

Keitaro Matsuo

List of Publications by Year in descending order

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842
papers

40,626
citations

2963

93
h-index

7496

151
g-index

869
all docs

869
docs citations

869
times ranked

45413
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
3	Association between Body-Mass Index and Risk of Death in More Than 1 Million Asians. <i>New England Journal of Medicine</i> , 2011, 364, 719-729.	13.9	730
4	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	9.4	513
5	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
6	High Complete Remission Rate and Promising Outcome by Combination of Imatinib and Chemotherapy for Newly Diagnosed BCR-ABL ⁺ Positive Acute Lymphoblastic Leukemia: A Phase II Study by the Japan Adult Leukemia Study Group. <i>Journal of Clinical Oncology</i> , 2006, 24, 460-466.	0.8	430
7	Genome-wide association studies identify four ER negative ⁺ specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
8	Meta-Analysis of Randomized Clinical Trials Comparing Cisplatin to Carboplatin in Patients With Advanced Non ⁺ Small-Cell Lung Cancer. <i>Journal of Clinical Oncology</i> , 2004, 22, 3852-3859.	0.8	373
9	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
10	Improved survival of gastric cancer with tumour Epstein ⁺ Barr virus positivity: an international pooled analysis. <i>Gut</i> , 2014, 63, 236-243.	6.1	309
11	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , 2013, 144, 799-807.e24.	0.6	292
12	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
13	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	9.4	286
14	Meta-analysis identifies nine new loci associated with rheumatoid arthritis in the Japanese population. <i>Nature Genetics</i> , 2012, 44, 511-516.	9.4	285
15	Genome-wide array-based CGH for mantle cell lymphoma: identification of homozygous deletions of the proapoptotic gene BIM. <i>Oncogene</i> , 2005, 24, 1348-1358.	2.6	282
16	Genetic abnormalities in myelodysplasia and secondary acute myeloid leukemia: impact on outcome of stem cell transplantation. <i>Blood</i> , 2017, 129, 2347-2358.	0.6	268
17	Prognostic significance of FLT3 internal tandem duplication and tyrosine kinase domain mutations for acute myeloid leukemia: a meta-analysis. <i>Leukemia</i> , 2005, 19, 1345-1349.	3.3	267
18	Transplantation of allogeneic hematopoietic stem cells for adult T-cell leukemia: a nationwide retrospective study. <i>Blood</i> , 2010, 116, 1369-1376.	0.6	255

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19	Differences in incidence and trends of haematological malignancies in Japan and the United States. <i>British Journal of Haematology</i> , 2014, 164, 536-545.	1.2	250
20	High-risk HLA allele mismatch combinations responsible for severe acute graft-versus-host disease and implication for its molecular mechanism. <i>Blood</i> , 2007, 110, 2235-2241.	0.6	246
21	Association between body mass index and cardiovascular disease mortality in east Asians and south Asians: pooled analysis of prospective data from the Asia Cohort Consortium. <i>BMJ</i> , 2013, 347, f5446-f5446.	3.0	239
22	Genetic Polymorphisms of the Human PNPLA3 Gene Are Strongly Associated with Severity of Non-Alcoholic Fatty Liver Disease in Japanese. <i>PLoS ONE</i> , 2012, 7, e38322.	1.1	228
23	Phase III Trial Comparing Docetaxel and Cisplatin Combination Chemotherapy With Mitomycin, Vindesine, and Cisplatin Combination Chemotherapy With Concurrent Thoracic Radiotherapy in Locally Advanced Non-Small-Cell Lung Cancer: OLCSG 0007. <i>Journal of Clinical Oncology</i> , 2010, 28, 3299-3306.	0.8	225
24	Prognostic relevance of genetic alterations in diffuse lower-grade gliomas. <i>Neuro-Oncology</i> , 2018, 20, 66-77.	0.6	225
25	Relationship between Smoking and Obesity: A Cross-Sectional Study of 499,504 Middle-Aged Adults in the UK General Population. <i>PLoS ONE</i> , 2015, 10, e0123579.	1.1	222
26	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
27	Heterogeneous Distribution of EGFR Mutations Is Extremely Rare in Lung Adenocarcinoma. <i>Journal of Clinical Oncology</i> , 2011, 29, 2972-2977.	0.8	218
28	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. <i>Nature Genetics</i> , 2014, 46, 533-542.	9.4	212
29	Polymerase Chain Reaction with Confronting Two-pair Primers for Polymorphism Genotyping. <i>Japanese Journal of Cancer Research</i> , 2000, 91, 865-868.	1.7	210
30	Effect of hepatitis C virus infection on the risk of non-Hodgkin's lymphoma: A meta-analysis of epidemiological studies. <i>Cancer Science</i> , 2004, 95, 745-752.	1.7	208
31	Role of Adjuvant Chemotherapy in Patients With Resected Non-Small-Cell Lung Cancer: Reappraisal With a Meta-Analysis of Randomized Controlled Trials. <i>Journal of Clinical Oncology</i> , 2004, 22, 3860-3867.	0.8	202
32	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	2.6	201
33	Comparison of genome profiles for identification of distinct subgroups of diffuse large B-cell lymphoma. <i>Blood</i> , 2005, 106, 1770-1777.	0.6	196
34	Relapse-Related Molecular Signature in Lung Adenocarcinomas Identifies Patients With Dismal Prognosis. <i>Journal of Clinical Oncology</i> , 2009, 27, 2793-2799.	0.8	194
35	Association between polymorphisms of folate- and methionine-metabolizing enzymes and susceptibility to malignant lymphoma. <i>Blood</i> , 2001, 97, 3205-3209.	0.6	185
36	Biological significance of HLA locus matching in unrelated donor bone marrow transplantation. <i>Blood</i> , 2015, 125, 1189-1197.	0.6	185

