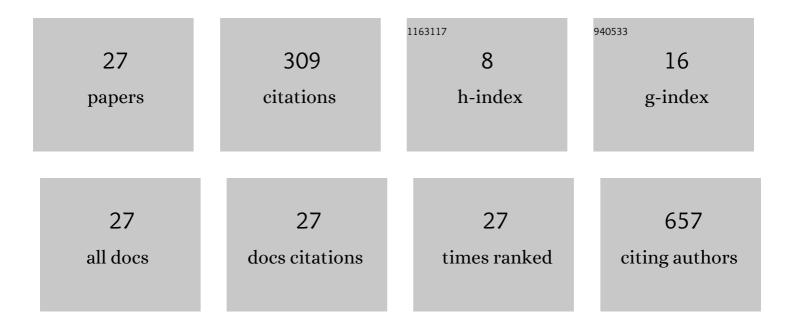
Yoshiki Yasukochi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Upregulation of cathepsin L gene under mild cold conditions in young Japanese male adults. Journal of Physiological Anthropology, 2021, 40, 16.	2.6	3
2	Individual variations and sex differences in hemodynamics with percutaneous arterial oxygen saturation (SpO2) in young Andean highlanders in Bolivia. Journal of Physiological Anthropology, 2020, 39, 31.	2.6	6
3	Transcriptomic Changes in Young Japanese Males After Exposure to Acute Hypobaric Hypoxia. Frontiers in Genetics, 2020, 11, 559074.	2.3	8
4	Effect of EGLN1 Genetic Polymorphisms on Hemoglobin Concentration in Andean Highlanders. BioMed Research International, 2020, 2020, 1-16.	1.9	3
5	Effects of acute hypobaric hypoxia on thermoregulatory and circulatory responses during cold air exposure. Journal of Physiological Anthropology, 2020, 39, 28.	2.6	4
6	Development of a novel monoclonal antibody that binds to most HLA-A allomorphs in a conformation-dependent yet peptide-promiscuous fashion. Immunogenetics, 2020, 72, 143-153.	2.4	0
7	Identification of six novel susceptibility loci for dyslipidemia using longitudinal exome-wide association studies in a Japanese population. Genomics, 2019, 111, 520-533.	2.9	2
8	Evolutionary history of diseaseâ€susceptibility loci identified in longitudinal exomeâ€wide association studies. Molecular Genetics & Genomic Medicine, 2019, 7, e925.	1.2	1
9	Two novel susceptibility loci for type 2 diabetes mellitus identified by longitudinal exome-wide association studies in a Japanese population. Genomics, 2019, 111, 34-42.	2.9	5
10	Identification of CDC42BPG as a novel susceptibility locus for hyperuricemia in a Japanese population. Molecular Genetics and Genomics, 2018, 293, 371-379.	2.1	12
11	Six novel susceptibility loci for coronary artery disease and cerebral infarction identified by longitudinal exomeâ€wide association studies in a Japanese population. Biomedical Reports, 2018, 9, 123-134.	2.0	8
12	Identification of three genetic variants as novel susceptibility loci for body mass index in a Japanese population. Physiological Genomics, 2018, 50, 179-189.	2.3	8
13	Association of EGLN1 genetic polymorphisms with SpO2 responses to acute hypobaric hypoxia in a Japanese cohort. Journal of Physiological Anthropology, 2018, 37, 9.	2.6	15
14	Identification of nine novel loci related to hematological traits in a Japanese population. Physiological Genomics, 2018, 50, 758-769.	2.3	5
15	Evolution of Fseg/Cseg dimorphism in region III of the Plasmodium falciparum eba-175 gene. Infection, Genetics and Evolution, 2017, 49, 251-255.	2.3	1
16	Longitudinal exome-wide association study to identify genetic susceptibility loci for hypertension in a Japanese population. Experimental and Molecular Medicine, 2017, 49, e409-e409.	7.7	8
17	Elucidating the origin of HLA-B*73 allelic lineage: Did modern humans benefit by archaic introgression?. Immunogenetics, 2017, 69, 63-67.	2.4	7
18	Genetic evidence for contribution of human dispersal to the genetic diversity of EBA-175 in Plasmodium falciparum. Malaria Journal, 2015, 14, 293.	2.3	4

Үознікі Үазикосні

#	Article	IF	CITATIONS
19	Molecular Evolution of the CYP2D Subfamily in Primates: Purifying Selection on Substrate Recognition Sites without the Frequent or Long-Tract Gene Conversion. Genome Biology and Evolution, 2015, 7, 1053-1067.	2.5	14
20	Nonsynonymous Substitution Rate Heterogeneity in the Peptide-Binding Region Among Different <i>HLA-DRB1</i> Lineages in Humans. G3: Genes, Genomes, Genetics, 2014, 4, 1217-1226.	1.8	6
21	A human-specific allelic group of the MHC DRB1 gene in primates. Journal of Physiological Anthropology, 2014, 33, 14.	2.6	9
22	Current perspectives on the intensity of natural selection of MHC loci. Immunogenetics, 2013, 65, 479-483.	2.4	30
23	MHC class II DQB diversity in the Japanese black bear, Ursus thibetanus japonicus. BMC Evolutionary Biology, 2012, 12, 230.	3.2	17
24	Evolution of the CYP2D gene cluster in humans and four non-human primates. Genes and Genetic Systems, 2011, 86, 109-116.	0.7	32
25	Identification of the expressed MHC class II DQB gene of the Asiatic black bear, Ursus thibetanus, in Japan. Genes and Genetic Systems, 2010, 85, 147-155.	0.7	7
26	Genetic Structure of the Asiatic Black Bear in Japan Using Mitochondrial DNA Analysis. Journal of Heredity, 2009, 100, 297-308.	2.4	70
27	Tandem duplication of mitochondrial DNA in the black-faced spoonbill, Platalea minor. Genes and Genetic Systems, 2009, 84, 297-305.	0.7	24