

Thirumalaisamy P Velavan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7089932/publications.pdf>

Version: 2024-02-01

177
papers

7,103
citations

109321

35
h-index

79698

73
g-index

181
all docs

181
docs citations

181
times ranked

12104
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of PD-L1 gene polymorphisms and circulating sPD-L1 levels with HBV infection susceptibility and related liver disease progression. <i>Gene</i> , 2022, 806, 145935.	2.2	2
2	Emergence of B.1.1.318 SARS-CoV-2 viral lineage and high incidence of alpha B.1.1.7 variant of concern in the Republic of Gabon. <i>International Journal of Infectious Diseases</i> , 2022, 114, 151-154.	3.3	10
3	Emergence of new SARS-CoV-2 Variant of Concern Omicron (B.1.1.529) - highlights Africa's research capabilities, but exposes major knowledge gaps, inequities of vaccine distribution, inadequacies in global COVID-19 response and control efforts. <i>International Journal of Infectious Diseases</i> , 2022, 114, 268-272.	3.3	136
4	Circulating level of sPD-1 and PD-1 genetic variants are associated with hepatitis B infection and related liver disease progression. <i>International Journal of Infectious Diseases</i> , 2022, 115, 229-236.	3.3	5
5	SARS-CoV-2 viral dynamics of the first 1000 sequences from Vietnam and neighbouring ASEAN countries. <i>IJID Regions</i> , 2022, 2, 175-179.	1.3	3
6	SARS-CoV-2 B.1.214.1, B.1.214.2 and B.1.620 are predominant lineages between December 2020 and July 2021 in the Republic of Congo. <i>IJID Regions</i> , 2022, 3, 106-113.	1.3	1
7	Evaluation of SARS-CoV-2 diagnostics and risk factors associated with SARS-CoV-2 infection in Zambia. <i>International Journal of Infectious Diseases</i> , 2022, 120, 150-157.	3.3	4
8	Monkeypox 2022 outbreak: An update. <i>Tropical Medicine and International Health</i> , 2022, 27, 604-605.	2.3	114
9	Host genetic loci LZTFL1 and CCL2 associated with SARS-CoV-2 infection and severity of COVID-19. <i>International Journal of Infectious Diseases</i> , 2022, 122, 427-436.	3.3	11
10	High prevalence of antibiotic-resistant <i>Escherichia coli</i> in Congolese students. <i>International Journal of Infectious Diseases</i> , 2021, 103, 119-123.	3.3	11
11	Reply to: Asymptomatic infection by SARS 2 coronavirus: invisible but invincible. <i>International Journal of Infectious Diseases</i> , 2021, 102, 85-86.	3.3	0
12	Targeting interleukin 6 signaling by monoclonal antibody siltuximab on cholangiocarcinoma. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2021, 36, 1334-1345.	2.8	18
13	Hepatitis E virus genome detection in commercial pork livers and pork meat products in Germany. <i>Journal of Viral Hepatitis</i> , 2021, 28, 196-204.	2.0	37
14	COVID-19 in Africa: between hope and reality. <i>Lancet Infectious Diseases</i> , The, 2021, 21, 315.	9.1	24
15	Longitudinal Monitoring of Lactate in Hospitalized and Ambulatory COVID-19 Patients. <i>American Journal of Tropical Medicine and Hygiene</i> , 2021, , .	1.4	15
16	Development of sustainable research excellence with a global perspective on infectious diseases: Centre de Recherches Médicales de Lambaréon (CERMEL), Gabon. <i>Wiener Klinische Wochenschrift</i> , 2021, 133, 500-508.	1.9	14
17	COVID-19: A PCR-defined pandemic. <i>International Journal of Infectious Diseases</i> , 2021, 103, 278-279.	3.3	17
18	Predominance of HBV Genotype B and HDV Genotype 1 in Vietnamese Patients with Chronic Hepatitis. <i>Viruses</i> , 2021, 13, 346.	3.3	7

