Thirumalaisamy P Velavan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The COVIDâ€19 epidemic. Tropical Medicine and International Health, 2020, 25, 278-280.	2.3	1,831
2	Mild versus severe COVID-19: Laboratory markers. International Journal of Infectious Diseases, 2020, 95, 304-307.	3.3	424
3	Swarm Learning for decentralized and confidential clinical machine learning. Nature, 2021, 594, 265-270.	27.8	375
4	Host-directed therapies for infectious diseases: current status, recent progress, and future prospects. Lancet Infectious Diseases, The, 2016, 16, e47-e63.	9.1	265
5	Asymptomatic SARS Coronavirus 2 infection: Invisible yet invincible. International Journal of Infectious Diseases, 2020, 100, 112-116.	3.3	177
6	A global metagenomic map of urban microbiomes and antimicrobial resistance. Cell, 2021, 184, 3376-3393.e17.	28.9	164
7	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. International Journal of Infectious Diseases, 2022, 114, 268-272.	3.3	136
8	Cartography of opportunistic pathogens and antibiotic resistance genes in a tertiary hospital environment. Nature Medicine, 2020, 26, 941-951.	30.7	130
9	Host genetic factors determining COVID-19 susceptibility and severity. EBioMedicine, 2021, 72, 103629.	6.1	126
10	Monkeypox 2022 outbreak: An update. Tropical Medicine and International Health, 2022, 27, 604-605.	2.3	114
11	Parasite Infection, Carcinogenesis and Human Malignancy. EBioMedicine, 2017, 15, 12-23.	6.1	108
12	Safety and immunogenicity of an mRNA-lipid nanoparticle vaccine candidate against SARS-CoV-2. Wiener Klinische Wochenschrift, 2021, 133, 931-941.	1.9	79
13	Return of chloroquine-sensitive Plasmodium falciparum parasites and emergence of chloroquine-resistant Plasmodium vivax in Ethiopia. Malaria Journal, 2014, 13, 244.	2.3	67
14	Antimicrobial resistance preparedness in sub-Saharan African countries. Antimicrobial Resistance and Infection Control, 2020, 9, 145.	4.1	64
15	Hepatitis E Virus Infection: Circulation, Molecular Epidemiology, and Impact on Global Health. Pathogens, 2020, 9, 856.	2.8	63
16	Hepatitis E: An update on One Health and clinical medicine. Liver International, 2021, 41, 1462-1473.	3.9	63
17	Hepatitis <scp>B</scp> virusâ€induced hepatocellular carcinoma: functional roles of <i><scp>MICA</scp></i> variants. Journal of Viral Hepatitis, 2013, 20, 687-698.	2.0	61
18	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

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19	2019-nCoV in context: lessons learned?. Lancet Planetary Health, The, 2020, 4, e87-e88.	11.4	59
20	Occult Hepatitis B Virus Infection in Nigerian Blood Donors and Hepatitis B Virus Transmission Risks. PLoS ONE, 2015, 10, e0131912.	2.5	57
21	Human genetic factors in tuberculosis: an update. Tropical Medicine and International Health, 2017, 22, 1063-1071.	2.3	53
22	Hepatitis E Virus Mutations: Functional and Clinical Relevance. EBioMedicine, 2016, 11, 31-42.	6.1	52
23	Herd immunity and vaccination of children for COVID-19. International Journal of Infectious Diseases, 2020, 98, 14-15.	3.3	51
24	The blood transcriptome of childhood malaria. EBioMedicine, 2019, 40, 614-625.	6.1	47
25	Hepatitis E Virus Superinfection and Clinical Progression in Hepatitis B Patients. EBioMedicine, 2015, 2, 2080-2086.	6.1	46
26	LRRK2 and RIPK2 Variants in the NOD 2-Mediated Signaling Pathway Are Associated with Susceptibility to Mycobacterium leprae in Indian Populations. PLoS ONE, 2013, 8, e73103.	2.5	45
27	Adiponectin and proâ€inflammatory cytokines are modulated in Vietnamese patients with type 2 diabetes mellitus. Journal of Diabetes Investigation, 2017, 8, 295-305.	2.4	45
28	Ficolin-2 Levels and FCN2 Haplotypes Influence Hepatitis B Infection Outcome in Vietnamese Patients. PLoS ONE, 2011, 6, e28113.	2.5	44
29	Intramuscular Artesunate for Severe Malaria in African Children: A Multicenter Randomized Controlled Trial. PLoS Medicine, 2016, 13, e1001938.	8.4	44
30	Association of vitamin D deficiency with hepatitis B virus - related liver diseases. BMC Infectious Diseases, 2016, 16, 507.	2.9	44
