Mette Christiansen

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/75789/publications.pdf

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		471509	414414
43	1,158	17	32
papers	citations	h-index	g-index

43	43	43	2002
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Functional IRF3 deficiency in a patient with herpes simplex encephalitis. Journal of Experimental Medicine, 2015, 212, 1371-1379.	8.5	171
2	Report of the first nationally implemented clinical routine screening for fetal RHD in Dâ˜' pregnant women to ascertain the requirement for antenatal RhD prophylaxis. Transfusion, 2012, 52, 752-758.	1.6	140
3	Characterization of the clinical and immunologic phenotype and management of 157 individuals with 56 distinct heterozygous NFKB1 mutations. Journal of Allergy and Clinical Immunology, 2020, 146, 901-911.	2.9	78
4	Mutations in the TLR3 signaling pathway and beyond in adult patients with herpes simplex encephalitis. Genes and Immunity, 2015, 16, 552-566.	4.1	75
5	Routine noninvasive prenatal screening for fetal RHD in plasma of RhDâ€negative pregnant womenâ€"2â€‰years of screening experience from Denmark. Prenatal Diagnosis, 2014, 34, 1000-1005.	2.3	71
6	Novel STAT1 Alleles in a Patient with Impaired Resistance to Mycobacteria. Journal of Clinical Immunology, 2011, 31, 265-271.	3.8	58
7	Mutations in RNA Polymerase III genes and defective DNA sensing in adults with varicella-zoster virus CNS infection. Genes and Immunity, 2019, 20, 214-223.	4.1	54
8	Varicella-zoster virus CNS vasculitis and RNA polymerase III gene mutation in identical twins. Neurology: Neuroimmunology and NeuroInflammation, 2018, 5, e500.	6.0	49
9	Defective RNA sensing by RIG-I in severe influenza virus infection. Clinical and Experimental Immunology, 2018, 192, 366-376.	2.6	39
10	Genetic, Immunological, and Clinical Features of 32 Patients with Autosomal Recessive STAT1 Deficiency. Journal of Immunology, 2021, 207, 133-152.	0.8	33
11	Neonatal-onset T â˜' B â˜' NK + severe combined immunodeficiency and neutropenia caused by mutated phosphoglucomutase 3. Journal of Allergy and Clinical Immunology, 2016, 137, 321-324.	2.9	29
12	Identification of Novel Genetic Variants in CVID Patients With Autoimmunity, Autoinflammation, or Malignancy. Frontiers in Immunology, 2019, 10, 3022.	4.8	28
13	Defects in LC3B2 and ATG4A underlie HSV2 meningitis and reveal a critical role for autophagy in antiviral defense in humans. Science Immunology, 2020, 5, .	11.9	27
14	A STAT1-gain-of-function mutation causing Th17 deficiency with chronic mucocutaneous candidiasis, psoriasiform hyperkeratosis and dermatophytosis. BMJ Case Reports, 2015, 2015, bcr2015211372.	0.5	25
15	Correlation between serology and genetics of weak D types in Denmark. Transfusion, 2008, 48, 187-193.	1.6	24
16	RHD positive among C/E+ and Dâ˜' blood donors in Denmark. Transfusion, 2010, 50, 1460-1464.	1.6	24
17	The genetic component of preeclampsia: A whole-exome sequencing study. PLoS ONE, 2018, 13, e0197217.	2.5	21
18	The challenge of discriminating between <scp>HIV</scp>â€1, <scp>HIV</scp>â€2 and <scp>HIV</scp>â€1/2 dual infections. HIV Medicine, 2018, 19, 403-410.	2.2	20

