David Courtin

List of Publications by Year in descending order

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Version: 2024-02-01

		279778	3	330122	
59	1,624 citations	23		37	
papers	citations	h-index		g-index	
65	65	65		1954	
all docs	docs citations	times ranked		citing authors	

#	Article	IF	CITATIONS
1	Untreated Human Infections by Trypanosoma brucei gambiense Are Not 100% Fatal. PLoS Neglected Tropical Diseases, 2012, 6, e1691.	3.0	163
2	Do Cryptic Reservoirs Threaten Gambiense-Sleeping Sickness Elimination?. Trends in Parasitology, 2018, 34, 197-207.	3.3	139
3	Evidence that seasonal malaria chemoprevention with SPAQ influences blood and pre-erythrocytic stage antibody responses of Plasmodium falciparum infections in Niger. Malaria Journal, 2021, 20, 1.	2.3	106
4	The Quantity and Quality of African Children's IgG Responses to Merozoite Surface Antigens Reflect Protection against Plasmodium falciparum Malaria. PLoS ONE, 2009, 4, e7590.	2.5	91
5	Haptoglobin (HP) and Haptoglobin-related protein (HPR) copy number variation, natural selection, and trypanosomiasis. Human Genetics, 2014, 133, 69-83.	3.8	72
6	Worldwide genetic variation at the $3\hat{a}\in^2$ untranslated region of the HLA-G gene: balancing selection influencing genetic diversity. Genes and Immunity, 2014, 15, 95-106.	4.1	69
7	Aparasitemic serological suspects in Trypanosoma brucei gambiense human African trypanosomiasis: A potential human reservoir of parasites?. Acta Tropica, 2006, 98, 183-188.	2.0	59
8	Host genetics in African trypanosomiasis. Infection, Genetics and Evolution, 2008, 8, 229-238.	2.3	56
9	HLA-G $3\hat{a}\in^2$ UTR-2 haplotype is associated with Human African trypanosomiasis susceptibility. Infection, Genetics and Evolution, 2013, 17, 1-7.	2.3	42
10	Interest of tumor necrosis factor-alpha â^'308 G/A and interleukin-10 â^'592 C/A polymorphisms in human African trypanosomiasis. Infection, Genetics and Evolution, 2006, 6, 123-129.	2.3	41
11	<i>Schistosoma haematobium</i> infection affects <i>Plasmodium falciparum</i> â€specific IgG responses associated with protection against malaria. Parasite Immunology, 2011, 33, 124-131.	1.5	39
12	Human IgG Antibody Response to Glossina Saliva: An Epidemiologic Marker of Exposure to Glossina Bites. American Journal of Tropical Medicine and Hygiene, 2008, 78, 750-753.	1.4	39
13	Human African trypanosomiasis: connecting parasite and host genetics. Trends in Parasitology, 2006, 22, 405-409.	3.3	38
14	Association between human African trypanosomiasis and the IL6 gene in a Congolese population. Infection, Genetics and Evolution, 2007, 7, 60-68.	2.3	36
15	Genome Wide Linkage Study, Using a 250K SNP Map, of Plasmodium falciparum Infection and Mild Malaria Attack in a Senegalese Population. PLoS ONE, 2010, 5, e11616.	2.5	36
16	Association of HLA-G 3′UTR polymorphisms with response to malaria infection: A first insight. Infection, Genetics and Evolution, 2013, 16, 263-269.	2.3	35
17	Cattle as natural host for Schistosoma haematobium (Bilharz, 1852) Weinland, 1858 x Schistosoma bovis Sonsino, 1876 interactions, with new cercarial emergence and genetic patterns. Parasitology Research, 2020, 119, 2189-2205.	1.6	33
18	Associations between an IgG3 polymorphism in the binding domain for FcRn, transplacental transfer of malaria-specific IgG3, and protection against Plasmodium falciparum malaria during infancy: A birth cohort study in Benin. PLoS Medicine, 2017, 14, e1002403.	8.4	32