31	A common polymorphism in the mechanosensitive ion channel <i>PIEZO1</i> is associated with protection from severe malaria in humans. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 9074-9081.	7.1	43
32	Re-evaluation of microscopy confirmed Plasmodium falciparum and Plasmodium vivax malaria by nested PCR detection in southern Ethiopia. Malaria Journal, 2014, 13, 48.	2.3	42
33	Ficolin-2 Levels and FCN2 Genetic Polymorphisms as a Susceptibility Factor in Schistosomiasis. Journal of Infectious Diseases, 2012, 206, 562-570.	4.0	41
34	Efficacy and Safety of Fosmidomycin–Piperaquine as Nonartemisinin-Based Combination Therapy for Uncomplicated Falciparum Malaria: A Single-Arm, Age De-escalation Proof-of-Concept Study in Gabon. Clinical Infectious Diseases, 2018, 66, 1823-1830.	5.8	41
35	Vitamin D deficiency and hepatitis viruses-associated liver diseases: A literature review. World Journal of Gastroenterology, 2018, 24, 445-460.	3.3	40
36	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

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37	Development and characterization of novel microsatellite markers from the olive ridley sea turtle (Lepidochelys olivacea). Molecular Ecology Notes, 2004, 4, 77-79.	1.7	38
38	Association of L-Ficolin Levels and FCN2 Genotypes with Chronic Chagas Disease. PLoS ONE, 2013, 8, e60237.	2.5	38
39	Hepatitis E virus genome detection in commercial pork livers and pork meat products in Germany. Journal of Viral Hepatitis, 2021, 28, 196-204.	2.0	37
40	Opisthorchiasis: An Overlooked Danger. PLoS Neglected Tropical Diseases, 2015, 9, e0003563.	3.0	36
41	Genetic Evidence of Functional Ficolin-2 Haplotype as Susceptibility Factor in Cutaneous Leishmaniasis. PLoS ONE, 2012, 7, e34113.	2.5	35
42	Molecular surveillance of pfhrp2 and pfhrp3 genes deletion in Plasmodium falciparum isolates and the implications for rapid diagnostic tests in Nigeria. Acta Tropica, 2019, 196, 121-125.	2.0	34
43	Regulatory T Cells and Parasites. Journal of Biomedicine and Biotechnology, 2011, 2011, 1-8.	3.0	32
44	<i>MBL2</i> Variations and Malaria Susceptibility in Indian Populations. Infection and Immunity, 2014, 82, 52-61.	2.2	30
45	Molecular epidemiology of hepatitis D virus circulating in Southwestern Nigeria. Virology Journal, 2016, 13, 61.	3.4	30
46	High Prevalence and Significance of Hepatitis D Virus Infection among Treatment-NaÃ ⁻ ve HBsAg-Positive Patients in Northern Vietnam. PLoS ONE, 2013, 8, e78094.	2.5	30
47	Development and characterization of novel microsatellite markers for the common earthworm (Lumbricus terrestris L.). Molecular Ecology Notes, 2007, 7, 1060-1062.	1.7	28
48	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
49	Plasmodium falciparum histidine-rich protein (PfHRP2 and 3) diversity in Western and Coastal Kenya. Scientific Reports, 2019, 9, 1709.	3.3	28
50	Novel and functional regulatory SNPs in the promoter region of FOXP3 gene in a Gabonese population. Immunogenetics, 2011, 63, 409-415.	2.4	26
51	A reliable and rapid method for molecular detection of malarial parasites using microwave irradiation and loop mediated isothermal amplification. Malaria Journal, 2014, 13, 454.	2.3	26
52	IL-4 Haplotype -590T, -34T and Intron-3 VNTR R2 Is Associated with Reduced Malaria Risk among Ancestral Indian Tribal Populations. PLoS ONE, 2012, 7, e48136.	2.5	26
53	Genetic diversity and parasite defense in a fragmented urban metapopulation of earthworms. Animal Conservation, 2007, 10, 162-175.	2.9	24
54	Mannose-Binding Lectin and Susceptibility to Schistosomiasis. Journal of Infectious Diseases, 2013, 207, 1675-1683.	4.0	24

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55	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. Annals of Clinical Microbiology and Antimicrobials, 2015, 14, 23.	3.8	24
56	COVID-19 in Africa: between hope and reality. Lancet Infectious Diseases, The, 2021, 21, 315.	9.1	24
57	<i>SOCS3</i> genetic variants and promoter hypermethylation in patients with chronic hepatitis B. Oncotarget, 2017, 8, 17127-17139.	1.8	24
58	Identification of a natural intergenotypic recombinant hepatitis delta virus genotype 1 and 2 in <scp>V</scp> ietnamese <scp>HB</scp> sAgâ€positive patients. Journal of Viral Hepatitis, 2015, 22, 55-63.	2.0	23