#	ARTICLE	IF	CITATIONS
19	Pharmacogenetic considerations in the treatment of co-infections with HIV/AIDS, tuberculosis and malaria in Congolese populations of Central Africa. <i>International Journal of Infectious Diseases</i> , 2021, 104, 207-213.	3.3	5
20	Genetic Diversity of Enteric Viruses in Children under Five Years Old in Gabon. <i>Viruses</i> , 2021, 13, 545.	3.3	6
21	Viral and serological testing of SARS-CoV-2 among health care workers and patients in Vietnam. <i>The Lancet Regional Health - Western Pacific</i> , 2021, 8, 100113.	2.9	5
22	Impact of <i>VSIG4</i> gene polymorphisms on susceptibility and functional status of rheumatoid arthritis. <i>International Journal of Immunogenetics</i> , 2021, 48, 260-265.	1.8	3
23	Genetic variants of programmed cell death 1 are associated with HBV infection and liver disease progression. <i>Scientific Reports</i> , 2021, 11, 7772.	3.3	5
24	Genomic surveillance of SARS-CoV-2 in the Republic of Congo. <i>International Journal of Infectious Diseases</i> , 2021, 105, 735-738.	3.3	19
25	Hepatitis E: An update on One Health and clinical medicine. <i>Liver International</i> , 2021, 41, 1462-1473.	3.9	63
26	COVID-19 and syndemic challenges in "Battling the Big Three": HIV, TB and malaria. <i>International Journal of Infectious Diseases</i> , 2021, 106, 29-32.	3.3	19
27	Swarm Learning for decentralized and confidential clinical machine learning. <i>Nature</i> , 2021, 594, 265-270.	27.8	375
28	Maternal Vaginal Colonization and Extended-Spectrum Beta-Lactamase-Producing Bacteria in Vietnamese Pregnant Women. <i>Antibiotics</i> , 2021, 10, 572.	3.7	4
29	A call to caution when hydroxychloroquine is given to elderly patients with COVID-19. <i>International Journal of Infectious Diseases</i> , 2021, 106, 265-268.	3.3	1
30	A global metagenomic map of urban microbiomes and antimicrobial resistance. <i>Cell</i> , 2021, 184, 3376-3393.e17.	28.9	164
31	Pharmacogene Sequencing of a Gabonese Population with Severe <i>Plasmodium falciparum</i> Malaria Reveals Multiple Novel Variants with Putative Relevance for Antimalarial Treatment. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0027521.	3.2	6
32	Longitudinal monitoring of laboratory markers characterizes hospitalized and ambulatory COVID-19 patients. <i>Scientific Reports</i> , 2021, 11, 14471.	3.3	15
33	Prevalence of urogenital and intestinal schistosomiasis among school children in South-west Nigeria. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009628.	3.0	16
34	Host factors facilitating SARS-CoV-2 virus infection and replication in the lungs. <i>Cellular and Molecular Life Sciences</i> , 2021, 78, 5953-5976.	5.4	19
35	Diagnosis of Chikungunya Virus in Febrile Patients From a Malaria Holoendemic Area. <i>International Journal of Infectious Diseases</i> , 2021, 109, 247-252.	3.3	9
36	Association Between Soluble Notch Ligand Delta-like Ligand 1 and Bleeding Complications in Patients With Dengue Fever Infection. <i>Journal of Infectious Diseases</i> , 2021, , .	4.0	1

