

Victoria J Wright

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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 | Genome-wide association study identifies FCGR2A as a susceptibility locus for Kawasaki disease. <i>Nature Genetics</i> , 2011 , 43, 1241-6 | 36.3 | 236 |
| 32 | Diagnosis of childhood tuberculosis and host RNA expression in Africa. <i>New England Journal of Medicine</i> , 2014 , 370, 1712-1723 | 59.2 | 229 |
| 31 | 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 | 11.6 | 224 |
| 30 | Genome-wide association study identifies variants in the CFH region associated with host susceptibility to meningococcal disease. <i>Nature Genetics</i> , 2010 , 42, 772-6 | 36.3 | 221 |
| 29 | A genome-wide association study identifies three new risk loci for Kawasaki disease. <i>Nature Genetics</i> , 2012 , 44, 517-21 | 36.3 | 217 |
| 28 | A genome-wide association study identifies novel and functionally related susceptibility Loci for Kawasaki disease. <i>PLoS Genetics</i> , 2009 , 5, e1000319 | 6 | 188 |
| 27 | 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, 835-45 | 27.4 | 166 |
| 26 | Global gene expression profiling identifies new therapeutic targets in acute Kawasaki disease. <i>Genome Medicine</i> , 2014 , 6, 541 | 14.4 | 83 |
| 25 | Diagnosis of Kawasaki Disease Using a Minimal Whole-Blood Gene Expression Signature. <i>JAMA Pediatrics</i> , 2018 , 172, e182293 | 8.3 | 52 |
| 24 | Transcriptomic profiling in childhood H1N1/09 influenza reveals reduced expression of protein synthesis genes. <i>Journal of Infectious Diseases</i> , 2013 , 208, 1664-8 | 7 | 51 |
| 23 | 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 | 14.5 | 40 |
| 22 | PRINCESS: Privacy-protecting Rare disease International Network Collaboration via Encryption through Software guard extensionS. <i>Bioinformatics</i> , 2017 , 33, 871-878 | 7.2 | 36 |
| 21 | 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 | | 33 |
| 20 | 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 | 4.3 | 29 |
| 19 | Extensive Ethnic Variation and Linkage Disequilibrium at the Locus: Different Genetic Associations Revealed in Kawasaki Disease. <i>Frontiers in Immunology</i> , 2019 , 10, 185 | 8.4 | 27 |
| 18 | Genetic polymorphisms in host response to meningococcal infection: the role of susceptibility and severity genes. <i>Vaccine</i> , 2009 , 27 Suppl 2, B90-102 | 4.1 | 27 |
| 17 | Natural resistance to Meningococcal Disease related to CFH loci: Meta-analysis of genome-wide association studies. <i>Scientific Reports</i> , 2016 , 6, 35842 | 4.9 | 26 |

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| 16 | 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, 1577-1578 | 27.4 | 25 |
| 15 | Mycobacterium tuberculosis Exploits a Molecular Off Switch of the Immune System for Intracellular Survival. <i>Scientific Reports</i> , 2018 , 8, 661 | 4.9 | 25 |
| 14 | Host RNA signatures for diagnostics: an example from paediatric tuberculosis in Africa. <i>Journal of Infection</i> , 2014 , 69 Suppl 1, S28-31 | 18.9 | 15 |
| 13 | Childhood tuberculosis is associated with decreased abundance of T cell gene transcripts and impaired T cell function. <i>PLoS ONE</i> , 2017 , 12, e0185973 | 3.7 | 9 |
| 12 | Biosynthetic homeostasis and resilience of the complement system in health and infectious disease. <i>EBioMedicine</i> , 2019 , 45, 303-313 | 8.8 | 7 |
| 11 | Biomarkers for the Discrimination of Acute Kawasaki Disease From Infections in Childhood. <i>Frontiers in Pediatrics</i> , 2020 , 8, 355 | 3.4 | 7 |
| 10 | 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> , 2021 , 2, e594-e603 | 22.2 | 5 |
| 9 | 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 | 2.3 | 4 |
| 8 | Chronic portal-systemic shunt encephalopathy (CPSE) in a hemodialysis patient: A case report.. <i>Nihon Toseki Igakkai Zasshi</i> , 1997 , 30, 999-1005 | 0.3 | 4 |
| 7 | A Rare Mutation in SPLUNC1 Affects Bacterial Adherence and Invasion in Meningococcal Disease. <i>Clinical Infectious Diseases</i> , 2020 , 70, 2045-2053 | 11.6 | 4 |
| 6 | Identification of Reduced Host Transcriptomic Signatures for Tuberculosis Disease and Digital PCR-Based Validation and Quantification. <i>Frontiers in Immunology</i> , 2021 , 12, 637164 | 8.4 | 4 |
| 5 | Identification of regulatory variants associated with genetic susceptibility to meningococcal disease. <i>Scientific Reports</i> , 2019 , 9, 6966 | 4.9 | 3 |
| 4 | 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 | 5.3 | 2 |
| 3 | A Novel Framework for Phenotyping Children With Suspected or Confirmed Infection for Future Biomarker Studies. <i>Frontiers in Pediatrics</i> , 2021 , 9, 688272 | 3.4 | 2 |
| 2 | 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 | 3 | 1 |
| 1 | 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 | 0.3 | |