59	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. Open Forum Infectious Diseases, 2019, 6, ofz306.	0.9	23
60	NTCP S267F variant associates with decreased susceptibility to HBV and HDV infection and decelerated progression of related liver diseases. International Journal of Infectious Diseases, 2019, 80, 147-152.	3.3	23
61	Enrichment of bacterial DNA for the diagnosis of blood stream infections. BMC Infectious Diseases, 2016, 16, 235.	2.9	22
62	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
63	Vitamin D receptor Apal polymorphism associated with progression of liver disease in Vietnamese patients chronically infected with hepatitis B virus. BMC Medical Genetics, 2019, 20, 201.	2.1	22
64	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. Malaria Journal, 2013, 12, 9.	2.3	21
65	Co-infection of human parvovirus B19 with Plasmodium falciparum contributes to malaria disease severity in Gabonese patients. BMC Infectious Diseases, 2013, 13, 375.	2.9	21
66	A trivial role of STAT4 variant in chronic hepatitis B induced hepatocellular carcinoma. Infection, Genetics and Evolution, 2013, 18, 257-261.	2.3	20
67	Triggering receptor expressed on myeloid cells 1 (<scp>TREM</scp> â€1) and cytokine gene variants in complicated malaria. Tropical Medicine and International Health, 2016, 21, 1592-1601.	2.3	20
68	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
69	Complement receptor 1 (CR1, CD35) association with susceptibility to leprosy. PLoS Neglected Tropical Diseases, 2018, 12, e0006705.	3.0	19
70	Genomic surveillance of SARS-CoV-2 in the Republic of Congo. International Journal of Infectious Diseases, 2021, 105, 735-738.	3.3	19
71	COVID-19 and syndemic challenges in â€~Battling the Big Three': HIV, TB and malaria. International Journal of Infectious Diseases, 2021, 106, 29-32.	3.3	19
72	Host factors facilitating SARSâ€CoVâ€2 virus infection and replication in the lungs. Cellular and Molecular Life Sciences, 2021, 78, 5953-5976.	5.4	19

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73	Genetic evidence of TAP1 gene variant as a susceptibility factor in Indian leprosy patients. Human Immunology, 2013, 74, 803-807.	2.4	18
74	Genetic insights on host and hepatitis B virus in liver diseases. Mutation Research - Reviews in Mutation Research, 2014, 762, 65-75.	5.5	18
75	Lectin Complement Protein Collectin 11 (CL-K1) and Susceptibility to Urinary Schistosomiasis. PLoS Neglected Tropical Diseases, 2015, 9, e0003647.	3.0	18
76	Monitoring of efficacy, tolerability and safety of artemether–lumefantrine and artesunate–amodiaquine for the treatment of uncomplicated Plasmodium falciparum malaria in Lambaréné, Gabon: an open-label clinical trial. Malaria Journal, 2019, 18, 424.	2.3	18
77	Targeting interleukin 6 signaling by monoclonal antibody siltuximab on cholangiocarcinoma. Journal of Gastroenterology and Hepatology (Australia), 2021, 36, 1334-1345.	2.8	18
78	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
79	Microsatellite markers for the droughtâ€resistant earthworm <i>Hormogaster elisae</i> . Molecular Ecology Resources, 2008, 8, 901-903.	4.8	17
80	Reliable and rapid characterization of functional FCN2 gene variants reveals diverse geographical patterns. BMC Medical Genetics, 2012, 13, 37.	2.1	17
81	Mannose Binding Lectin and Susceptibility to Rheumatoid Arthritis in Brazilian Patients and Their Relatives. PLoS ONE, 2014, 9, e95519.	2.5	17
82	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
83	COVID-19: A PCR-defined pandemic. International Journal of Infectious Diseases, 2021, 103, 278-279.	3.3	17
84	Analysis of Plasmodium falciparum Pfcrt and Pfmdr1 genes in parasite isolates from asymptomatic individuals in Southeast Nigeria 11Âyears after withdrawal of chloroquine. Malaria Journal, 2019, 18, 343.	2.3	16
85	Prevalence of urogenital and intestinal schistosomiasis among school children in South-west Nigeria. PLoS Neglected Tropical Diseases, 2021, 15, e0009628.	3.0	16
86	Prevalence and genotype distribution of hepatitis delta virus among chronic hepatitis B carriers in Central Vietnam. PLoS ONE, 2017, 12, e0175304.	2.5	16
