

Daehee Kang

List of Publications by Year in descending order

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Version: 2024-02-01

372
papers

23,618
citations

10351

72
h-index

11899

134
g-index

385
all docs

385
docs citations

385
times ranked

30812
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	13.7	2,165
2	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
3	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
4	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
5	A common coding variant in <i>CASP8</i> is associated with breast cancer risk. <i>Nature Genetics</i> , 2007, 39, 352-358.	9.4	591
6	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
7	Multiple independent variants at the <i>TERT</i> locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
8	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	9.4	434
9	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
10	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
11	Large-scale genome-wide association studies in east Asians identify new genetic loci influencing metabolic traits. <i>Nature Genetics</i> , 2011, 43, 990-995.	9.4	270
12	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	9.4	256
13	Meta-analysis identifies multiple loci associated with kidney function-related traits in east Asian populations. <i>Nature Genetics</i> , 2012, 44, 904-909.	9.4	254
14	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> , The, 2013, 347, f5446-f5446.	3.0	239
15	Pooled Analysis and Meta-analysis of Glutathione S-Transferase M1 and Bladder Cancer: A HuGE Review. <i>American Journal of Epidemiology</i> , 2002, 156, 95-109.	1.6	209
16	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
17	Young age: an independent risk factor for disease-free survival in women with operable breast cancer. <i>BMC Cancer</i> , 2004, 4, 82.	1.1	197
18	Meta-analysis of genome-wide association studies in East Asian-ancestry populations identifies four new loci for body mass index. <i>Human Molecular Genetics</i> , 2014, 23, 5492-5504.	1.4	192

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19	Correlates associated with participation in physical activity among adults: a systematic review of reviews and update. BMC Public Health, 2017, 17, 356.	1.2	192
20	Identification of Serum MicroRNAs as Novel Non-Invasive Biomarkers for Early Detection of Gastric Cancer. PLoS ONE, 2012, 7, e33608.	1.1	169
21	Genome-wide association study of Crohn's disease in Koreans revealed three new susceptibility loci and common attributes of genetic susceptibility across ethnic populations. Gut, 2014, 63, 80-87.	6.1	157
22	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
23	Urinary 1-hydroxypyrene and other PAH metabolites as biomarkers of exposure to environmental PAH in air particulate matter. Toxicology Letters, 1999, 108, 191-199.	0.4	154
24	Body Mass Index and Diabetes in Asia: A Cross-Sectional Pooled Analysis of 900,000 Individuals in the Asia Cohort Consortium. PLoS ONE, 2011, 6, e19930.	1.1	154
25	Effect of Diet and Helicobacter pylori Infection to the Risk of Early Gastric Cancer.. Journal of Epidemiology, 2003, 13, 162-168.	1.1	139
26	Genome-Wide Association Study in East Asians Identifies Novel Susceptibility Loci for Breast Cancer. PLoS Genetics, 2012, 8, e1002532.	1.5	137
27	Genome-wide association analysis in East Asians identifies breast cancer susceptibility loci at 1q32.1, 5q14.3 and 15q26.1. Nature Genetics, 2014, 46, 886-890.	9.4	135
28	Association between type 2 diabetes and risk of cancer mortality: a pooled analysis of over 771,000 individuals in the Asia Cohort Consortium. Diabetologia, 2017, 60, 1022-1032.	2.9	132
29	Meta- and Pooled Analysis of GSTT1 and Lung Cancer: A HuGE-GSEC Review. American Journal of Epidemiology, 2006, 164, 1027-1042.	1.6	130
30	Interindividual differences in the concentration of 1-hydroxypyrene-glucuronide in urine and polycyclic aromatic hydrocarbon-DNA adducts in peripheral white blood cells after charbroiled beef consumption. Carcinogenesis, 1995, 16, 1079-1085.	1.3	129
31	Functional Variant of Manganese Superoxide Dismutase (<i>SOD2 V16A</i>) Polymorphism Is Associated with Prostate Cancer Risk in the Prostate, Lung, Colorectal, and Ovarian Cancer Study. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 1581-1586.	1.1	129
32	Paternal smoking, genetic polymorphisms in CYP1A1 and childhood leukemia risk. Leukemia Research, 2009, 33, 250-258.	0.4	129
33	Female Breast Cancer Incidence Among Asian and Western Populations: More Similar Than Expected. Journal of the National Cancer Institute, 2015, 107, .	3.0	127
34	Gene-environment interactions between the smoking habit and polymorphisms in the DNA repair genes, APE1 Asp148Glu and XRCC1 Arg399Gln, in Japanese lung cancer risk. Carcinogenesis, 2004, 25, 1395-1401.	1.3	126
35	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
36	Effects of bisphenol A on breast cancer and its risk factors. Archives of Toxicology, 2009, 83, 281-285.	1.9	121

