# 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 | 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 |

| # | 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 DifferentHLA-DRB1Lineages 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 |