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37	Gene-environment interaction between an aldehyde dehydrogenase-2 (ALDH2) polymorphism and alcohol consumption for the risk of esophageal cancer. <i>Carcinogenesis</i> , 2001, 22, 913-916.	1.3	176
38	Replication of Lung Cancer Susceptibility Loci at Chromosomes 15q25, 5p15, and 6p21: A Pooled Analysis From the International Lung Cancer Consortium. <i>Journal of the National Cancer Institute</i> , 2010, 102, 959-971.	3.0	174
39	Combination of intensive chemotherapy and imatinib can rapidly induce high-quality complete remission for a majority of patients with newly diagnosed BCR-ABL-positive acute lymphoblastic leukemia. <i>Blood</i> , 2004, 104, 3507-3512.	0.6	173
40	Genome-wide association analyses in east Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013, 45, 191-196.	9.4	173
41	Efficacy of allogeneic hematopoietic stem cell transplantation depends on cytogenetic risk for acute myeloid leukemia in first disease remission. <i>Cancer</i> , 2005, 103, 1652-1658.	2.0	169
42	Smoking Increases the Treatment Failure for Helicobacter pylori Eradication. <i>American Journal of Medicine</i> , 2006, 119, 217-224.	0.6	164
43	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	7.7	157
44	Variant ALDH2 is associated with accelerated progression of bone marrow failure in Japanese Fanconi anemia patients. <i>Blood</i> , 2013, 122, 3206-3209.	0.6	156
45	Body Mass Index and Diabetes in Asia: A Cross-Sectional Pooled Analysis of 900,000 Individuals in the Asia Cohort Consortium. <i>PLoS ONE</i> , 2011, 6, e19930.	1.1	154
46	Allogeneic hematopoietic stem cell transplantation as part of postremission therapy improves survival for adult patients with high-risk acute lymphoblastic leukemia. <i>Cancer</i> , 2006, 106, 2657-2663.	2.0	153
47	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv279.	3.0	152
48	Rb Loss and KRAS Mutation Are Predictors of the Response to Platinum-Based Chemotherapy in Pancreatic Neuroendocrine Neoplasm with Grade 3: A Japanese Multicenter Pancreatic NEN-G3 Study. <i>Clinical Cancer Research</i> , 2017, 23, 4625-4632.	3.2	150
49	Teeth Loss and Risk of Cancer at 14 Common Sites in Japanese. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 1222-1227.	1.1	149
50	Mutational and Epigenetic Evidence for Independent Pathways for Lung Adenocarcinomas Arising in Smokers and Never Smokers. <i>Cancer Research</i> , 2006, 66, 1371-1375.	0.4	147
51	Feasibility of HLA-haploidentical hematopoietic stem cell transplantation between noninherited maternal antigen (NIMA)-mismatched family members linked with long-term fetomaternal microchimerism. <i>Blood</i> , 2004, 104, 3821-3828.	0.6	145
52	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013, 4, 1628.	5.8	144
53	Increased risk of lung cancer in individuals with a family history of the disease: A pooled analysis from the International Lung Cancer Consortium. <i>European Journal of Cancer</i> , 2012, 48, 1957-1968.	1.3	143
54	Interleukin 1 Polymorphisms, Lifestyle Factors, and Helicobacter pylori Infection. <i>Japanese Journal of Cancer Research</i> , 2001, 92, 383-389.	1.7	138

