## Jun Ohashi

## List of Publications by Year in descending order

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623734 552781 1,000 27 14 26 citations g-index h-index papers 28 28 28 1911 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Admixture with indigenous people helps local adaptation: admixture-enabled selection in Polynesians. Bmc Ecology and Evolution, 2021, 21, 179.	1.6	2
2	Admixture and natural selection shaped genomes of an Austronesian-speaking population in the Solomon Islands. Scientific Reports, 2020, 10, 6872.	3.3	7
3	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
4	The Impact of Natural Selection on an ABCC11 SNP Determining Earwax Type. Molecular Biology and Evolution, 2011, 28, 849-857.	8.9	44
5	Linkage disequilibrium structure of the 5q31-33 region in a Thai population. Journal of Human Genetics, 2008, 53, 850-856.	2.3	6
6	Appropriate data cleaning methods for genome-wide association study. Journal of Human Genetics, 2008, 53, 886-893.	2.3	40
7	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
8	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
9	A Practical Genome Scan for Population-Specific Strong Selective Sweeps That Have Reached Fixation. PLoS ONE, 2007, 2, e286.	2.5	378
10	The genotypes of GYPA and GYPB carrying the MNSs antigens are not associated with cerebral malaria. Journal of Human Genetics, 2007, 52, 476-479.	2.3	1
11	FTO polymorphisms in oceanic populations. Journal of Human Genetics, 2007, 52, 1031-1035.	2.3	127
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	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
14	HLA-DRB1 polymorphism on Ha'ano island of the Kingdom of Tonga. Anthropological Science, 2006, 114, 193-198.	0.4	7
15	Gene Expression Profiles of CD133-positive Fractions Predict the Survival of Individuals with Acute Myeloid Leukemia. Cancer Genomics and Proteomics, 2006, 3, 169-181.	2.0	O
16	Strong linkage disequilibrium of a HbE variant with the (AT)9(T)5 repeat in the BP1 binding site upstream of the $\hat{l}^2$ -globin gene in the Thai population. Journal of Human Genetics, 2005, 50, 7-11.	2.3	4
17	Application of the Stepwise Focusing Method to Optimize the Costâ€effectiveness of Genomeâ€wide Association Studies with Limited Research Budgets for Genotyping and Phenotyping. Annals of Human Genetics, 2005, 69, 323-328.	0.8	6
18	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

#	Article	IF	Citations
19	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
20	Extended Linkage Disequilibrium Surrounding the Hemoglobin E Variant Due to Malarial Selection. American Journal of Human Genetics, 2004, 74, 1198-1208.	6.2	117
21	Power of genome-wide linkage disequilibrium testing by using microsatellite markers. Journal of Human Genetics, 2003, 48, 487-491.	2.3	31
22	Power of association test for detecting minor histocompatibility gene causing graft-versus-host disease following bone barrow transplantation. Journal of Human Genetics, 2003, 48, 502-507.	2.3	2
23	HLA-DRB1 Polymorphism of Balopa Islanders in Papua New Guinea. Anthropological Science, 2003, 111, 157-164.	0.4	2
24	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
25	Lack of association between interleukin-10 gene promoter polymorphism, -1082G/A, and severe malaria in Thailand. Southeast Asian Journal of Tropical Medicine and Public Health, 2002, 33 Suppl 3, 5-7.	1.0	2
26	Computer Simulation Analysis Suggests Weak Balancing Selection Operative at the MICALocus. Hereditas, 2000, 133, 25-28.	1.4	4
27	MHC (Major Histocompatibility Complex)-DRB Genes and Polymorphisms in Common Marmoset. Journal of Molecular Evolution, 2000, 51, 214-222.	1.8	15