Victoria J Wright

List of Publications by Year in Descending Order

33 papers	2,002 citations	19 h-index	34 g-index
34 ext. papers	2,557 ext. citations	12.4 avg, IF	3.59 L-index

#	Paper	IF	Citations
33	Identification of novel locus associated with coronary artery aneurysms and validation of loci for susceptibility to Kawasaki disease. *European Journal of Human Genetics*, **2021**, 29, 1734-1744	5.3	2
32	A Novel Framework for Phenotyping Children With Suspected or Confirmed Infection for Future Biomarker Studies. *Frontiers in Pediatrics*, **2021**, 9, 688272	3.4	2
31	Identification of Reduced Host Transcriptomic Signatures for Tuberculosis Disease and Digital PCR-Based Validation and Quantification. *Frontiers in Immunology*, **2021**, 12, 637164	8.4	4
30	Discovery and validation of a three-gene signature to distinguish COVID-19 and other viral infections in emergency infectious disease presentations: a case-control and observational cohort study. *Lancet Microbe, The*, **2021**, 2, e594-e603	22.2	5
29	Biomarkers for the Discrimination of Acute Kawasaki Disease From Infections in Childhood. *Frontiers in Pediatrics*, **2020**, 8, 355	3.4	7
28	A Rare Mutation in SPLUNC1 Affects Bacterial Adherence and Invasion in Meningococcal Disease. *Clinical Infectious Diseases*, **2020**, 70, 2045-2053	11.6	4
27	HLA-C variants associated with amino acid substitutions in the peptide binding groove influence susceptibility to Kawasaki disease. *Human Immunology*, **2019**, 80, 731-738	2.3	4
26	Identification of regulatory variants associated with genetic susceptibility to meningococcal disease. *Scientific Reports*, **2019**, 9, 6966	4.9	3
25	Extensive Ethnic Variation and Linkage Disequilibrium at the Locus: Different Genetic Associations Revealed in Kawasaki Disease. *Frontiers in Immunology*, **2019**, 10, 185	8.4	27
24	Biosynthetic homeostasis and resilience of the complement system in health and infectious disease. *EBioMedicine*, **2019**, 45, 303-313	8.8	7
23	Mycobacterium tuberculosis Exploits a Molecular Off Switch of the Immune System for Intracellular Survival. *Scientific Reports*, **2018**, 8, 661	4.9	25
22	Diagnosis of Kawasaki Disease Using a Minimal Whole-Blood Gene Expression Signature. *JAMA Pediatrics*, **2018**, 172, e182293	8.3	52
21	Cohort profile of the Biomarkers of Acute Serious Illness in Children (BASIC) study: a prospective multicentre cohort study in critically ill children. *BMJ Open*, **2018**, 8, e024729	3	1
20	Life-threatening infections in children in Europe (the EUCLIDS Project): a prospective cohort study. *The Lancet Child and Adolescent Health*, **2018**, 2, 404-414	14.5	40
19	PRINCESS: Privacy-protecting Rare disease International Network Collaboration via Encryption through Software guard extensionS. *Bioinformatics*, **2017**, 33, 871-878	7.2	36
18	Diagnosis of Bacterial Infection Using a 2-Transcript Host RNA Signature in Febrile Infants 60 Days or Younger. *JAMA - Journal of the American Medical Association*, **2017**, 317, 1577-1578	27.4	25
17	A genome-wide association analysis identifies NMNAT2 and HCP5 as susceptibility loci for Kawasaki disease. *Journal of Human Genetics*, **2017**, 62, 1023-1029	4.3	29

List of Publications

16	Childhood tuberculosis is associated with decreased abundance of T cell gene transcripts and impaired T cell function. *PLoS ONE*, **2017**, 12, e0185973	3.7	9
15	Diagnostic Test Accuracy of a 2-Transcript Host RNA Signature for Discriminating Bacterial vs Viral Infection in Febrile Children. *JAMA - Journal of the American Medical Association*, **2016**, 316, 835-45	27.4	166
14	Genetic Variation in the SLC8A1 Calcium Signaling Pathway Is Associated With Susceptibility to Kawasaki Disease and Coronary Artery Abnormalities. *Circulation: Cardiovascular Genetics*, **2016**, 9, 559-568		33
13	Natural resistance to Meningococcal Disease related to CFH loci: Meta-analysis of genome-wide association studies. *Scientific Reports*, **2016**, 6, 35842	4.9	26
12	Host RNA signatures for diagnostics: an example from paediatric tuberculosis in Africa. *Journal of Infection*, **2014**, 69 Suppl 1, S28-31	18.9	15
11	Diagnosis of childhood tuberculosis and host RNA expression in Africa. *New England Journal of Medicine*, **2014**, 370, 1712-1723	59.2	229
10	Global gene expression profiling identifies new therapeutic targets in acute Kawasaki disease. *Genome Medicine*, **2014**, 6, 541	14.4	83
9	Transcriptomic profiling in childhood H1N1/09 influenza reveals reduced expression of protein synthesis genes. *Journal of Infectious Diseases*, **2013**, 208, 1664-8	7	51
8	Detection of tuberculosis in HIV-infected and -uninfected African adults using whole blood RNA expression signatures: a case-control study. *PLoS Medicine*, **2013**, 10, e1001538	11.6	224
7	A genome-wide association study identifies three new risk loci for Kawasaki disease. *Nature Genetics*, **2012**, 44, 517-21	36.3	217
6	Genome-wide association study identifies FCGR2A as a susceptibility locus for Kawasaki disease. *Nature Genetics*, **2011**, 43, 1241-6	36.3	236
5	Genome-wide association study identifies variants in the CFH region associated with host susceptibility to meningococcal disease. *Nature Genetics*, **2010**, 42, 772-6	36.3	221
4	A genome-wide association study identifies novel and functionally related susceptibility Loci for Kawasaki disease. *PLoS Genetics*, **2009**, 5, e1000319	6	188
3	Genetic polymorphisms in host response to meningococcal infection: the role of susceptibility and severity genes. *Vaccine*, **2009**, 27 Suppl 2, B90-102	4.1	27
2	Angiotensin I converting enzyme inhibitor and worsening of anemia in hemodialysis patients: Prevention with rHuEPO.. *Nihon Toseki Igakkai Zasshi*, **1997**, 30, 315-320	0.3	
1	Chronic portal-systemic shunt encephalopathy (CPSE) in a hemodialysis patient: A case report.. *Nihon Toseki Igakkai Zasshi*, **1997**, 30, 999-1005	0.3	4