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55	Allogeneic hematopoietic stem cell transplantation from family members other than HLA-identical siblings over the last decade (1991-2000). <i>Blood</i> , 2003, 102, 1541-1547.	0.6	138
56	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	5.8	138
57	Genome-Wide Association Study in East Asians Identifies Novel Susceptibility Loci for Breast Cancer. <i>PLoS Genetics</i> , 2012, 8, e1002532.	1.5	137
58	Genome-wide association analysis in East Asians identifies breast cancer susceptibility loci at 1q32.1, 5q14.3 and 15q26.1. <i>Nature Genetics</i> , 2014, 46, 886-890.	9.4	135
59	Cigarette smoking and gastric cancer in the Stomach Cancer Pooling (StoP) Project. <i>European Journal of Cancer Prevention</i> , 2018, 27, 124-133.	0.6	134
60	Prospective Validation for Prediction of Gefitinib Sensitivity by Epidermal Growth Factor Receptor Gene Mutation in Patients with Non-Small Cell Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2007, 2, 22-28.	0.5	134
61	Association between type 2 diabetes and risk of cancer mortality: a pooled analysis of over 771,000 individuals in the Asia Cohort Consortium. <i>Diabetologia</i> , 2017, 60, 1022-1032.	2.9	132
62	A Genome-Wide Association Analysis Identified a Novel Susceptible Locus for Pathological Myopia at 11q24.1. <i>PLoS Genetics</i> , 2009, 5, e1000660.	1.5	131
63	HLA mismatch combinations associated with decreased risk of relapse: implications for the molecular mechanism. <i>Blood</i> , 2009, 113, 2851-2858.	0.6	129
64	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	3.0	129
65	Risk factors for head and neck cancer in young adults: a pooled analysis in the INHANCE consortium. <i>International Journal of Epidemiology</i> , 2015, 44, 169-185.	0.9	128
66	Gene-environment interactions between the smoking habit and polymorphisms in the DNA repair genes, APE1 Asp148Glu and XRCC1 Arg399Gln, in Japanese lung cancer risk. <i>Carcinogenesis</i> , 2004, 25, 1395-1401.	1.3	126
67	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
68	Hepatic toxicity and prognosis in hepatitis C virus-infected patients with diffuse large B-cell lymphoma treated with rituximab-containing chemotherapy regimens: a Japanese multicenter analysis. <i>Blood</i> , 2010, 116, 5119-5125.	0.6	123
69	Identification of epigenetic aberrant promoter methylation in serum DNA is useful for early detection of lung cancer. <i>Clinical Cancer Research</i> , 2005, 11, 1219-25.	3.2	123
70	DNA Methylation in Peripheral Blood: A Potential Biomarker for Cancer <i>Molecular Epidemiology</i> . <i>Journal of Epidemiology</i> , 2012, 22, 384-394.	1.1	121
71	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
72	A genome-wide association study identifies a breast cancer risk variant in ERBB4 at 2q34: results from the Seoul Breast Cancer Study. <i>Breast Cancer Research</i> , 2012, 14, R56.	2.2	118

