Masaaki Muramatsu

List of Publications by Year
in descending order

48	805	623734 14	526287 27
papers	citations	h-index	g-index

53	53	53	1551
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Dâ€˜amino acid oxidase (DAO) rare genetic missense variant p.Pro103Leu and gastric cancer. Molecular and Clinical Oncology, 2021, 14, 58.	1.0	2
2	Trimester-specific associations between extracellular vesicle microRNAs and fetal growth. Journal of Maternal-Fetal and Neonatal Medicine, 2021, , 1-7.	1.5	2
3	COL17A1 germline variant p.Ser1029Ala and mucosal malignant melanoma: An autopsy study. Molecular and Clinical Oncology, 2021, 16, 32.	1.0	0
4	Association study of long non-coding RNA HOTAIR rs920778 polymorphism with the risk of cancer in an elderly Japanese population. Gene, 2020, 729, 144263.	2.2	11
5	Clinicopathological characteristics of gastric cancer with carbohydrate antigen 19â€9 expression occurring in elderly individuals: An autopsy study. Pathology International, 2020, 70, 92-100.	1.3	4
6	Association of CYP2A6 gene deletion with cancers in Japanese elderly: an autopsy study. BMC Cancer, 2020, 20, 186.	2.6	4
7	Association between rs1229984 in ADH1B and cancer prevalence in a Japanese population. Molecular and Clinical Oncology, 2020, 12, 503-510.	1.0	3
8	The stem cellâ€specific intermediate filament nestin missense variation p.A1199P is associated with pancreatic cancer. Oncology Letters, 2019, 17, 4647-4654.	1.8	2
9	Metabolic and Immunological Shifts during Mid-to-Late Gestation Influence Maternal Blood Methylation of CPT1A and SREBF1. International Journal of Molecular Sciences, 2019, 20, 1066.	4.1	13
10	Self-organizing scale-free patterns in a phase-modulated periodic connecting system. BMC Research Notes, 2019, 12, 122.	1.4	1
11	Effectiveness of personal genomic testing for disease-prevention behavior when combined with careful consultation with a physician: a preliminary study. BMC Research Notes, 2018, 11, 223.	1.4	3
12	Relationship between pancreatic intraepithelial neoplasias, pancreatic ductal adenocarcinomas, and single nucleotide polymorphisms in autopsied elderly patients. Genes Chromosomes and Cancer, 2018, 57, 12-18.	2.8	6
13	Inverse Association Between Height-Increasing Alleles and Extreme Longevity in Japanese Women. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2018, 73, 588-595.	3.6	9
14	Long noncoding RNA variations in cardiometabolic diseases. Journal of Human Genetics, 2017, 62, 97-104.	2.3	40
15	Identification of EGFLAM, SPATC1L and RNASE13 as novel susceptibility loci for aortic aneurysm in Japanese individuals by exome-wide association studies. International Journal of Molecular Medicine, 2017, 39, 1091-1100.	4.0	15
16	Identification of six polymorphisms as novel susceptibility loci for ischemic or hemorrhagic stroke by exome-wide association studies. International Journal of Molecular Medicine, 2017, 39, 1477-1491.	4.0	16
17	Association of ZFHX3 gene variation with atrial fibrillation, cerebral infarction, and lung thromboembolism: An autopsy study. Journal of Cardiology, 2017, 70, 180-184.	1.9	18
18	Identification of TNFSF13, SPATC1L, SLC22A25 and SALL4 as novel susceptibility loci for atrial fibrillation by an exome-wide association study. Molecular Medicine Reports, 2017, 16, 5823-5832.	2.4	8

