

Maude E Phipps

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

2,946
citations

331670

21
h-index

276875

41
g-index

48
all docs

48
docs citations

48
times ranked

3934
citing authors

#	ARTICLE	IF	CITATIONS
1	Allelic and haplotypic HLA diversity in indigenous Malaysian populations explored using Next Generation Sequencing. <i>Human Immunology</i> , 2022, 83, 17-26.	2.4	4
2	The gut virome in two indigenous populations from Malaysia. <i>Scientific Reports</i> , 2022, 12, 1824.	3.3	8
3	Insights into the demographic history of Asia from common ancestry and admixture in the genomic landscape of present-day Austroasiatic speakers. <i>BMC Biology</i> , 2021, 19, 61.	3.8	8
4	Shared Signature of Recent Positive Selection on the <i>TSBP1</i> and <i>BTNL2</i> HLA-DRA Genes in Five Native Populations from North Borneo. <i>Genome Biology and Evolution</i> , 2020, 12, 2245-2257.	2.5	2
5	Seroprevalence of Nipah Virus Infection in Peninsular Malaysia. <i>Journal of Infectious Diseases</i> , 2020, 221, S370-S374.	4.0	6
6	Analysis of five deep-sequenced trio-genomes of the Peninsular Malaysia Orang Asli and North Borneo populations. <i>BMC Genomics</i> , 2019, 20, 842.	2.8	3
7	Metabolic syndrome and cardiometabolic risk factors among indigenous Malaysians. <i>Public Health</i> , 2019, 176, 106-113.	2.9	10
8	Health and saliva microbiomes of a semi-urbanized indigenous tribe in Peninsular Malaysia. <i>F1000Research</i> , 2019, 8, 175.	1.6	9
9	Health and saliva microbiomes of a semi-urbanized indigenous tribe in Peninsular Malaysia. <i>F1000Research</i> , 2019, 8, 175.	1.6	8
10	Genetic relatedness of indigenous ethnic groups in northern Borneo to neighboring populations from Southeast Asia, as inferred from genome-wide SNP data. <i>Annals of Human Genetics</i> , 2018, 82, 216-226.	0.8	13
11	Genomic structure of the native inhabitants of Peninsular Malaysia and North Borneo suggests complex human population history in Southeast Asia. <i>Human Genetics</i> , 2018, 137, 161-173.	3.8	20
12	The prehistoric peopling of Southeast Asia. <i>Science</i> , 2018, 361, 88-92.	12.6	291
13	Discerning the Origins of the Negritos, First Sundaland People: Deep Divergence and Archaic Admixture. <i>Genome Biology and Evolution</i> , 2017, 9, 2013-2022.	2.5	54
14	A genomic history of Aboriginal Australia. <i>Nature</i> , 2016, 538, 207-214.	27.8	439
15	Unravelling the Genetic History of Negritos and Indigenous Populations of Southeast Asia. <i>Genome Biology and Evolution</i> , 2015, 7, 1206-1215.	2.5	63
16	Differential positive selection of malaria resistance genes in three indigenous populations of Peninsular Malaysia. <i>Human Genetics</i> , 2015, 134, 375-392.	3.8	19
17	Cardio-metabolic health risks in indigenous populations of Southeast Asia and the influence of urbanization. <i>BMC Public Health</i> , 2015, 15, 47.	2.9	36
18	Novel Population Specific Autosomal Copy Number Variation and Its Functional Analysis amongst Negritos from Peninsular Malaysia. <i>PLoS ONE</i> , 2014, 9, e100371.	2.5	6