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19	High plasma levels of HLA-G are associated with low birth weight and with an increased risk of malaria in infancy. Malaria Journal, 2014, 13, 312.	2.3	31
20	Association of <i>HLAâ€G</i> 3′ untranslated region polymorphisms with antibody response against <i>Plasmodium falciparum</i> antigens: preliminary results. Tissue Antigens, 2013, 82, 53-58.	1.0	28
21	Human IgG antibody response to Glossina saliva: an epidemiologic marker of exposure to Glossina bites. American Journal of Tropical Medicine and Hygiene, 2008, 78, 750-3.	1.4	28
22	Balancing immunity and tolerance: genetic footprint of natural selection in the transcriptional regulatory region of HLA-G. Genes and Immunity, 2015, 16, 57-70.	4.1	24
23	HUMAN/VECTOR RELATIONSHIPS DURING HUMAN AFRICAN TRYPANOSOMIASIS: INITIAL SCREENING OF IMMUNOGENIC SALIVARY PROTEINS OF GLOSSINA SPECIES. American Journal of Tropical Medicine and Hygiene, 2007, 76, 327-333.	1.4	23
24	Association of IL-4 and IL-10 maternal haplotypes with immune responses to P. falciparum in mothers and newborns. BMC Infectious Diseases, 2013, 13, 215.	2.9	20
25	Specific antibodies to Anopheles gSG6-P1 salivary peptide to assess early childhood exposure to malaria vector bites. Malaria Journal, 2015, 14, 285.	2.3	20
26	Acquisition of natural humoral immunity to P. falciparum in early life in Benin: impact of clinical, environmental and host factors. Scientific Reports, 2016, 6, 33961.	3.3	20
27	The role of HLAâ€G in parasitic diseases. Hla, 2018, 91, 255-270.	0.6	20
28	Soluble human leukocyte antigen -G during pregnancy and infancy in Benin: Mother/child resemblance and association with the risk of malaria infection and low birth weight. PLoS ONE, 2017, 12, e0171117.	2.5	19
29	Comparison of cytokine plasma levels in human African trypanosomiasis. Tropical Medicine and International Health, 2006, 11, 647-653.	2.3	18
30	Human Leukocyte Antigen-G: A Promising Prognostic Marker of Disease Progression to Improve the Control of Human African Trypanosomiasis. Clinical Infectious Diseases, 2016, 63, ciw505.	5.8	15
31	NLRP7 and the genetics of post-molar choriocarcinomas in Senegal. Molecular Human Reproduction, 2012, 18, 52-56.	2.8	14
32	HLA-G, -E and -F regulatory and coding region variability and haplotypes in the Beninese Toffin population sample. Molecular Immunology, 2018, 104, 108-127.	2.2	14
33	First genome-wide association study of non-severe malaria in two birth cohorts in Benin. Human Genetics, 2019, 138, 1341-1357.	3.8	14
34	Plasmodium falciparum merozoite surface antigen-specific cytophilic IgG and control of malaria infection in a Beninese birth cohort. Malaria Journal, 2019, 18, 194.	2.3	14
35	Intravenous Artesunate for the Treatment of Severe Imported Malaria: Implementation, Efficacy, and Safety in 1391 Patients. Clinical Infectious Diseases, 2021, 73, 1795-1804.	5.8	13
36	Soil-transmitted helminth infection in pregnancy and long-term child neurocognitive and behavioral development: A prospective mother-child cohort in Benin. PLoS Neglected Tropical Diseases, 2021, 15, e0009260.	3.0	13