#	ARTICLE	IF	CITATIONS
37	Safety and immunogenicity of an mRNA-lipid nanoparticle vaccine candidate against SARS-CoV-2. Wiener Klinische Wochenschrift, 2021, 133, 931-941.	1.9	79
38	Molecular detection of blaCTX-M gene to predict phenotypic cephalosporin resistance and clinical outcome of Escherichia coli bloodstream infections in Vietnam. Annals of Clinical Microbiology and Antimicrobials, 2021, 20, 60.	3.8	4
39	How to (ab)use a COVID-19 antigen rapid test with soft drinks?. International Journal of Infectious Diseases, 2021, 111, 28-30.	3.3	7
40	Low Prevalence of HEV Infection and No Associated Risk of HEV Transmission from Mother to Child among Pregnant Women in Vietnam. Pathogens, 2021, 10, 1340.	2.8	1
41	Host genetic factors determining COVID-19 susceptibility and severity. EBioMedicine, 2021, 72, 103629.	6.1	126
42	Molecular surveillance and genetic divergence of rotavirus A antigenic epitopes in Gabonese children with acute gastroenteritis. EBioMedicine, 2021, 73, 103648.	6.1	6
43	Absence of Plasmodium falciparum artemisinin resistance gene mutations eleven years after the adoption of artemisinin-based combination therapy in Nigeria. Malaria Journal, 2021, 20, 434.	2.3	3
44	Neopterin levels and Kyn/Trp ratios were significantly increased in dengue virus patients and subsequently decreased after recovery. International Journal of Infectious Diseases, 2020, 91, 162-168.	3.3	17
45	The COVID-19 epidemic. Tropical Medicine and International Health, 2020, 25, 278-280.	2.3	1,831
46	Analysis of sulphadoxine-pyrimethamine resistance-associated mutations in Plasmodium falciparum isolates obtained from asymptomatic pregnant women in Ogun State, Southwest Nigeria. Infection, Genetics and Evolution, 2020, 85, 104503.	2.3	12
47	Hepatitis E Virus Infection: Circulation, Molecular Epidemiology, and Impact on Global Health. Pathogens, 2020, 9, 856.	2.8	63
48	Asymptomatic SARS Coronavirus 2 infection: Invisible yet invincible. International Journal of Infectious Diseases, 2020, 100, 112-116.	3.3	177
49	Predominant secondary dengue infection among Vietnamese adults mostly without warning signs and severe disease. International Journal of Infectious Diseases, 2020, 100, 316-323.	3.3	6
50	Complement protein levels and MBL2 polymorphisms are associated with dengue and disease severity. Scientific Reports, 2020, 10, 14923.	3.3	9
51	Antimicrobial resistance preparedness in sub-Saharan African countries. Antimicrobial Resistance and Infection Control, 2020, 9, 145.	4.1	64
52	Natural killer cell receptor variants and chronic hepatitis B virus infection in the Vietnamese population. International Journal of Infectious Diseases, 2020, 96, 541-547.	3.3	11
53	Molecular surveillance of the Pfmdr1 N86Y allele among Congolese pregnant women with asymptomatic malaria. Malaria Journal, 2020, 19, 178.	2.3	3
54	Herd immunity and vaccination of children for COVID-19. International Journal of Infectious Diseases, 2020, 98, 14-15.	3.3	51

#	ARTICLE	IF	CITATIONS
55	Cartography of opportunistic pathogens and antibiotic resistance genes in a tertiary hospital environment. <i>Nature Medicine</i> , 2020, 26, 941-951.	30.7	130
56	Upregulation of Enzymes involved in ISGylation and Ubiquitination in patients with hepatocellular carcinoma. <i>International Journal of Medical Sciences</i> , 2020, 17, 347-353.	2.5	9
57	2019-nCoV in context: lessons learned?. <i>Lancet Planetary Health</i> , The, 2020, 4, e87-e88.	11.4	59
58	Upregulation of SMYD3 and SMYD3 VNTR 3/3 polymorphism increase the risk of hepatocellular carcinoma. <i>Scientific Reports</i> , 2020, 10, 2797.	3.3	6
59	Association of FCN2 polymorphisms and Ficolin-2 levels with dengue fever in Vietnamese patients. <i>International Journal of Infectious Diseases</i> , 2020, 95, 253-261.	3.3	8
60	A common polymorphism in the mechanosensitive ion channel <i>PIEZO1</i> is associated with protection from severe malaria in humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 9074-9081.	7.1	43
61	Mild versus severe COVID-19: Laboratory markers. <i>International Journal of Infectious Diseases</i> , 2020, 95, 304-307.	3.3	424
62	Hepatitis B Virus Infection Among Leprosy Patients: A Case for Polymorphisms Compromising Activation of the Lectin Pathway and Complement Receptors. <i>Frontiers in Immunology</i> , 2020, 11, 574457.	4.8	4
63	White Paper: Bridging the gap between human and animal surveillance data, antibiotic policy and stewardship in the hospital sector—practical guidance from the JPIAMR ARCH and COMBACTE-MAGNET EPI-Net networks. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, ii20-ii32.	3.0	13
64	White Paper: Bridging the gap between surveillance data and antimicrobial stewardship in the outpatient sector—practical guidance from the JPIAMR ARCH and COMBACTE-MAGNET EPI-Net networks. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, ii42-ii51.	3.0	12
65	White Paper: Bridging the gap between surveillance data and antimicrobial stewardship in the animal sector—practical guidance from the JPIAMR ARCH and COMBACTE-MAGNET EPI-Net networks. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, ii52-ii66.	3.0	7
66	Molecular surveillance and temporal monitoring of malaria parasites in focal Vietnamese provinces. <i>Malaria Journal</i> , 2020, 19, 458.	2.3	8
67	Rapid, low cost and sensitive detection of Calreticulin mutations by a PCR based amplicon length differentiation assay for diagnosis of myeloproliferative neoplasms. <i>BMC Medical Genetics</i> , 2019, 20, 115.	2.1	3
68	High Hepatitis E virus (HEV) Positivity Among Domestic Pigs and Risk of HEV Infection of Individuals Occupationally Exposed to Pigs and Pork Meat in Hanoi, Vietnam. <i>Open Forum Infectious Diseases</i> , 2019, 6, ofz306.	0.9	23
69	Analysis of <i>Plasmodium falciparum</i> Pfcr1 and Pfmdr1 genes in parasite isolates from asymptomatic individuals in Southeast Nigeria 11 years after withdrawal of chloroquine. <i>Malaria Journal</i> , 2019, 18, 343.	2.3	16
70	DNA recovery from archived RDTs for genetic characterization of <i>Plasmodium falciparum</i> in a routine setting in Lambaré, Gabon. <i>Malaria Journal</i> , 2019, 18, 336.	2.3	11
71	Distribution of the cytochrome P450 CYP2C8*2 allele in Brazzaville, Republic of Congo. <i>International Journal of Infectious Diseases</i> , 2019, 85, 49-53.	3.3	6
72	An alternative dogma on reduced artemisinin susceptibility: A new shadow from east to west. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12611-12612.	7.1	14

