Translational Research, 2012, 160, 283-290.	5.0	122
40	Sequence Analysis of the Human Virome in Febrile and Afebrile Children. PLoS ONE, 2012, 7, e27735.	2.5	159
41	Novel Bacterial Taxa in the Human Microbiome. PLoS ONE, 2012, 7, e35294.	2.5	86
42	Increased Prevalence of Anellovirus in Pediatric Patients with Fever. PLoS ONE, 2012, 7, e50937.	2.5	61
43	Astrovirus MLB2 Viremia in Febrile Child. Emerging Infectious Diseases, 2011, 17, 2050-2.	4.3	57
44	The Human Virome in Children and its Relationship to Febrile Illness. Nature Precedings, 2010, , .	0.1	0
45	A physical map of the human Y chromosome. Nature, 2001, 409, 943-945.	27.8	239