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73	Alcohol Dehydrogenase 2 His47Arg Polymorphism Influences Drinking Habit Independently of Aldehyde Dehydrogenase 2 Glu487Lys Polymorphism: Analysis of 2,299 Japanese Subjects. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1009-1013.	1.1	116
74	Effects of HLA Allele and Killer Immunoglobulin-Like Receptor Ligand Matching on Clinical Outcome in Leukemia Patients Undergoing Transplantation With T-cellâ€“Replete Marrow From an Unrelated Donor. <i>Biology of Blood and Marrow Transplantation</i> , 2007, 13, 315-328.	2.0	116
75	Epigenetic Profiles Distinguish Malignant Pleural Mesothelioma from Lung Adenocarcinoma. <i>Cancer Research</i> , 2009, 69, 9073-9082.	0.4	116
76	Diet and the risk of head and neck cancer: a pooled analysis in the INHANCE consortium. <i>Cancer Causes and Control</i> , 2012, 23, 69-88.	0.8	116
77	A Functional Single Nucleotide Polymorphism in Mucin 1, at Chromosome 1q22, Determines Susceptibility to Diffuse-Type Gastric Cancer. <i>Gastroenterology</i> , 2011, 140, 892-902.	0.6	114
78	A genome-wide association study identifies two susceptibility loci for duodenal ulcer in the Japanese population. <i>Nature Genetics</i> , 2012, 44, 430-434.	9.4	114
79	<i>CFH</i>and<i>ARMS2</i>Variations in Age-Related Macular Degeneration, Polypoidal Choroidal Vasculopathy, and Retinal Angiomatic Proliferation. , 2010, 51, 5914.		112
80	Obesity and survival among women with ovarian cancer: results from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2015, 113, 817-826.	2.9	111
81	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , 2019, 156, 1455-1466.	0.6	111
82	Impact of graft-versus-host disease on outcomes after allogeneic hematopoietic cell transplantation for adult T-cell leukemia: a retrospective cohort study. <i>Blood</i> , 2012, 119, 2141-2148.	0.6	110
83	Gene-environment Interactions and Polymorphism Studies of Cancer Risk in the Hospital-based Epidemiologic Research Program at Aichi Cancer Center II (HERPACC-II). <i>Asian Pacific Journal of Cancer Prevention</i> , 2001, 2, 99-107.	0.5	109
84	Alcohol Drinking and Colorectal Cancer in Japanese: A Pooled Analysis of Results from Five Cohort Studies. <i>American Journal of Epidemiology</i> , 2008, 167, 1397-1406.	1.6	107
85	Effect of selective lymph node dissection based on patterns of lobe-specific lymph node metastases on patient outcome in patients with resectable nonâ€“small cell lung cancer: A large-scale retrospective cohort study applying a propensity score. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2010, 139, 1001-1006.	0.4	107
86	Identification of a Functional Genetic Variant at 16q12.1 for Breast Cancer Risk: Results from the Asia Breast Cancer Consortium. <i>PLoS Genetics</i> , 2010, 6, e1001002.	1.5	107
87	ABCA Transporter Gene Expression and Poor Outcome in Epithelial Ovarian Cancer. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	3.0	107
88	Gene-environment interaction involved in oral carcinogenesis: molecular epidemiological study for metabolic and DNA repair gene polymorphisms. <i>Journal of Oral Pathology and Medicine</i> , 2006, 35, 11-18.	1.4	105
89	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
90	Nonâ€“Hodgkin lymphoma and obesity: A pooled analysis from the InterLymph Consortium. <i>International Journal of Cancer</i> , 2008, 122, 2062-2070.	2.3	104

