

# Victoria J Wright

## List of Publications by Year in descending order

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Version: 2024-02-01

34  
papers

2,894  
citations

361296  
20  
h-index

395590  
33  
g-index

34  
all docs

34  
docs citations

34  
times ranked

4351  
citing authors

#	ARTICLE	IF	CITATIONS
1	Diagnosis of Childhood Tuberculosis and Host RNA Expression in Africa. <i>New England Journal of Medicine</i> , 2014, 370, 1712-1723.	13.9	324
2	Detection of Tuberculosis in HIV-Infected and -Uninfected African Adults Using Whole Blood RNA Expression Signatures: A Case-Control Study. <i>PLoS Medicine</i> , 2013, 10, e1001538.	3.9	314
3	Genome-wide association study identifies FCGR2A as a susceptibility locus for Kawasaki disease. <i>Nature Genetics</i> , 2011, 43, 1241-1246.	9.4	297
4	A genome-wide association study identifies three new risk loci for Kawasaki disease. <i>Nature Genetics</i> , 2012, 44, 517-521.	9.4	284
5	Genome-wide association study identifies variants in the CFH region associated with host susceptibility to meningococcal disease. <i>Nature Genetics</i> , 2010, 42, 772-776.	9.4	275
6	Diagnostic Test Accuracy of a 2-Transcript Host RNA Signature for Discriminating Bacterial vs Viral Infection in Febrile Children. <i>JAMA - Journal of the American Medical Association</i> , 2016, 316, 835.	3.8	263
7	A Genome-Wide Association Study Identifies Novel and Functionally Related Susceptibility Loci for Kawasaki Disease. <i>PLoS Genetics</i> , 2009, 5, e1000319.	1.5	234
8	Global gene expression profiling identifies new therapeutic targets in acute Kawasaki disease. <i>Genome Medicine</i> , 2014, 6, 541.	3.6	126
9	Diagnosis of Kawasaki Disease Using a Minimal Whole-Blood Gene Expression Signature. <i>JAMA Pediatrics</i> , 2018, 172, e182293.	3.3	92
10	Transcriptomic Profiling in Childhood H1N1/09 Influenza Reveals Reduced Expression of Protein Synthesis Genes. <i>Journal of Infectious Diseases</i> , 2013, 208, 1664-1668.	1.9	84
11	PRINCESS: Privacy-protecting Rare disease International Network Collaboration via Encryption through Software guard extensionS. <i>Bioinformatics</i> , 2017, 33, 871-878.	1.8	75
12	Life-threatening infections in children in Europe (the EUCLIDS Project): a prospective cohort study. <i>The Lancet Child and Adolescent Health</i> , 2018, 2, 404-414.	2.7	69
13	Diagnosis of Bacterial Infection Using a 2-Transcript Host RNA Signature in Febrile Infants 60 Days or Younger. <i>JAMA - Journal of the American Medical Association</i> , 2017, 317, 1577.	3.8	46
14	Genetic Variation in the SLC8A1 Calcium Signaling Pathway Is Associated With Susceptibility to Kawasaki Disease and Coronary Artery Abnormalities. <i>Circulation: Cardiovascular Genetics</i> , 2016, 9, 559-568.	5.1	45
15	Extensive Ethnic Variation and Linkage Disequilibrium at the FCGR2/3 Locus: Different Genetic Associations Revealed in Kawasaki Disease. <i>Frontiers in Immunology</i> , 2019, 10, 185.	2.2	43
16	A genome-wide association analysis identifies NMNAT2 and HCP5 as susceptibility loci for Kawasaki disease. <i>Journal of Human Genetics</i> , 2017, 62, 1023-1029.	1.1	40
17	Genetic polymorphisms in host response to meningococcal infection: The role of susceptibility and severity genes. <i>Vaccine</i> , 2009, 27, B90-B102.	1.7	35
18	A Novel Framework for Phenotyping Children With Suspected or Confirmed Infection for Future Biomarker Studies. <i>Frontiers in Pediatrics</i> , 2021, 9, 688272.	0.9	34

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19	Natural resistance to Meningococcal Disease related to CFH loci: Meta-analysis of genome-wide association studies. <i>Scientific Reports</i> , 2016, 6, 35842.	1.6	33
20	<i>Mycobacterium tuberculosis</i> Exploits a Molecular Off Switch of the Immune System for Intracellular Survival. <i>Scientific Reports</i> , 2018, 8, 661.	1.6	33
21	Identification of Reduced Host Transcriptomic Signatures for Tuberculosis Disease and Digital PCR-Based Validation and Quantification. <i>Frontiers in Immunology</i> , 2021, 12, 637164.	2.2	25
22	Biosynthetic homeostasis and resilience of the complement system in health and infectious disease. <i>EBioMedicine</i> , 2019, 45, 303-313.	2.7	20
23	Biomarkers for the Discrimination of Acute Kawasaki Disease From Infections in Childhood. <i>Frontiers in Pediatrics</i> , 2020, 8, 355.	0.9	17
24	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. <i>Lancet Microbe</i> , The, 2021, 2, e594-e603.	3.4	17
25	Host RNA signatures for diagnostics: An example from paediatric tuberculosis in Africa. <i>Journal of Infection</i> , 2014, 69, S28-S31.	1.7	16
26	Childhood tuberculosis is associated with decreased abundance of T cell gene transcripts and impaired T cell function. <i>PLoS ONE</i> , 2017, 12, e0185973.	1.1	15
27	Identification of novel locus associated with coronary artery aneurysms and validation of loci for susceptibility to Kawasaki disease. <i>European Journal of Human Genetics</i> , 2021, 29, 1734-1744.	1.4	10
28	A Rare Mutation in <i>SPLUNC1</i> Affects Bacterial Adherence and Invasion in Meningococcal Disease. <i>Clinical Infectious Diseases</i> , 2020, 70, 2045-2053.	2.9	6
29	Kawasaki Disease Patient Stratification and Pathway Analysis Based on Host Transcriptomic and Proteomic Profiles. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5655.	1.8	6
30	HLA-C variants associated with amino acid substitutions in the peptide binding groove influence susceptibility to Kawasaki disease. <i>Human Immunology</i> , 2019, 80, 731-738.	1.2	5
31	Cohort profile of the Biomarkers of Acute Serious Illness in Children (BASIC) study: a prospective multicentre cohort study in critically ill children. <i>BMJ Open</i> , 2018, 8, e024729.	0.8	4
32	Chronic portal-systemic shunt encephalopathy (CPSE) in a hemodialysis patient: A case report.. <i>Nihon Toseki Igakkai Zasshi</i> , 1997, 30, 999-1005.	0.2	4
33	Identification of regulatory variants associated with genetic susceptibility to meningococcal disease. <i>Scientific Reports</i> , 2019, 9, 6966.	1.6	3
34	Angiotensin I converting enzyme inhibitor and worsening of anemia in hemodialysis patients: Prevention with rHuEPO.. <i>Nihon Toseki Igakkai Zasshi</i> , 1997, 30, 315-320.	0.2	0