#	ARTICLE	IF	CITATIONS
73	Molecular surveillance of pfhrp2 and pfhrp3 genes deletion in Plasmodium falciparum isolates and the implications for rapid diagnostic tests in Nigeria. <i>Acta Tropica</i> , 2019, 196, 121-125.	2.0	34
74	No expression of HBV-human chimeric fusion transcript (HBx-LINE1) among Vietnamese patients with HBV-associated hepatocellular carcinoma. <i>Annals of Hepatology</i> , 2019, 18, 404-405.	1.5	3
75	Genetic diversity and population structure of <i>Plasmodium falciparum</i> in Kenyan-Ugandan border areas. <i>Tropical Medicine and International Health</i> , 2019, 24, 647-656.	2.3	10
76	An update on glucose-6-phosphate dehydrogenase deficiency in children from Brazzaville, Republic of Congo. <i>Malaria Journal</i> , 2019, 18, 57.	2.3	13
77	Cytochrome P450 CYP2B6*6 distribution among Congolese individuals with HIV, Tuberculosis and Malaria infection. <i>International Journal of Infectious Diseases</i> , 2019, 82, 111-116.	3.3	7
78	NTCP S267F variant associates with decreased susceptibility to HBV and HDV infection and decelerated progression of related liver diseases. <i>International Journal of Infectious Diseases</i> , 2019, 80, 147-152.	3.3	23
79	<i>Plasmodium falciparum</i> histidine-rich protein (PfHRP2 and 3) diversity in Western and Coastal Kenya. <i>Scientific Reports</i> , 2019, 9, 1709.	3.3	28
80	Monitoring of efficacy, tolerability and safety of artemether-lumefantrine and artesunate-amodiaquine for the treatment of uncomplicated <i>Plasmodium falciparum</i> malaria in Lambaré, Gabon: an open-label clinical trial. <i>Malaria Journal</i> , 2019, 18, 424.	2.3	18
81	Complement 5a Receptor Polymorphisms Are Associated With Panton-Valentine Leukocidin-positive <i>Staphylococcus aureus</i> Colonization in African Pygmies. <i>Clinical Infectious Diseases</i> , 2019, 68, 854-856.	5.8	6
82	Epstein-Barr virus, malaria and endemic Burkitt lymphoma. <i>EBioMedicine</i> , 2019, 39, 13-14.	6.1	6
83	The blood transcriptome of childhood malaria. <i>EBioMedicine</i> , 2019, 40, 614-625.	6.1	47
84	Vitamin D receptor Apal polymorphism associated with progression of liver disease in Vietnamese patients chronically infected with hepatitis B virus. <i>BMC Medical Genetics</i> , 2019, 20, 201.	2.1	22
85	Silencing of Kangai 1 C-terminal interacting tetraspanin suppresses progression of cholangiocarcinoma. <i>Experimental Cell Research</i> , 2018, 364, 59-67.	2.6	1
86	High prevalence of dihydrofolate reductase gene mutations in <i>Plasmodium falciparum</i> parasites among pregnant women in Nigeria after reported use of sulfadoxine-pyrimethamine. <i>Pathogens and Global Health</i> , 2018, 112, 86-92.	2.3	6
87	Human complement receptor type 1 (CR1) protein levels and genetic variants in chronic Chagas Disease. <i>Scientific Reports</i> , 2018, 8, 526.	3.3	9
88	Efficacy and Safety of Fosmidomycin-Piperaquine as Nonartemisinin-Based Combination Therapy for Uncomplicated <i>Falciparum</i> Malaria: A Single-Arm, Age De-escalation Proof-of-Concept Study in Gabon. <i>Clinical Infectious Diseases</i> , 2018, 66, 1823-1830.	5.8	41
89	KIR-HLA distribution in a Vietnamese population from Hanoi. <i>Human Immunology</i> , 2018, 79, 93-100.	2.4	6
90	Differential contribution of interleukin-10 promoter variants in malaria and schistosomiasis mono- and co-infections among Nigerian children. <i>Tropical Medicine and International Health</i> , 2018, 23, 45-52.	2.3	9