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37	<scp><i>HLA </i></scp> genetic diversity and evolutionary insights in two samples from Brazil and Benin. Hla, 2020, 96, 468-486.	0.6	12
38	<i>Schistosoma haematobium</i> infection modulates <i>Plasmodium falciparum</i> parasite density and antimalarial antibody responses. Parasite Immunology, 2020, 42, e12702.	1.5	12
39	Is Placental Malaria a Long-term Risk Factor for Mild Malaria Attack in Infancy? Revisiting a Paradigm. Clinical Infectious Diseases, 2018, 66, 930-935.	5.8	11
40	Evolution of the levels of human leukocyte antigen G (HLA-G) in Beninese infant during the first year of life in a malaria endemic area: using latent class analysis. Malaria Journal, 2016, 15, 78.	2.3	10
41	Genome-wide association study of antibody responses to Plasmodium falciparum candidate vaccine antigens. Genes and Immunity, 2016, 17, 110-117.	4.1	10
42	High level of soluble human leukocyte antigen (HLA)-G at beginning of pregnancy as predictor of risk of malaria during infancy. Scientific Reports, 2019, 9, 9160.	3.3	10
43	Susceptibility to Plasmodium falciparum Malaria: Influence of Combined Polymorphisms of IgG3 Gm Allotypes and Fc Gamma Receptors IIA, IIIA, and IIIB. Frontiers in Immunology, 2020, 11, 608016.	4.8	10
44	Human/vector relationships during human African trypanosomiasis: initial screening of immunogenic salivary proteins of Glossina species. American Journal of Tropical Medicine and Hygiene, 2007, 76, 327-33.	1.4	10
45	Trypanosome-induced Interferon- \hat{i}^3 production in whole blood stimulation assays is associated with latent Trypanosoma brucei gambiense infections. Microbes and Infection, 2016, 18, 436-440.	1.9	9
46	G6PD Aâ $^{\circ}$ variant influences the antibody responses to Plasmodium falciparum MSP2. Infection, Genetics and Evolution, 2011, 11, 1287-1292.	2.3	8
47	Comparison of growth models to describe growth from birth to 6 years in a Beninese cohort of children with repeated measurements. BMJ Open, 2020, 10, e035785.	1.9	6
48	HLA-G expression during hookworm infection in pregnant women. Acta Tropica, 2019, 196, 52-59.	2.0	5
49	Increased Risk of Malaria During the First Year of Life in Small-for-Gestational-Age Infants: A Longitudinal Study in Benin. Journal of Infectious Diseases, 2019, 219, 1642-1651.	4.0	5
50	Plasmodium falciparum infection and age influence parasite growth inhibition mediated by IgG in Beninese infants. Acta Tropica, 2016, 159, 111-119.	2.0	4
51	Placental Malaria is Associated with Higher LILRB2 Expression in Monocyte Subsets and Lower Anti-Malarial IgG Antibodies During Infancy. Frontiers in Immunology, 0, 13, .	4.8	4
52	Blood lead level in infants and subsequent risk of malaria: A prospective cohort study in Benin, Sub-Saharan Africa. PLoS ONE, 2019, 14, e0220023.	2.5	3
53	Mixed logistic regression in genome-wide association studies. BMC Bioinformatics, 2020, 21, 536.	2.6	3
54	Red Blood Cell Deformability, Age, Ethnicity and Susceptibility to Malaria in Africa. Blood, 2016, 128, 2441-2441.	1.4	3

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55	The Impact of Maternal Depression and Parent–Child Interactions on Risk of Parasitic Infections in Early Childhood: A Prospective Cohort in Benin. Maternal and Child Health Journal, 2022, 26, 1049-1058.	1.5	2
56	Genotyping complex structural variation at the malariaâ€associated human glycophorin locus using a PCRâ€based strategy. Annals of Human Genetics, 2021, 85, 7-17.	0.8	1
57	Editorial: The Role of Gene Polymorphisms in Modulating the Immune Responses Against Tropical Infectious Diseases. Frontiers in Immunology, 2021, 12, 714237.	4.8	1
58	<i>> Plasmodium falciparum $$ </i> $$ coinfection is associated with improved IgE and IgG3 response against hookworm antigens. Health Science Reports, 2022, 5, .	1.5	1
59	Human leukocyte antigen (HLA)-F and -G gene polymorphisms and haplotypes are associated with malaria susceptibility in the Beninese Toffin children. Infection, Genetics and Evolution, 2021, 92, 104828.	2.3	0