Yoshiki Yasukochi

List of Publications by Year
in descending order

27	309	1163117 8	940533 16
papers	citations	h-index	g-index

27	27	27	657
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genetic Structure of the Asiatic Black Bear in Japan Using Mitochondrial DNA Analysis. Journal of Heredity, 2009, 100, 297-308.	2.4	70
2	Evolution of the CYP2D gene cluster in humans and four non-human primates. Genes and Genetic Systems, 2011, 86, 109-116.	0.7	32
3	Current perspectives on the intensity of natural selection of MHC loci. Immunogenetics, 2013, 65, 479-483.	2.4	30
4	Tandem duplication of mitochondrial DNA in the black-faced spoonbill, Platalea minor. Genes and Genetic Systems, 2009, 84, 297-305.	0.7	24
5	MHC class II DQB diversity in the Japanese black bear, Ursus thibetanus japonicus. BMC Evolutionary Biology, 2012, 12, 230.	3.2	17
6	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
7	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
8	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
9	A human-specific allelic group of the MHC DRB1 gene in primates. Journal of Physiological Anthropology, 2014, 33, 14.	2.6	9
10	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
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	Transcriptomic Changes in Young Japanese Males After Exposure to Acute Hypobaric Hypoxia. Frontiers in Genetics, 2020, 11, 559074.	2.3	8
14	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
15	Elucidating the origin of HLA-B*73 allelic lineage: Did modern humans benefit by archaic introgression?. Immunogenetics, 2017, 69, 63-67.	2.4	7
16	Nonsynonymous Substitution Rate Heterogeneity in the Peptide-Binding Region Among DifferentHLA-DRB1Lineages in Humans. G3: Genes, Genomes, Genetics, 2014, 4, 1217-1226.	1.8	6
17	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
18	Identification of nine novel loci related to hematological traits in a Japanese population. Physiological Genomics, 2018, 50, 758-769.	2.3	5

#	Article	IF	Citations
19	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
20	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
21	Effects of acute hypobaric hypoxia on thermoregulatory and circulatory responses during cold air exposure. Journal of Physiological Anthropology, 2020, 39, 28.	2.6	4
22	Effect of EGLN1 Genetic Polymorphisms on Hemoglobin Concentration in Andean Highlanders. BioMed Research International, 2020, 2020, 1-16.	1.9	3
23	Upregulation of cathepsin L gene under mild cold conditions in young Japanese male adults. Journal of Physiological Anthropology, 2021, 40, 16.	2.6	3
24	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
25	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
26	Evolutionary history of diseaseâ€susceptibility loci identified in longitudinal exomeâ€wide association studies. Molecular Genetics & Genomic Medicine, 2019, 7, e925.	1.2	1
27	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