#	ARTICLE	IF	CITATIONS
91	Soluble fibrinogen-like protein 2 levels in patients with hepatitis B virus-related liver diseases. BMC Infectious Diseases, 2018, 18, 553.	2.9	11
92	Vitamin D deficiency and hepatitis viruses-associated liver diseases: A literature review. World Journal of Gastroenterology, 2018, 24, 445-460.	3.3	40
93	HDV infection rates in northern Vietnam. Scientific Reports, 2018, 8, 8047.	3.3	10
94	Complement receptor 1 (CR1, CD35) association with susceptibility to leprosy. PLoS Neglected Tropical Diseases, 2018, 12, e0006705.	3.0	19
95	PfHRP2-PfHRP3 diversity among Kenyan isolates and comparative evaluation of PfHRP2/pLDH malaria RDT with microscopy and nested PCR methodologies. Parasitology International, 2018, 67, 793-799.	1.3	15
96	Optimisation of quantitative miRNA panels to consolidate the diagnostic surveillance of HBV-related hepatocellular carcinoma. PLoS ONE, 2018, 13, e0196081.	2.5	12
97	Effect of AdipoR agonist in cholangiocarcinoma.. Journal of Clinical Oncology, 2018, 36, 323-323.	1.6	2
98	Interferon-stimulated gene 20 kDa protein serum levels and clinical outcome of hepatitis B virus-related liver diseases. Oncotarget, 2018, 9, 27858-27871.	1.8	5
99	Genetic variants of interferon regulatory factor 5 associated with chronic hepatitis B infection. World Journal of Gastroenterology, 2018, 24, 248-256.	3.3	4
100	Genetic Diversity of the Plasmodium falciparum Glutamate-Rich Protein R2 Region Before and Twelve Years after Introduction of Artemisinin Combination Therapies among Febrile Children in Nigeria. American Journal of Tropical Medicine and Hygiene, 2018, 98, 667-676.	1.4	6
101	Pyruvate Kinase and Fcγ3 Receptor Gene Copy Numbers Associated With Malaria Phenotypes. Journal of Infectious Diseases, 2017, 216, 276-282.	4.0	12
102	Parasite Infection, Carcinogenesis and Human Malignancy. EBioMedicine, 2017, 15, 12-23.	6.1	108
103	Mannose-binding lectin deficiency and miscarriages in rheumatoid arthritis. Autoimmunity, 2017, 50, 409-413.	2.6	7
104	Kangai 1 C-terminal interacting tetraspanin plays an important role in cholangiocarcinogenesis. Journal of Hepatology, 2017, 66, S460.	3.7	0
105	Human genetic factors in tuberculosis: an update. Tropical Medicine and International Health, 2017, 22, 1063-1071.	2.3	53
106	Molecular surveillance of Plasmodium falciparum drug resistance in the Republic of Congo: four and nine years after the introduction of artemisinin-based combination therapy. Malaria Journal, 2017, 16, 155.	2.3	22
107	Molecular markers of anti-malarial drug resistance in Central, West and East African children with severe malaria. Malaria Journal, 2017, 16, 217.	2.3	20
108	Adiponectin and pro-inflammatory cytokines are modulated in Vietnamese patients with type 2 diabetes mellitus. Journal of Diabetes Investigation, 2017, 8, 295-305.	2.4	45