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91	One-carbon metabolism-related gene polymorphisms and risk of breast cancer. <i>Carcinogenesis</i> , 2008, 29, 356-362.	1.3	104
92	Risk estimation model for nonalcoholic fatty liver disease in the Japanese using multiple genetic markers. <i>PLoS ONE</i> , 2018, 13, e0185490.	1.1	104
93	Association of Diabetes With All-Cause and Cause-Specific Mortality in Asia. <i>JAMA Network Open</i> , 2019, 2, e192696.	2.8	103
94	Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 632-640.	0.5	103
95	Clinicoradiologic characteristics of patients with lung adenocarcinoma harboring EML4-ALK fusion oncogene. <i>Lung Cancer</i> , 2012, 77, 319-325.	0.9	102
96	Tobacco Smoking and Mortality in Asia. <i>JAMA Network Open</i> , 2019, 2, e191474.	2.8	102
97	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
98	The role of oral hygiene in head and neck cancer: results from International Head and Neck Cancer Epidemiology (INHANCE) consortium. <i>Annals of Oncology</i> , 2016, 27, 1619-1625.	0.6	101
99	Alcohol and head and neck cancer. <i>Cancer and Metastasis Reviews</i> , 2017, 36, 425-434.	2.7	101
100	Meta-analysis of the effects of prokinetic agents in patients with functional dyspepsia. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2007, 22, 304-310.	1.4	100
101	Body Mass Index and Mortality From All Causes and Major Causes in Japanese: Results of a Pooled Analysis of 7 Large-Scale Cohort Studies. <i>Journal of Epidemiology</i> , 2011, 21, 417-430.	1.1	100
102	Exposure to secondhand tobacco smoke and lung cancer by histological type: A pooled analysis of the International Lung Cancer Consortium (ILCCO). <i>International Journal of Cancer</i> , 2014, 135, 1918-1930.	2.3	100
103	Novel Germline Mutation in the Transmembrane Domain of HER2 in Familial Lung Adenocarcinomas. <i>Journal of the National Cancer Institute</i> , 2014, 106, djt338.	3.0	99
104	Alcohol and cigarette consumption predict mortality in patients with head and neck cancer: a pooled analysis within the International Head and Neck Cancer Epidemiology (INHANCE) Consortium. <i>Annals of Oncology</i> , 2017, 28, 2843-2851.	0.6	99
105	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	2.6	98
106	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98
107	Burden of Total and Cause-Specific Mortality Related to Tobacco Smoking among Adults Aged ≥45 Years in Asia: A Pooled Analysis of 21 Cohorts. <i>PLoS Medicine</i> , 2014, 11, e1001631.	3.9	98
108	Winner's Curse Correction and Variable Thresholding Improve Performance of Polygenic Risk Modeling Based on Genome-Wide Association Study Summary-Level Data. <i>PLoS Genetics</i> , 2016, 12, e1006493.	1.5	98

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109	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	0.6	97
110	Pre-transplant imatinib-based therapy improves the outcome of allogeneic hematopoietic stem cell transplantation for BCR-ABL-positive acute lymphoblastic leukemia. <i>Leukemia</i> , 2011, 25, 41-47.	3.3	96
111	Genetic Variants on Chromosome 1q41 Influence Ocular Axial Length and High Myopia. <i>PLoS Genetics</i> , 2012, 8, e1002753.	1.5	95
112	Diabetes mellitus and cancer risk: Pooled analysis of eight cohort studies in Japan. <i>Cancer Science</i> , 2013, 104, 1499-1507.	1.7	94
113	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
114	NAD(P)H: quinone oxidoreductase 1 (NQO1) C609T polymorphism and the risk of eight cancers for Japanese. <i>International Journal of Clinical Oncology</i> , 2002, 7, 103-108.	1.0	93
115	International Lung Cancer Consortium: Pooled Analysis of Sequence Variants in DNA Repair and Cell Cycle Pathways. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 3081-3089.	1.1	93
116	Profile of Participants and Genotype Distributions of 108 Polymorphisms in a Cross-Sectional Study of Associations of Genotypes With Lifestyle and Clinical Factors: A Project in the Japan Multi-Institutional Collaborative Cohort (J-MICC) Study. <i>Journal of Epidemiology</i> , 2011, 21, 223-235.	1.1	92
117	Genome-wide association study identifies breast cancer risk variant at 10q21.2: results from the Asia Breast Cancer Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4991-4999.	1.4	92
118	Two Aldehyde Clearance Systems Are Essential to Prevent Lethal Formaldehyde Accumulation in Mice and Humans. <i>Molecular Cell</i> , 2020, 80, 996-1012.e9.	4.5	92
119	Meta- and Pooled Analyses of the Methylenetetrahydrofolate Reductase (MTHFR) C677T Polymorphism and Colorectal Cancer: A HuGE-GSEC Review. <i>American Journal of Epidemiology</i> , 2009, 170, 1207-1221.	1.6	91
120	The human AIRE gene at chromosome 21q22 is a genetic determinant for the predisposition to rheumatoid arthritis in Japanese population. <i>Human Molecular Genetics</i> , 2011, 20, 2680-2685.	1.4	90
121	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	1.4	90
122	Addition of platinum compounds to a new agent in patients with advanced non-small-cell lung cancer: a literature based meta-analysis of randomised trials. <i>Annals of Oncology</i> , 2004, 15, 1782-1789.	0.6	89
123	Markedly improved outcomes and acceptable toxicity in adolescents and young adults with acute lymphoblastic leukemia following treatment with a pediatric protocol: a phase II study by the Japan Adult Leukemia Study Group. <i>Blood Cancer Journal</i> , 2014, 4, e252-e252.	2.8	88
124	HLA-Haploidentical Peripheral Blood Stem Cell Transplantation with Post-Transplant Cyclophosphamide after Busulfan-Containing Reduced-Intensity Conditioning. <i>Biology of Blood and Marrow Transplantation</i> , 2015, 21, 1646-1652.	2.0	88
125	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.	5.8	88
126	No associations of p73 G4C14-to-A4T14 at exon 2 and p53 Arg72Pro polymorphisms with the risk of digestive tract cancers in Japanese. <i>Cancer Letters</i> , 2002, 181, 81-85.	3.2	86