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19	The population genomic landscape of human genetic structure, admixture history and local adaptation in Peninsular Malaysia. <i>Human Genetics</i> , 2014, 133, 1169-1185.	3.8	30
20	Admixture Patterns and Genetic Differentiation in Negrito Groups from West Malaysia Estimated from Genome-wide SNP Data. <i>Human Biology</i> , 2013, 85, 173-188.	0.2	21
21	HLA variants rs9271366 and rs9275328 are associated with systemic lupus erythematosus susceptibility in Malays and Chinese. <i>Lupus</i> , 2013, 22, 198-204.	1.6	9
22	Admixture Patterns and Genetic Differentiation in Negrito Groups from West Malaysia Estimated from Genome-wide SNP Data. <i>Human Biology</i> , 2013, 85, 173.	0.2	3
23	Evolutionary History of Continental Southeast Asians: "Early Train" Hypothesis Based on Genetic Analysis of Mitochondrial and Autosomal DNA Data. <i>Molecular Biology and Evolution</i> , 2012, 29, 3513-3527.	8.9	122
24	Denisova Admixture and the First Modern Human Dispersals into Southeast Asia and Oceania. <i>American Journal of Human Genetics</i> , 2011, 89, 516-528.	6.2	525
25	Contribution of VKORC1 and CYP2C9 polymorphisms in the interethnic variability of warfarin dose in Malaysian populations. <i>Annals of Hematology</i> , 2011, 90, 635-641.	1.8	41
26	Molecular analysis of HLA Class I and Class II genes in four indigenous Malaysian populations. <i>Tissue Antigens</i> , 2010, 75, 151-158.	1.0	19
27	MICA polymorphism: biology and importance in immunity and disease. <i>Trends in Molecular Medicine</i> , 2010, 16, 97-106.	6.7	89
28	Mapping Human Genetic Diversity in Asia. <i>Science</i> , 2009, 326, 1541-1545.	12.6	557
29	The association between HLA genes and radiological erosions in Malaysian patients with rheumatoid arthritis. <i>Autoimmunity</i> , 2007, 40, 187-190.	2.6	3
30	The distribution of major histocompatibility complex class I polymorphic Alu insertions and their associations with HLA alleles in a Chinese population from Malaysia. <i>Tissue Antigens</i> , 2007, 70, 136-143.	1.0	21
31	Possible Polyphyletic Origin of Major Histocompatibility Complex Class I Chain-Related Gene A (MICA) Alleles. <i>Journal of Molecular Evolution</i> , 2003, 57, 38-43.	1.8	6
32	HLA-DRB1 Genes and Susceptibility to Rheumatoid Arthritis in Three Ethnic Groups from Malaysia. <i>Autoimmunity</i> , 2002, 35, 235-239.	2.6	37
33	Human Fc gamma receptor IIA (Fc γ RIIA) genotyping and association with systemic lupus erythematosus (SLE) in Chinese and Malays in Malaysia. <i>Lupus</i> , 1999, 8, 305-310.	1.6	32
34	Fc gamma receptor IIIB-NA gene frequencies in patients with systemic lupus erythematosus and healthy individuals of Malay and Chinese ethnicity. <i>Immunology Letters</i> , 1999, 68, 295-300.	2.5	21
35	Molecular genetic analysis of the 3p " syndrome. <i>Human Molecular Genetics</i> , 1994, 3, 903-908.	2.9	42
36	Detailed mapping of germline deletions of the von Hippel-Lindau disease tumour suppressor gene. <i>Human Molecular Genetics</i> , 1994, 3, 595-598.	2.9	81

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37	Molecular genetic investigations of the mechanism of tumourigenesis in von Hippel-Lindau disease: analysis of allele loss in VHL tumours. Human Genetics, 1994, 93, 53-8.	3.8	129
38	Physical mapping of chromosome 3p25-p26 by fluorescence in situ hybridisation (FISH). Human Genetics, 1993, 92, 18-22.	3.8	10
39	Detailed genetic mapping of the von Hippel-Lindau disease tumour suppressor gene.. Journal of Medical Genetics, 1993, 30, 104-107.	3.2	51
40	Genetic linkage between Von Hippel-Lindau disease and three microsatellite polymorphisms refines the localisation of the VHL locus. Human Molecular Genetics, 1993, 2, 279-282.	2.9	45
41	Mapping the Von Hippel-Lindau disease tumour suppressor gene: identification of germline deletions by pulsed field gel electrophoresis. Human Molecular Genetics, 1993, 2, 879-882.	2.9	53