#	ARTICLE	IF	CITATIONS
109	Prevalence and genotype distribution of hepatitis delta virus among chronic hepatitis B carriers in Central Vietnam. PLoS ONE, 2017, 12, e0175304.	2.5	16
110	Geographical distribution of complement receptor type 1 variants and their associated disease risk. PLoS ONE, 2017, 12, e0175973.	2.5	9
111	<i>SOCS3</i> genetic variants and promoter hypermethylation in patients with chronic hepatitis B. Oncotarget, 2017, 8, 17127-17139.	1.8	24
112	Intramuscular Artesunate for Severe Malaria in African Children: A Multicenter Randomized Controlled Trial. PLoS Medicine, 2016, 13, e1001938.	8.4	44
113	Molecular epidemiology and surveillance of circulating rotavirus and adenovirus in Congolese children with gastroenteritis. Journal of Medical Virology, 2016, 88, 596-605.	5.0	39
114	Glucose-6-phosphate dehydrogenase deficiency and reduced haemoglobin levels in African children with severe malaria. Malaria Journal, 2016, 15, 346.	2.3	12
115	Triggering receptor expressed on myeloid cells 1 (TREM1) and cytokine gene variants in complicated and uncomplicated malaria. Tropical Medicine and International Health, 2016, 21, 1592-1601.	2.3	20
116	Characterization of <i>Vibrio cholerae</i> Strains Isolated from the Nigerian Cholera Outbreak in 2010. Journal of Clinical Microbiology, 2016, 54, 2618-2621.	3.9	7
117	Hepatitis E Virus Mutations: Functional and Clinical Relevance. EBioMedicine, 2016, 11, 31-42.	6.1	52
118	Association of vitamin D deficiency with hepatitis B virus - related liver diseases. BMC Infectious Diseases, 2016, 16, 507.	2.9	44
119	Enrichment of bacterial DNA for the diagnosis of blood stream infections. BMC Infectious Diseases, 2016, 16, 235.	2.9	22
120	Molecular epidemiology of hepatitis D virus circulating in Southwestern Nigeria. Virology Journal, 2016, 13, 61.	3.4	30
121	Host-directed therapies for infectious diseases: current status, recent progress, and future prospects. Lancet Infectious Diseases, The, 2016, 16, e47-e63.	9.1	265
122	Interferon-stimulated gene 15 in hepatitis B-related liver diseases. Oncotarget, 2016, 7, 67777-67787.	1.8	14
123	Hepatitis E Virus Superinfection and Clinical Progression in Hepatitis B Patients. EBioMedicine, 2015, 2, 2080-2086.	6.1	46
124	Low MBL-associated serine protease 2 (MASP2) levels correlate with urogenital schistosomiasis in Nigerian children. Tropical Medicine and International Health, 2015, 20, 1311-1319.	2.3	2
125	Association of Ficolin-2 Serum Levels and FCN2 Genetic Variants with Indian Visceral Leishmaniasis. PLoS ONE, 2015, 10, e0125940.	2.5	9
126	Correlation of Interleukin-6 levels and lectins during <i>Schistosoma haematobium</i> infection. Cytokine, 2015, 76, 152-155.	3.2	10

