

Patrick R Shea

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/863318/publications.pdf>

Version: 2024-02-01

26
papers

1,277
citations

471509

17
h-index

677142

22
g-index

26
all docs

26
docs citations

26
times ranked

2486
citing authors

#	ARTICLE	IF	CITATIONS
1	A genome-wide screen for variants influencing certolizumab pegol response in a moderate to severe rheumatoid arthritis population. <i>PLoS ONE</i> , 2022, 17, e0261165.	2.5	2
2	Genome-wide association study (GWAS) of human host factors influencing viral severity of herpes simplex virus type 2 (HSV-2). <i>Genes and Immunity</i> , 2019, 20, 112-120.	4.1	17
3	Exome sequencing of an adolescent with nonalcoholic fatty liver disease identifies a clinically actionable case of Wilson disease. <i>Journal of Physical Education and Sports Management</i> , 2018, 4, a003087.	1.2	3
4	Association of CYP2B6 Single-Nucleotide Polymorphisms Altering Efavirenz Metabolism With Hepatitis C Virus (HCV) Treatment Relapse Among Human Immunodeficiency Virus/HCVâ€“Coinfected African Americans Receiving Ledipasvir/Sofosbuvir in the ION-4 Trial. <i>Clinical Infectious Diseases</i> , 2018, 66, 1953-1956.	5.8	0
5	Killer cell immunoglobulinâ€“like receptor 3DL1 variation modifies HLA-B*57 protection against HIV-1. <i>Journal of Clinical Investigation</i> , 2018, 128, 1903-1912.	8.2	52
6	No Evidence for Association of Î²â€“Defensin Genomic Copy Number with HIV Susceptibility, HIV Load during Clinical Latency, or Progression to AIDS. <i>Annals of Human Genetics</i> , 2017, 81, 27-34.	0.8	4
7	Evaluating the Impact of Functional Genetic Variation on HIV-1 Control. <i>Journal of Infectious Diseases</i> , 2017, 216, 1063-1069.	4.0	20
8	Immune perturbations in HIV-1â€“infected individuals who make broadly neutralizing antibodies. <i>Science Immunology</i> , 2016, 1, aag0851.	11.9	120
9	Genome-Wide Association Study of the Human Genetic Factors Influencing the Risk of Adverse Events during Idelalisib Therapy in Patients with Relapsed Indolent Lymphoma. <i>Blood</i> , 2016, 128, 5284-5284.	1.4	0
10	Polymorphisms of large effect explain the majority of the host genetic contribution to variation of HIV-1 virus load. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 14658-14663.	7.1	154
11	Testing for risk and protective trends in genetic analyses of HIV acquisition. <i>Biostatistics</i> , 2015, 16, 268-280.	1.5	0
12	Host Genetics of HIV Acquisition and Viral Control. <i>Annual Review of Medicine</i> , 2013, 64, 203-217.	12.2	29
13	Association Study of Common Genetic Variants and HIV-1 Acquisition in 6,300 Infected Cases and 7,200 Controls. <i>PLoS Pathogens</i> , 2013, 9, e1003515.	4.7	109
14	Polymorphisms in Regulator of Protease B (RopB) Alter Disease Phenotype and Strain Virulence of Serotype M3 Group A Streptococcus. <i>Journal of Infectious Diseases</i> , 2012, 205, 1719-1729.	4.0	25
15	Full-Genome Dissection of an Epidemic of Severe Invasive Disease Caused by a Hypervirulent, Recently Emerged Clone of Group A Streptococcus. <i>American Journal of Pathology</i> , 2012, 180, 1522-1534.	3.8	70
16	Polymorphisms in regulator of protease B (RopB) alter disease phenotype and strain virulence of serotype M3 Group A Streptococcus. <i>FASEB Journal</i> , 2012, 26, 1034.3.	0.5	0
17	Group A Streptococcus emm Gene Types in Pharyngeal Isolates, Ontario, Canada, 2002â€“2010. <i>Emerging Infectious Diseases</i> , 2011, 17, 2010-7.	4.3	65
18	Distinct signatures of diversifying selection revealed by genome analysis of respiratory tract and invasive bacterial populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 5039-5044.	7.1	90

#	ARTICLE	IF	CITATIONS
19	Naturally occurring single amino acid replacements in a regulatory protein alter streptococcal gene expression and virulence in mice. <i>Journal of Clinical Investigation</i> , 2011, 121, 1956-1968.	8.2	63
20	Molecular complexity of successive bacterial epidemics deconvoluted by comparative pathogenomics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 4371-4376.	7.1	153
21	Interactome analysis of longitudinal pharyngeal infection of cynomolgus macaques by group A <i>Streptococcus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 4693-4698.	7.1	30
22	Lower respiratory tract infection in cynomolgus macaques (<i>Macaca fascicularis</i>) infected with group A <i>Streptococcus</i> . <i>Microbial Pathogenesis</i> , 2010, 49, 336-347.	2.9	13
23	RNASEL and RNASEL inhibitor variation and prostate cancer risk in Afro-Caribbeans. <i>Prostate</i> , 2008, 68, 354-359.	2.3	23
24	Androgen receptor CAG repeat polymorphism is associated with fat-free mass in men. <i>Journal of Applied Physiology</i> , 2005, 98, 132-137.	2.5	92
25	Vitamin D Receptor Genotype Is Associated With Fat-Free Mass and Sarcopenia in Elderly Men. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2004, 59, B10-B15.	3.6	126
26	ELAC2 and prostate cancer risk in Afro-Caribbeans of Tobago. <i>Human Genetics</i> , 2002, 111, 398-400.	3.8	17