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127	Association of p53 codon arg72pro and p73 G4C14-to-A4T14 at exon 2 genetic polymorphisms with the risk of Japanese breast cancer. <i>Breast Cancer</i> , 2003, 10, 307-311.	1.3	86
128	Risk factors differ for non-small-cell lung cancers with and without EGFR mutation: assessment of smoking and sex by a case-control study in Japanese. <i>Cancer Science</i> , 2007, 98, 96-101.	1.7	86
129	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. <i>Human Molecular Genetics</i> , 2013, 22, 2539-2550.	1.4	86
130	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	5.8	86
131	Dominant negative isoform of the Ikaros gene in patients with adult B-cell acute lymphoblastic leukemia. <i>Cancer Research</i> , 2000, 60, 4062-5.	0.4	86
132	Alcohol consumption and gastric cancer risk: A pooled analysis within the StoP project consortium. <i>International Journal of Cancer</i> , 2017, 141, 1950-1962.	2.3	85
133	Prospective monitoring of <i>BCR-ABL1</i> transcript levels in patients with Philadelphia chromosome-positive acute lymphoblastic leukaemia undergoing imatinib-combined chemotherapy. <i>British Journal of Haematology</i> , 2008, 143, 503-510.	1.2	84
134	A genetic risk predictor for breast cancer using a combination of low-penetrance polymorphisms in a Japanese population. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 711-721.	1.1	84
135	Associations between polymorphisms in the thymidylate synthase and serine hydroxymethyltransferase genes and susceptibility to malignant lymphoma. <i>Haematologica</i> , 2003, 88, 159-66.	1.7	83
136	Analysis of Genetic Variants in Never-Smokers with Lung Cancer Facilitated by an Internet-Based Blood Collection Protocol: A Preliminary Report. <i>Clinical Cancer Research</i> , 2010, 16, 755-763.	3.2	82
137	The INHANCE consortium: toward a better understanding of the causes and mechanisms of head and neck cancer. <i>Oral Diseases</i> , 2015, 21, 685-693.	1.5	82
138	Dietary Risk Factors for Colon and Rectal Cancers: A Comparative Case-Control Study. <i>Journal of Epidemiology</i> , 2006, 16, 125-135.	1.1	81
139	The Impact of Sex and Smoking Status on the Mutational Spectrum of Epidermal Growth Factor Receptor Gene in Non-small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2007, 13, 5763-5768.	3.2	81
140	One-carbon metabolism related gene polymorphisms interact with alcohol drinking to influence the risk of colorectal cancer in Japan. <i>Carcinogenesis</i> , 2005, 26, 2164-2171.	1.3	80
141	Prognostic Significance of T-Cell or Cytotoxic Molecules Phenotype in Classical Hodgkin's Lymphoma: A Clinicopathologic Study. <i>Journal of Clinical Oncology</i> , 2006, 24, 4626-4633.	0.8	80
142	Insulin-like growth factor (IGF)-I and IGF-binding protein 3 and the risk of premenopausal breast cancer: A meta-analysis of literature. <i>International Journal of Cancer</i> , 2004, 111, 293-297.	2.3	79
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421	Impact of being overweight on outcomes of hematopoietic SCT: a meta-analysis. <i>Bone Marrow Transplantation</i> , 2014, 49, 66-72.	1.3	25
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