#	ARTICLE	IF	CITATIONS
127	FOXO3A regulatory polymorphism and susceptibility to severe malaria in Gabonese children. <i>Immunogenetics</i> , 2015, 67, 67-71.	2.4	14
128	Opisthorchiasis: An Overlooked Danger. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003563.	3.0	36
129	Lectin Complement Protein Collectin 11 (CL-K1) and Susceptibility to Urinary Schistosomiasis. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003647.	3.0	18
130	Soluble MICB protein levels and platelet counts during hepatitis B virus infection and response to hepatocellular carcinoma treatment. <i>BMC Infectious Diseases</i> , 2015, 15, 25.	2.9	8
131	Simple multiplex PCR assays to detect common pathogens and associated genes encoding for acquired extended spectrum betalactamases (ESBL) or carbapenemases from surgical site specimens in Vietnam. <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2015, 14, 23.	3.8	24
132	Mannose-binding Lectin (MBL) as a susceptible host factor influencing Indian Visceral Leishmaniasis. <i>Parasitology International</i> , 2015, 64, 591-596.	1.3	11
133	Identification of a natural intergenotypic recombinant hepatitis delta virus genotype 1 and 2 in Vietnamese HBsAg-positive patients. <i>Journal of Viral Hepatitis</i> , 2015, 22, 55-63.	2.0	23
134	Occult Hepatitis B Virus Infection in Nigerian Blood Donors and Hepatitis B Virus Transmission Risks. <i>PLoS ONE</i> , 2015, 10, e0131912.	2.5	57
135	Mannose Binding Lectin and Susceptibility to Rheumatoid Arthritis in Brazilian Patients and Their Relatives. <i>PLoS ONE</i> , 2014, 9, e95519.	2.5	17
136	A reliable and rapid method for molecular detection of malarial parasites using microwave irradiation and loop mediated isothermal amplification. <i>Malaria Journal</i> , 2014, 13, 454.	2.3	26
137	Analysis of genetic variants in the IL4 promoter and VNTR loci in Indian patients with Visceral Leishmaniasis. <i>Human Immunology</i> , 2014, 75, 1177-1181.	2.4	8
138	Return of chloroquine-sensitive Plasmodium falciparum parasites and emergence of chloroquine-resistant Plasmodium vivax in Ethiopia. <i>Malaria Journal</i> , 2014, 13, 244.	2.3	67
139	MBL2 Variations and Malaria Susceptibility in Indian Populations. <i>Infection and Immunity</i> , 2014, 82, 52-61.	2.2	30
140	Re-evaluation of microscopy confirmed Plasmodium falciparum and Plasmodium vivax malaria by nested PCR detection in southern Ethiopia. <i>Malaria Journal</i> , 2014, 13, 48.	2.3	42
141	Differential ability to resist to complement lysis and invade host cells mediated by MBL in R4 and 860 strains of Trypanosoma cruzi. <i>FEBS Letters</i> , 2014, 588, 956-961.	2.8	14
142	Analysis of polymorphic sites in the promoter of the nitric oxide synthase 2 gene in Brazilian patients with leprosy. <i>International Journal of Immunogenetics</i> , 2014, 41, 231-235.	1.8	2
143	Genetic insights on host and hepatitis B virus in liver diseases. <i>Mutation Research - Reviews in Mutation Research</i> , 2014, 762, 65-75.	5.5	18
144	Genetic evidence of regulatory gene variants of the STAT6, IL10R and FOXP3 locus as a susceptibility factor in uncomplicated malaria and parasitaemia in Congolese children. <i>Malaria Journal</i> , 2013, 12, 9.	2.3	21

#	ARTICLE	IF	CITATIONS
145	A trivial role of STAT4 variant in chronic hepatitis B induced hepatocellular carcinoma. <i>Infection, Genetics and Evolution</i> , 2013, 18, 257-261.	2.3	20
146	Association of IP-10 and PDGF-BB levels with clinical forms of chronic Chagas disease. <i>International Journal of Cardiology</i> , 2013, 169, e53-e55.	1.7	12
147	Co-infection of human parvovirus B19 with <i>Plasmodium falciparum</i> contributes to malaria disease severity in Gabonese patients. <i>BMC Infectious Diseases</i> , 2013, 13, 375.	2.9	21
148	Genetic evidence of TAP1 gene variant as a susceptibility factor in Indian leprosy patients. <i>Human Immunology</i> , 2013, 74, 803-807.	2.4	18
149	Mannose-Binding Lectin and Susceptibility to Schistosomiasis. <i>Journal of Infectious Diseases</i> , 2013, 207, 1675-1683.	4.0	24
150	Hepatitis B virus-induced hepatocellular carcinoma: functional roles of MICA variants. <i>Journal of Viral Hepatitis</i> , 2013, 20, 687-698.	2.0	61
151	Association of L-Ficolin Levels and FCN2 Genotypes with Chronic Chagas Disease. <i>PLoS ONE</i> , 2013, 8, e60237.	2.5	38
152	LRRK2 and RIPK2 Variants in the NOD 2-Mediated Signaling Pathway Are Associated with Susceptibility to <i>Mycobacterium leprae</i> in Indian Populations. <i>PLoS ONE</i> , 2013, 8, e73103.	2.5	45
153	High Prevalence and Significance of Hepatitis D Virus Infection among Treatment-Naïve HBsAg-Positive Patients in Northern Vietnam. <i>PLoS ONE</i> , 2013, 8, e78094.	2.5	30
154	Ficolin-2 Levels and FCN2 Genetic Polymorphisms as a Susceptibility Factor in Schistosomiasis. <i>Journal of Infectious Diseases</i> , 2012, 206, 562-570.	4.0	41
155	Reliable and rapid characterization of functional FCN2 gene variants reveals diverse geographical patterns. <i>BMC Medical Genetics</i> , 2012, 13, 37.	2.1	17
156	In-vitro characterization of novel and functional regulatory SNPs in the promoter region of IL2 and IL2R alpha in a Gabonese population. <i>BMC Medical Genetics</i> , 2012, 13, 117.	2.1	4
157	Genetic Evidence of Functional Ficolin-2 Haplotype as Susceptibility Factor in Cutaneous Leishmaniasis. <i>PLoS ONE</i> , 2012, 7, e34113.	2.5	35
158	Combined promoter haplotypes of the IL10R genes are associated with protection against severe malaria in Gabonese children. <i>Immunogenetics</i> , 2012, 64, 87-95.	2.4	15
159	Association of CISH -292A/T genetic variant with hepatitis B virus infection. <i>Immunogenetics</i> , 2012, 64, 261-265.	2.4	14
160	IL-4 Haplotype -590T, -34T and Intron-3 VNTR R2 Is Associated with Reduced Malaria Risk among Ancestral Indian Tribal Populations. <i>PLoS ONE</i> , 2012, 7, e48136.	2.5	26
161	Molecular characterization of regulatory polymorphisms in the promoter region of the STAT6 gene in a Gabonese population. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2011, 106, 65-69.	1.6	7
162	Novel regulatory SNPs in the promoter region of the TNFRSF18 gene in a Gabonese population. <i>Brazilian Journal of Medical and Biological Research</i> , 2011, 44, 418-420.	1.5	8

