

Jun Ohashi

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

1,000
citations

623734

14
h-index

552781

26
g-index

28
all docs

28
docs citations

28
times ranked

1911
citing authors

#	ARTICLE	IF	CITATIONS
1	A Practical Genome Scan for Population-Specific Strong Selective Sweeps That Have Reached Fixation. PLoS ONE, 2007, 2, e286.	2.5	378
2	FTO polymorphisms in oceanic populations. Journal of Human Genetics, 2007, 52, 1031-1035.	2.3	127
3	Extended Linkage Disequilibrium Surrounding the Hemoglobin E Variant Due to Malarial Selection. American Journal of Human Genetics, 2004, 74, 1198-1208.	6.2	117
4	Significant Association of Longer Forms of CCTTT Microsatellite Repeat in the Inducible Nitric Oxide Synthase Promoter with Severe Malaria in Thailand. Journal of Infectious Diseases, 2002, 186, 578-581.	4.0	52
5	Gene Flow and Natural Selection in Oceanic Human Populations Inferred from Genome-Wide SNP Typing. Molecular Biology and Evolution, 2008, 25, 1750-1761.	8.9	47
6	The Impact of Natural Selection on an ABCC11 SNP Determining Earwax Type. Molecular Biology and Evolution, 2011, 28, 849-857.	8.9	44
7	Appropriate data cleaning methods for genome-wide association study. Journal of Human Genetics, 2008, 53, 886-893.	2.3	40
8	Association and interaction analyses of NRG1 and ERBB4 genes with schizophrenia in a Japanese population. Journal of Human Genetics, 2008, 53, 929-935.	2.3	33
9	Power of genome-wide linkage disequilibrium testing by using microsatellite markers. Journal of Human Genetics, 2003, 48, 487-491.	2.3	31
10	Brief communication: Mitochondrial DNA variation suggests extensive gene flow from Polynesian ancestors to indigenous Melanesians in the northwestern Bismarck Archipelago. American Journal of Physical Anthropology, 2006, 130, 551-556.	2.1	19
11	The Origin and Composition of Korean Ethnicity Analyzed by Ancient and Present-Day Genome Sequences. Genome Biology and Evolution, 2020, 12, 553-565.	2.5	17
12	Polymorphisms in the ABO blood group gene in three populations in the New Georgia group of the Solomon Islands. Journal of Human Genetics, 2006, 51, 407-411.	2.3	16
13	MHC (Major Histocompatibility Complex)-DRB Genes and Polymorphisms in Common Marmoset. Journal of Molecular Evolution, 2000, 51, 214-222.	1.8	15
14	A functional polymorphism in the IL1B gene promoter, IL1B -31C>T, is not associated with cerebral malaria in Thailand. Malaria Journal, 2005, 4, 38.	2.3	14
15	Comparative study of the haplotype structure and linkage disequilibrium of chromosome 1p36.2 region in the Korean and Japanese populations. Journal of Human Genetics, 2004, 49, 603-609.	2.3	7
16	Admixture and natural selection shaped genomes of an Austronesian-speaking population in the Solomon Islands. Scientific Reports, 2020, 10, 6872.	3.3	7
17	HLA-DRB1 polymorphism on Ha'ano island of the Kingdom of Tonga. Anthropological Science, 2006, 114, 193-198.	0.4	7
18	Application of the Stepwise Focusing Method to Optimize the Cost-effectiveness of Genome-wide Association Studies with Limited Research Budgets for Genotyping and Phenotyping. Annals of Human Genetics, 2005, 69, 323-328.	0.8	6

#	ARTICLE	IF	CITATIONS
19	Linkage disequilibrium structure of the 5q31-33 region in a Thai population. <i>Journal of Human Genetics</i> , 2008, 53, 850-856.	2.3	6
20	Computer Simulation Analysis Suggests Weak Balancing Selection Operative at the MICALocus. <i>Hereditas</i> , 2000, 133, 25-28.	1.4	4
21	Strong linkage disequilibrium of a HbE variant with the (AT) ₉ (T) ₅ repeat in the BP1 binding site upstream of the β^2 -globin gene in the Thai population. <i>Journal of Human Genetics</i> , 2005, 50, 7-11.	2.3	4
22	Power of association test for detecting minor histocompatibility gene causing graft-versus-host disease following bone marrow transplantation. <i>Journal of Human Genetics</i> , 2003, 48, 502-507.	2.3	2
23	Admixture with indigenous people helps local adaptation: admixture-enabled selection in Polynesians. <i>Bmc Ecology and Evolution</i> , 2021, 21, 179.	1.6	2
24	HLA-DRB1 Polymorphism of Balopa Islanders in Papua New Guinea. <i>Anthropological Science</i> , 2003, 111, 157-164.	0.4	2
25	Lack of association between interleukin-10 gene promoter polymorphism, -1082G/A, and severe malaria in Thailand. <i>Southeast Asian Journal of Tropical Medicine and Public Health</i> , 2002, 33 Suppl 3, 5-7.	1.0	2
26	The genotypes of GYPA and GYPB carrying the MNSs antigens are not associated with cerebral malaria. <i>Journal of Human Genetics</i> , 2007, 52, 476-479.	2.3	1
27	Gene Expression Profiles of CD133-positive Fractions Predict the Survival of Individuals with Acute Myeloid Leukemia. <i>Cancer Genomics and Proteomics</i> , 2006, 3, 169-181.	2.0	0