87	High genetic diversity and heterogeneous parasite load in the earthworm Lumbricus terrestris on a German meadow. Soil Biology and Biochemistry, 2009, 41, 1591-1595.	8.8	15
88	Combined promoter haplotypes of the IL10R genes are associated with protection against severe malaria in Gabonese children. Immunogenetics, 2012, 64, 87-95.	2.4	15
89	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
90	Longitudinal Monitoring of Lactate in Hospitalized and Ambulatory COVID-19 Patients. American Journal of Tropical Medicine and Hygiene, 2021, , .	1.4	15

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91	Longitudinal monitoring of laboratory markers characterizes hospitalized and ambulatory COVID-19 patients. Scientific Reports, 2021, 11, 14471.	3.3	15
92	Association of CISH -292A/T genetic variant with hepatitis B virus infection. Immunogenetics, 2012, 64, 261-265.	2.4	14
93	Differential ability to resist to complement lysis and invade host cells mediated by MBL in R4 and 860 strains of <i>Trypanosoma cruzi</i> . FEBS Letters, 2014, 588, 956-961.	2.8	14
94	FOXO3A regulatory polymorphism and susceptibility to severe malaria in Gabonese children. Immunogenetics, 2015, 67, 67-71.	2.4	14
95	An alternative dogma on reduced artemisinin susceptibility: A new shadow from east to west. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 12611-12612.	7.1	14
96	Development of sustainable research excellence with a global perspective on infectious diseases: Centre de Recherches Médicales de Lambaréné (CERMEL), Gabon. Wiener Klinische Wochenschrift, 2021, 133, 500-508.	1.9	14
97	Interferon-stimulated gene 15 in hepatitis B-related liver diseases. Oncotarget, 2016, 7, 67777-67787.	1.8	14
98	Development and characterization of ten novel microsatellite markers from olive ridley sea turtle (Lepidochelys olivacea). Conservation Genetics, 2008, 9, 981-984.	1.5	13
99	An update on glucose-6-phosphate dehydrogenase deficiency in children from Brazzaville, Republic of Congo. Malaria Journal, 2019, 18, 57.	2.3	13
100	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. Journal of Antimicrobial Chemotherapy, 2020, 75, ii20-ii32.	3.0	13
101	Association of IP-10 and PDGF-BB levels with clinical forms of chronic Chagas disease. International Journal of Cardiology, 2013, 169, e53-e55.	1.7	12
102	Glucose-6-phosphate dehydrogenase deficiency and reduced haemoglobin levels in African children with severe malaria. Malaria Journal, 2016, 15, 346.	2.3	12
103	Pyruvate Kinase and Fcl ³ Receptor Gene Copy Numbers Associated With Malaria Phenotypes. Journal of Infectious Diseases, 2017, 216, 276-282.	4.0	12
104	Optimisation of quantitative miRNA panels to consolidate the diagnostic surveillance of HBV-related hepatocellular carcinoma. PLoS ONE, 2018, 13, e0196081.	2.5	12
105	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
106	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. Journal of Antimicrobial Chemotherapy, 2020, 75, ii42-ii51.	3.0	12
107	A New Frog Species from the Central Western Ghats of India, and Its Phylogenetic Position. Zoological Science, 2007, 24, 525-534.	0.7	11
108	Mannose-binding Lectin (MBL) as a susceptible host factor influencing Indian Visceral Leishmaniasis. Parasitology International, 2015, 64, 591-596.	1.3	11

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109	Soluble fibrinogen-like protein 2 levels in patients with hepatitis B virus-related liver diseases. BMC Infectious Diseases, 2018, 18, 553.	2.9	11
110	DNA recovery from archived RDTs for genetic characterization of Plasmodium falciparum in a routine setting in Lambaréné, Gabon. Malaria Journal, 2019, 18, 336.	2.3	11
111	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
112	High prevalence of antibiotic-resistant Escherichia coli in Congolese students. International Journal of Infectious Diseases, 2021, 103, 119-123.	3.3	11
113	Host genetic loci LZTFL1 and CCL2 associated with SARS-CoV-2 infection and severity of COVID-19. International Journal of Infectious Diseases, 2022, 122, 427-436.	3.3	11
114	Correlation of Interleukin-6 levels and lectins during Schistosoma haematobium infection. Cytokine, 2015, 76, 152-155.	3.2	10