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19	Early gestational maternal low-protein diet diminishes hepatic response to fasting in young adult male mice. Scientific Reports, 2017, 7, 9812.	3.3	2
20	Identification of STXBP2 as a novel susceptibility locus for myocardial infarction in Japanese individuals by an exome-wide association study. Oncotarget, 2017, 8, 33527-33535.	1.8	14
21	Identification of C21orf59 and ATG2A as novel determinants of renal function-related traits in Japanese by exome-wide association studies. Oncotarget, 2017, 8, 45259-45273.	1.8	9
22	Identification of rs7350481 at chromosome 11q23.3 as a novel susceptibility locus for metabolic syndrome in Japanese individuals by an exome-wide association study. Oncotarget, 2017, 8, 39296-39308.	1.8	13
23	Identification of eight genetic variants as novel determinants of dyslipidemia in Japanese by exome-wide association studies. Oncotarget, 2017, 8, 38950-38961.	1.8	6
24	Identification of polymorphisms in 12q24.1, ACAD10, and BRAP as novel genetic determinants of blood pressure in Japanese by exome-wide association studies. Oncotarget, 2017, 8, 43068-43079.	1.8	13
25	Identification of five genetic variants as novel determinants of type 2 diabetes mellitus in Japanese by exome-wide association studies. Oncotarget, 2017, 8, 80492-80505.	1.8	3
26	An Alpha-kinase 2 Gene Variant Disrupts Filamentous Actin Localization in the Surface Cells of Colorectal Cancer Spheroids. Anticancer Research, 2017, 37, 3855-3862.	1.1	5
27	Association of polymorphisms of the transporter associated with antigen processing (TAP2) gene with pulmonary tuberculosis in an elderly Japanese population. Apmis, 2016, 124, 675-680.	2.0	8
28	Exome-wide Association Study IdentifiesCLEC3BMissense Variant p.S106G as Being Associated With Extreme Longevity in East Asian Populations. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2016, 72, glw074.	3.6	13
29	Genetic risk score based on the lifetime prevalence of femoral fracture in 924 consecutive autopsies of Japanese males. Journal of Bone and Mineral Metabolism, 2016, 34, 685-691.	2.7	6
30	Association of non-synonymous variants in and genes with abdominal aortic aneurysm: an autopsy study. Journal of Geriatric Cardiology, 2016, 13, 960-967.	0.2	1
31	Associations between the orexin (hypocretin) receptor 2 gene polymorphism Val308Ile and nicotine dependence in genome-wide and subsequent association studies. Molecular Brain, 2015, 8, 50.	2.6	23
32	Association of the chromodomain helicase DNAâ€binding protein 4 (CHD4) missense variation p.D140E with cancer: Potential interaction with smoking. Genes Chromosomes and Cancer, 2015, 54, 122-128.	2.8	14
33	A missense single nucleotide polymorphism, V114I of the Werner syndrome gene, is associated with risk of osteoporosis and femoral fracture in the Japanese population. Journal of Bone and Mineral Metabolism, 2015, 33, 694-700.	2.7	6
34	Empirical analysis of scale-free patterns of connectivity in medical term occurrence. Journal of Japan Society for Fuzzy Theory and Intelligent Informatics, 2015, 27, 616-620.	0.0	1
35	No association between catechol-O-methyltransferase (COMT) genotype and attention deficit hyperactivity disorder (ADHD) in Japanese children. Brain and Development, 2014, 36, 620-625.	1.1	13
36	Association of the RYR3 gene polymorphisms with atherosclerosis in elderly Japanese population. BMC Cardiovascular Disorders, 2014, 14, 6.	1.7	5

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37	Epistasis effects of COMT and MTHFR on inter-individual differences in mental health: Under the inverted U-shaped prefrontal dopamine model. Biochemical and Biophysical Research Communications, 2014, 451, 574-579.	2.1	13
38	Independent strong association of HLA-A*02:06 and HLA-B*44:03 with cold medicine-related Stevens-Johnson syndrome with severe mucosal involvement. Scientific Reports, 2014, 4, 4862.	3.3	83
39	Association Analysis of Single Nucleotide Polymorphisms in miR-146a and miR-196a2 on the Prevalence of Cancer in Elderly Japanese: A Case-Control Study. Asian Pacific Journal of Cancer Prevention, 2014, 15, 2101-2107.	1.2	39
40	Polymorphisms of the formylpeptide receptor gene (FPR1) and susceptibility to stomach cancer in 1531 consecutive autopsy cases. Biochemical and Biophysical Research Communications, 2011, 405, 356-361.	2.1	28
41	Lymphotoxin-alpha polymorphisms and presence of cancer in 1,536 consecutive autopsy cases. BMC Cancer, 2008, 8, 235.	2.6	22
42	Functional polymorphisms in carboxylesterase1A2 (CES1A2) gene involves specific protein 1 (Sp1) binding sites. Biochemical and Biophysical Research Communications, 2008, 369, 939-942.	2.1	44
43	Polymorphisms in pro- and anti-inflammatory cytokine genes and susceptibility to atherosclerosis: a pathological study of 1503 consecutive autopsy cases. Human Molecular Genetics, 2007, 16, 592-599.	2.9	55
44	Bone morphogenetic protein-7 and interferon-alpha synergistically suppress hepatitis C virus replicon. Biochemical and Biophysical Research Communications, 2007, 357, 467-473.	2.1	16
45	Dyslipidemia is a major determinant of systemic atherosclerosis in the elderly: An autopsy study. Geriatrics and Gerontology International, 2007, 7, 229-237.	1.5	7
46	Association of Transforming Growth Factorâ€“Î²1 Functional Polymorphisms with Natural Clearance of Hepatitis C Virus. Journal of Infectious Diseases, 2006, 193, 1371-1374.	4.0	47
47	A Single Nucleotide Polymorphism in the Carboxylesterase Gene Is Associated with the Responsiveness to Imidapril Medication and the Promoter Activity. Hypertension Research, 2005, 28, 719-725.	2.7	89
48	Genetic variations in humans associated with differences in the course of hepatitis C. Biochemical and Biophysical Research Communications, 2004, 317, 335-341.	2.1	50