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19	Defective interferon priming and impaired antiviral responses in a patient with an IRF7 variant and severe influenza. Medical Microbiology and Immunology, 2019, 208, 869-876.	4.8	19
20	Noninvasive fetal <scp>RHD</scp> genotyping to guide targeted antiâ€Ð prophylaxisâ€“an external quality assessment workshop. Vox Sanguinis, 2019, 114, 386-393.	1.5	14
21	Identification of an IRF3 variant and defective antiviral interferon responses in a patient with severe influenza. European Journal of Immunology, 2019, 49, 2111-2114.	2.9	13
22	Whole Exome Sequencing of HIV-1 long-term non-progressors identifies rare variants in genes encoding innate immune sensors and signaling molecules. Scientific Reports, 2018, 8, 15253.	3.3	12
23	Frequently used bioinformatics tools overestimate the damaging effect of allelic variants. Genes and Immunity, 2019, 20, 10-22.	4.1	12
24	Systemic juvenile idiopathic arthritis and recurrent macrophage activation syndrome due to a CASP1 variant causing inflammasome hyperactivation. Rheumatology, 2020, 59, 3099-3105.	1.9	12
25	Ectodermal dysplasia with immunodeficiency caused by a branch-point mutation in IKBKG/NEMO. Journal of Allergy and Clinical Immunology, 2016, 138, 1706-1709.e4.	2.9	11
26	Assessing a single targeted next generation sequencing for human leukocyte antigen typing protocol for interoperability, as performed by users with variable experience. Human Immunology, 2017, 78, 642-648.	2.4	11
27	The influence of human leukocyte antigen-types on disease progression among HIV-2 infected patients in Guinea-Bissau. Aids, 2018, 32, 721-728.	2.2	10
28	XIAP deficiency and MEFV variants resulting in an autoinflammatory lymphoproliferative syndrome. BMJ Case Reports, 2016, 2016, bcr2016216922.	0.5	9
29	Phylogeny of human T-lymphotropic virus-1 subtypes in Guinea-Bissau. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2018, 112, 175-180.	1.8	9
30	Whole-Exome Sequencing of Patients With Recurrent HSV-2 Lymphocytic Mollaret Meningitis. Journal of Infectious Diseases, 2021, 223, 1776-1786.	4.0	9
31	Vox Sanguinis International Forum on application of fetal blood grouping. Vox Sanguinis, 2018, 113, e26-e35.	1.5	8
32	Impaired immune responses to herpesviruses and microbial ligands in patients with Mono MAC. British Journal of Haematology, 2019, 186, 471-476.	2.5	8
33	Pattern Recognition Molecules of the Lectin Pathwayâ€”Screening of Patients with Suspected Immunodeficiency. Journal of Clinical Immunology, 2019, 39, 668-677.	3.8	7
34	Host Genetics, Innate Immune Responses, and Cellular Death Pathways in Poliomyelitis Patients. Frontiers in Microbiology, 2019, 10, 1495.	3.5	7
35	Discovery of a novel <scp>HLA</scp>â€Ƀ*15 allele, <scp>HLA</scp>â€Ƀ*15:379, in a patient from Guineaâ€Ƀissau. Hla, 2016, 88, 203-204.	0.6	4
36	Protein a Immunoadsorption May Hamper the Decision to Transplant Due to Interference With CDC Crossmatch Results. Journal of Clinical Apheresis, 2017, 32, 163-169.	1.3	3

#	Article	IF	Citations
37	HLAâ€ÐQB1*06:276 , a novel HLA allele fund in a patient from Guineaâ€Bissau. Hla, 2019, 93, 243-244.	0.6	3
38	Detecting mismatched donor HLA types from allograft biopsies â€“ An easily applicable tool for improved individualized risk assessment. Human Immunology, 2020, 81, 337-341.	2.4	3
39	Vox Sanguinis International forum on the selection and preparation of blood components for intrauterine transfusion. Vox Sanguinis, 2020, 115, e18-e38.	1.5	3
40	Two novel HLAâ€B alleles, HLAâ€B*53:01:17 and â€B*58:83, found in patients from Guineaâ€Bissau. Hla, 2018, 92, 417-418.	0.6	2
41	HTLV infected individuals have increased B-cell activation and proinflammatory regulatory T-cells. Immunobiology, 2020, 225, 151878.	1.9	2
42	Discriminatory rapid tests cause HIV-type misclassificationâ€"evaluation of three rapid tests using clinical samples from Guinea-Bissau. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2019, 113, 555-559.	1.8	1
43	Very early onset inflammatory bowel disease with compound heterozygous variants in Nuclear Factor of Activated T cell 5. European Journal of Immunology, 2021, 51, 999-1001.	2.9	0