115	HDV infection rates in northern Vietnam. Scientific Reports, 2018, 8, 8047.	3.3	10
116	Genetic diversity and population structure of <i>Plasmodium falciparum</i> in Kenyan–Ugandan border areas. Tropical Medicine and International Health, 2019, 24, 647-656.	2.3	10
117	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. International Journal of Infectious Diseases, 2022, 114, 151-154.	3.3	10
118	Association of Ficolin-2 Serum Levels and FCN2 Genetic Variants with Indian Visceral Leishmaniasis. PLoS ONE, 2015, 10, e0125940.	2.5	9
119	Human complement receptor type 1 (CR1) protein levels and genetic variants in chronic Chagas Disease. Scientific Reports, 2018, 8, 526.	3.3	9
120	Differential contribution of interleukinâ€10 promoter variants in malaria and schistosomiasis monoâ€ and coâ€infections among Nigerian children. Tropical Medicine and International Health, 2018, 23, 45-52.	2.3	9
121	Complement protein levels and MBL2 polymorphisms are associated with dengue and disease severity. Scientific Reports, 2020, 10, 14923.	3.3	9
122	Upregulation of Enzymes involved in ISGylation and Ubiquitination in patients with hepatocellular carcinoma. International Journal of Medical Sciences, 2020, 17, 347-353.	2.5	9
123	Diagnosis of Chikungunya Virus in Febrile Patients From a Malaria Holoendemic Area. International Journal of Infectious Diseases, 2021, 109, 247-252.	3.3	9
124	Geographical distribution of complement receptor type 1 variants and their associated disease risk. PLoS ONE, 2017, 12, e0175973.	2.5	9
125	Novel regulatory SNPs in the promoter region of the TNFRSF18 gene in a Gabonese population. Brazilian Journal of Medical and Biological Research, 2011, 44, 418-420.	1.5	8
126	Analysis of genetic variants in the IL4 promoter and VNTR loci in Indian patients with Visceral Leishmaniasis. Human Immunology, 2014, 75, 1177-1181.	2.4	8

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127	Soluble MICB protein levels and platelet counts during hepatitis B virus infection and response to hepatocellular carcinoma treatment. BMC Infectious Diseases, 2015, 15, 25.	2.9	8
128	Association of FCN2 polymorphisms and Ficolin-2 levels with dengue fever in Vietnamese patients. International Journal of Infectious Diseases, 2020, 95, 253-261.	3.3	8
129	Molecular surveillance and temporal monitoring of malaria parasites in focal Vietnamese provinces. Malaria Journal, 2020, 19, 458.	2.3	8
130	Molecular characterization of regulatory polymorphisms in the promoter region of the STAT6 gene in a Gabonese population. Memorias Do Instituto Oswaldo Cruz, 2011, 106, 65-69.	1.6	7
131	Characterization of Vibrio cholerae Strains Isolated from the Nigerian Cholera Outbreak in 2010. Journal of Clinical Microbiology, 2016, 54, 2618-2621.	3.9	7
132	Mannose-binding lectin deficiency and miscarriages in rheumatoid arthritis. Autoimmunity, 2017, 50, 409-413.	2.6	7
133	Cytochrome P450 CYP2B6*6 distribution among Congolese individuals with HIV, Tuberculosis and Malaria infection. International Journal of Infectious Diseases, 2019, 82, 111-116.	3.3	7
134	Predominance of HBV Genotype B and HDV Genotype 1 in Vietnamese Patients with Chronic Hepatitis. Viruses, 2021, 13, 346.	3.3	7
135	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
136	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. Journal of Antimicrobial Chemotherapy, 2020, 75, ii52-ii66.	3.0	7
137	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
138	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
139	High prevalence of dihydrofolate reductase gene mutations in Plasmodium falciparum parasites among pregnant women in Nigeria after reported use of sulfadoxine-pyrimethamine. Pathogens and Global Health, 2018, 112, 86-92.	2.3	6
140	KIR-HLA distribution in a Vietnamese population from Hanoi. Human Immunology, 2018, 79, 93-100.	2.4	6
141	Distribution of the cytochrome P450 CYP2C8*2 allele in Brazzaville, Republic of Congo. International Journal of Infectious Diseases, 2019, 85, 49-53.	3.3	6
142	Complement 5a Receptor Polymorphisms Are Associated With Panton-Valentine Leukocidin–positive <i>Staphylococcus aureus</i> Colonization in African Pygmies. Clinical Infectious Diseases, 2019, 68, 854-856.	5.8	6
