

# Veron Ramsuran

## List of Publications by Year in descending order

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Version: 2024-02-01

42  
papers

1,073  
citations

516215

16  
h-index

454577

30  
g-index

43  
all docs

43  
docs citations

43  
times ranked

2018  
citing authors

#	ARTICLE	IF	CITATIONS
1	Hypermethylation at the <i>CXCR5</i> gene locus limits trafficking potential of CD8+ T cells into B-cell follicles during HIV-1 infection. <i>Blood Advances</i> , 2022, 6, 1904-1916.	2.5	6
2	High-Resolution Melting Analysis to Detect Antimicrobial Resistance Determinants in South African <i>Neisseria gonorrhoeae</i> Clinical Isolates and Specimens. <i>International Journal of Microbiology</i> , 2022, 1-9.	0.9	1
3	The challenges of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) testing in low-middle income countries and possible cost-effective measures in resource-limited settings. <i>Globalization and Health</i> , 2022, 18, 5.	2.4	16
4	The Effect of miRNA Gene Regulation on HIV Disease. <i>Frontiers in Genetics</i> , 2022, 13, .	1.1	7
5	Genetic variation that determines <i>TAPBP</i> expression levels associates with the course of malaria in an HLA allotype-dependent manner. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	3
6	Comparison of methods for the detection of <i>Neisseria gonorrhoeae</i> from South African women attending antenatal care. <i>International Journal of STD and AIDS</i> , 2021, 32, 396-402.	0.5	0
7	Genome Sequences of Five Novel <i>Neisseria gonorrhoeae</i> Sequence Types Isolated in KwaZulu-Natal, South Africa. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.3	3
8	Performance of TaqMan probes for the detection of sexually transmitted infections in South African women. <i>African Journal of Laboratory Medicine</i> , 2021, 10, 1124.	0.2	0
9	Epigenetic Regulation of BST-2 Expression Levels and the Effect on HIV-1 Pathogenesis. <i>Frontiers in Immunology</i> , 2021, 12, 669241.	2.2	4
10	Strong correlation between urine and vaginal swab samples for bacterial vaginosis. <i>Southern African Journal of Infectious Diseases</i> , 2021, 36, 199.	0.3	2
11	Transient association between semen exposure and biomarkers of genital inflammation in South African women at risk of HIV infection. <i>Journal of the International AIDS Society</i> , 2021, 24, e25766.	1.2	5
12	Antimicrobial Resistance Mechanisms, Multilocus Sequence Typing, and NG-STAR Sequence Types of Diverse <i>Neisseria gonorrhoeae</i> Isolates in KwaZulu-Natal, South Africa. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0075921.	1.4	10
13	SARS-CoV-2 and helminth co-infections, and environmental pollution exposure: An epidemiological and immunological perspective. <i>Environment International</i> , 2021, 156, 106695.	4.8	17
14	16S rRNA-based metagenomic profiling of microbes on contact surfaces within shared sanitation facilities. <i>Ecological Genetics and Genomics</i> , 2021, 21, 100095.	0.3	1
15	An HLA-I signature favouring KIR-educated Natural Killer cells mediates immune control of HIV in children and contrasts with the HLA-B-restricted CD8+ T-cell-mediated immune control in adults. <i>PLoS Pathogens</i> , 2021, 17, e1010090.	2.1	12
16	Deciphering DNA Methylation in HIV Infection. <i>Frontiers in Immunology</i> , 2021, 12, 795121.	2.2	5
17	The Role of Highly Active Antiretroviral Therapy (HAART) on Interleukin 17A (IL-17A) in Normotensive and Preeclamptic Black South African Women. <i>Infectious Diseases in Obstetrics and Gynecology</i> , 2020, 1-11.	0.4	4
18	Next Generation Sequencing and Bioinformatics Analysis of Family Genetic Inheritance. <i>Frontiers in Genetics</i> , 2020, 11, 544162.	1.1	41