#	ARTICLE	IF	CITATIONS
163	Ficolin-2 Levels and FCN2 Haplotypes Influence Hepatitis B Infection Outcome in Vietnamese Patients. PLoS ONE, 2011, 6, e28113.	2.5	44
164	Olfactory neuron-specific expression of A30P alpha-synuclein exacerbates dopamine deficiency and hyperactivity in a novel conditional model of early Parkinson's disease stages. Neurobiology of Disease, 2011, 44, 192-204.	4.4	28
165	Novel and functional regulatory SNPs in the promoter region of FOXP3 gene in a Gabonese population. Immunogenetics, 2011, 63, 409-415.	2.4	26
166	Regulatory T Cells and Parasites. Journal of Biomedicine and Biotechnology, 2011, 2011, 1-8.	3.0	32
167	Detection of multiple infections by <i>Monocystis</i> strains in a single earthworm host using ribosomal internal transcribed spacer sequence variation. Parasitology, 2010, 137, 45-51.	1.5	6
168	Variant alleles of the mannose binding lectin 2 gene (<i>MBL2</i>) confer heterozygote advantage within Crohn's families. Scandinavian Journal of Gastroenterology, 2010, 45, 1129-1130.	1.5	6
169	High genetic diversity and heterogeneous parasite load in the earthworm <i>Lumbricus terrestris</i> on a German meadow. Soil Biology and Biochemistry, 2009, 41, 1591-1595.	8.8	15
170	Development and characterization of ten novel microsatellite markers from olive ridley sea turtle (<i>Lepidochelys olivacea</i>). Conservation Genetics, 2008, 9, 981-984.	1.5	13
171	Microsatellite markers for the drought-resistant earthworm <i>Hormogaster elisae</i> . Molecular Ecology Resources, 2008, 8, 901-903.	4.8	17
172	A New Frog Species from the Central Western Ghats of India, and Its Phylogenetic Position. Zoological Science, 2007, 24, 525-534.	0.7	11
173	Genetic diversity and parasite defense in a fragmented urban metapopulation of earthworms. Animal Conservation, 2007, 10, 162-175.	2.9	24
174	Development and characterization of novel microsatellite markers for the common earthworm (<i>Lumbricus terrestris</i> L.). Molecular Ecology Notes, 2007, 7, 1060-1062.	1.7	28
175	Different population histories of the Mundari- and Mon-Khmer-speaking Austro-Asiatic tribes inferred from the mtDNA 9-bp deletion/insertion polymorphism in Indian populations. Human Genetics, 2006, 119, 223-224.	3.8	17
176	Different population histories of the Mundari- and Mon-Khmer-speaking Austro-Asiatic tribes inferred from the mtDNA 9-bp deletion/insertion polymorphism in Indian populations. Human Genetics, 2005, 116, 507-517.	3.8	60
177	Development and characterization of novel microsatellite markers from the olive ridley sea turtle (<i>Lepidochelys olivacea</i>). Molecular Ecology Notes, 2004, 4, 77-79.	1.7	38