143	Epstein-Barr virus, malaria and endemic Burkitt lymphoma. EBioMedicine, 2019, 39, 13-14.	6.1	6
144	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

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145	Upregulation of SMYD3 and SMYD3 VNTR 3/3 polymorphism increase the risk of hepatocellular carcinoma. Scientific Reports, 2020, 10, 2797.	3.3	6
146	Genetic Diversity of Enteric Viruses in Children under Five Years Old in Gabon. Viruses, 2021, 13, 545.	3.3	6
147	Pharmacogene Sequencing of a Gabonese Population with Severe Plasmodium falciparum Malaria Reveals Multiple Novel Variants with Putative Relevance for Antimalarial Treatment. Antimicrobial Agents and Chemotherapy, 2021, 65, e0027521.	3.2	6
148	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
149	Molecular surveillance and genetic divergence of rotavirus A antigenic epitopes in Gabonese children with acute gastroenteritis. EBioMedicine, 2021, 73, 103648.	6.1	6
150	Pharmacogenetic considerations in the treatment of co-infections with HIV/AIDS, tuberculosis and malaria in Congolese populations of Central Africa. International Journal of Infectious Diseases, 2021, 104, 207-213.	3.3	5
151	Viral and serological testing of SARS-CoV-2 among health care workers and patients in Vietnam. The Lancet Regional Health - Western Pacific, 2021, 8, 100113.	2.9	5
152	Genetic variants of programmed cell death 1 are associated with HBV infection and liver disease progression. Scientific Reports, 2021, 11, 7772.	3.3	5
153	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
154	Circulating level of sPD-1 and PD-1 genetic variants are associated with hepatitis B infection and related liver disease progression. International Journal of Infectious Diseases, 2022, 115, 229-236.	3.3	5
155	In-vitro characterization of novel and functional regulatory SNPs in the promoter region of IL2 and IL2R alpha in a Gabonese population. BMC Medical Genetics, 2012, 13, 117.	2.1	4
156	Hepatitis B Virus Infection Among Leprosy Patients: A Case for Polymorphisms Compromising Activation of the Lectin Pathway and Complement Receptors. Frontiers in Immunology, 2020, 11, 574457.	4.8	4
157	Maternal Vaginal Colonization and Extended-Spectrum Beta-Lactamase-Producing Bacteria in Vietnamese Pregnant Women. Antibiotics, 2021, 10, 572.	3.7	4
158	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
159	Genetic variants of interferon regulatory factor 5 associated with chronic hepatitis B infection. World Journal of Gastroenterology, 2018, 24, 248-256.	3.3	4
160	Evaluation of SARS-CoV-2 diagnostics and risk factors associated with SARS-CoV-2 infection in Zambia. International Journal of Infectious Diseases, 2022, 120, 150-157.	3.3	4
161	Rapid, low cost and sensitive detection of Calreticulin mutations by a PCR based amplicon length differentiation assay for diagnosis of myeloproliferative neoplasms. BMC Medical Genetics, 2019, 20, 115.	2.1	3
162	No expression of HBV-human chimeric fusion transcript (HBx-LINE1) among Vietnamese patients with HBV-associated hepatocellular carcinoma. Annals of Hepatology, 2019, 18, 404-405.	1.5	3

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163	Molecular surveillance of the Pfmdr1 N86Y allele among Congolese pregnant women with asymptomatic malaria. Malaria Journal, 2020, 19, 178.	2.3	3
164	Impact of <i>VSIG4</i> gene polymorphisms on susceptibility and functional status of rheumatoid arthritis. International Journal of Immunogenetics, 2021, 48, 260-265.	1.8	3
165	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
166	SARS-CoV-2 viral dynamics of the first 1000 sequences from Vietnam and neighbouring ASEAN countries. IJID Regions, 2022, 2, 175-179.	1.3	3
167	Analysis of polymorphic sites in the promoter of the nitric oxide synthase 2 gene in Brazilian patients with leprosy. International Journal of Immunogenetics, 2014, 41, 231-235.	1.8	2
168	Low <scp>MBL</scp> â€associated serine protease 2 (<scp>MASP</scp> â€2) levels correlate with urogenital schistosomiasis in Nigerian children. Tropical Medicine and International Health, 2015, 20, 1311-1319.	2.3	2
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