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19	Prevalence of Genotypes and Subtypes of <i>Gardnerella vaginalis</i> in South African Pregnant Women. <i>Infectious Diseases in Obstetrics and Gynecology</i> , 2020, 2020, 1-12.	0.4	1
20	HLA-G Polymorphisms Associated with HIV Infection and Preeclampsia in South Africans of African Ancestry. <i>BioMed Research International</i> , 2020, 2020, 1-14.	0.9	5
21	Assessing a diagnosis tool for bacterial vaginosis. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1481-1485.	1.3	10
22	Maternal variants within the apolipoprotein L1 gene are associated with preeclampsia in a South African cohort of African ancestry. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2020, 246, 129-133.	0.5	21
23	Genome-wide Association Study Identifies HLA-DPB1 as a Significant Risk Factor for Severe Aplastic Anemia. <i>American Journal of Human Genetics</i> , 2020, 106, 264-271.	2.6	25
24	Gene polymorphisms of uric acid are associated with pre-eclampsia in South Africans of African ancestry. <i>Hypertension in Pregnancy</i> , 2020, 39, 103-116.	0.5	2
25	Polymorphisms within vitamin D binding protein gene within a Preeclamptic South African population. <i>Hypertension in Pregnancy</i> , 2019, 38, 260-267.	0.5	5
26	Effects of genetic variability on rifampicin and isoniazid pharmacokinetics in South African patients with recurrent tuberculosis. <i>Pharmacogenomics</i> , 2019, 20, 225-240.	0.6	32
27	Pre-eclampsia: the role of highly active antiretroviral therapy and immune markers. <i>Inflammation Research</i> , 2019, 68, 47-57.	1.6	18
28	Effect of genetic variation in <i>UGT1A</i> and <i>ABCB1</i> on moxifloxacin pharmacokinetics in South African patients with tuberculosis. <i>Pharmacogenomics</i> , 2018, 19, 17-29.	0.6	16
29	Elevated <i>HLA-A</i> expression impairs HIV control through inhibition of NKG2A-expressing cells. <i>Science</i> , 2018, 359, 86-90.	6.0	135
30	Variation in the Untranslated Genome and Susceptibility to Infections. <i>Frontiers in Immunology</i> , 2018, 9, 2046.	2.2	17
31	Case report: mechanisms of HIV elite control in two African women. <i>BMC Infectious Diseases</i> , 2018, 18, 54.	1.3	82
32	Killer cell immunoglobulin-like receptor 3DL1 variation modifies HLA-B*57 protection against HIV-1. <i>Journal of Clinical Investigation</i> , 2018, 128, 1903-1912.	3.9	52
33	Sequence and Phylogenetic Analysis of the Untranslated Promoter Regions for <i>HLA</i> Class I Genes. <i>Journal of Immunology</i> , 2017, 198, 2320-2329.	0.4	42
34	Posttranscriptional Regulation of HLA-A Protein Expression by Alternative Polyadenylation Signals Involving the RNA-Binding Protein Syncrin. <i>Journal of Immunology</i> , 2017, 199, 3892-3899.	0.4	31
35	Hyperbilirubinemia in atazanavir-treated human immunodeficiency virus-infected patients: the impact of the <i>UGT1A1</i> *28 allele. <i>Pharmacogenomics and Personalized Medicine</i> , 2017, Volume 10, 233-234.	0.4	0
36	HLA-C Level Is Regulated by a Polymorphic Oct1 Binding Site in the HLA-C Promoter Region. <i>American Journal of Human Genetics</i> , 2016, 99, 1353-1358.	2.6	49

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37	Epigenetic regulation of differential HLA-A allelic expression levels. <i>Human Molecular Genetics</i> , 2015, 24, 4268-4275.	1.4	94
38	Epigenetic mechanisms, T-cell activation, and CCR5 genetics interact to regulate T-cell expression of CCR5, the major HIV-1 coreceptor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E4762-71.	3.3	48
39	Effect of Female Genital Schistosomiasis and Anti-Schistosomal Treatment on Monocytes, CD4+ T-Cells and CCR5 Expression in the Female Genital Tract. <i>PLoS ONE</i> , 2014, 9, e98593.	1.1	47
40	Genetic interplay between HLA-C and MIR148A in HIV control and Crohn disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 20705-20710.	3.3	109
41	Duffy-Null Associated Low Neutrophil Counts Influence HIV-1 Susceptibility in High-Risk South African Black Women. <i>Clinical Infectious Diseases</i> , 2011, 52, 1248-1256.	2.9	69
42	Influence of Variations in CCL3L1 and CCR5 on Tuberculosis in a Northwestern Colombian Population. <i>Journal of Infectious Diseases</i> , 2011, 203, 1590-1594.	1.9	26