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37	DNA Methylation in Peripheral Blood: A Potential Biomarker for Cancer Molecular Epidemiology. <i>Journal of Epidemiology</i> , 2012, 22, 384-394.	1.1	121
38	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
39	Intrauterine environments and breast cancer risk: meta-analysis and systematic review. <i>Breast Cancer Research</i> , 2008, 10, R8.	2.2	118
40	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
41	FGFR2 variants and breast cancer risk: fine-scale mapping using African American studies and analysis of chromatin conformation. <i>Human Molecular Genetics</i> , 2009, 18, 1692-1703.	1.4	110
42	Association between body size, weight change and depression: systematic review and meta-analysis. <i>British Journal of Psychiatry</i> , 2017, 211, 14-21.	1.7	110
43	Meat intake and cause-specific mortality: a pooled analysis of Asian prospective cohort studies. <i>American Journal of Clinical Nutrition</i> , 2013, 98, 1032-1041.	2.2	109
44	Rising prostate cancer rates in South Korea. <i>Prostate</i> , 2006, 66, 1285-1291.	1.2	108
45	Alcohol consumption, glutathione S-transferase M1 and T1 genetic polymorphisms and breast cancer risk. <i>Pharmacogenetics and Genomics</i> , 2000, 10, 301-309.	5.7	106
46	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
47	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. <i>Human Molecular Genetics</i> , 2015, 24, 1791-1800.	1.4	105
48	Identification of 1-hydroxypyrene glucuronide as a major pyrene metabolite in human urine by synchronous fluorescence spectroscopy and gas chromatography-mass spectrometry. <i>Carcinogenesis</i> , 1994, 15, 483-487.	1.3	103
49	Association of Diabetes With All-Cause and Cause-Specific Mortality in Asia. <i>JAMA Network Open</i> , 2019, 2, e192696.	2.8	103
50	Tobacco Smoking and Mortality in Asia. <i>JAMA Network Open</i> , 2019, 2, e191474.	2.8	102
51	Risk of Estrogen Receptor-Positive and -Negative Breast Cancer and Single-Nucleotide Polymorphism 2q35-rs13387042. <i>Journal of the National Cancer Institute</i> , 2009, 101, 1012-1018.	3.0	99
52	Genetic Polymorphisms of Selected DNA Repair Genes, Estrogen and Progesterone Receptor Status, and Breast Cancer Risk. <i>Clinical Cancer Research</i> , 2005, 11, 4620-4626.	3.2	98
53	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
54	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

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55	The Health Examinees (HEXA) Study: Rationale, Study Design and Baseline Characteristics. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 16, 1591-1597.	0.5	97
56	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
57	Relationship between the Val158Met polymorphism of catechol O-methyl transferase and breast cancer. <i>Pharmacogenetics and Genomics</i> , 2001, 11, 279-286.	5.7	92
58	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
59	Effect of BRCA1/2 mutation on short-term and long-term breast cancer survival: a systematic review and meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2010, 122, 11-25.	1.1	88
60	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
61	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
62	IARC Monographs: 40 Years of Evaluating Carcinogenic Hazards to Humans. <i>Environmental Health Perspectives</i> , 2015, 123, 507-514.	2.8	86
63	Nucleotide Excision Repair Gene Polymorphisms and Risk of Advanced Colorectal Adenoma: XPC Polymorphisms Modify Smoking-Related Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 306-311.	1.1	85
64	Nitric oxide synthase gene polymorphisms and prostate cancer risk. <i>Carcinogenesis</i> , 2009, 30, 621-625.	1.3	85
65	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1131-1139.	1.4	84
66	Polymorphisms in DNA repair genes and risk of non-Hodgkin lymphoma among women in Connecticut. <i>Human Genetics</i> , 2006, 119, 659-668.	1.8	81
67	Isoflavones from Phytoestrogens and Gastric Cancer Risk: A Nested Case-Control Study within the Korean Multicenter Cancer Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1292-1300.	1.1	80
68	XRCC1 genetic polymorphism and breast cancer risk. <i>Pharmacogenetics and Genomics</i> , 2002, 12, 335-338.	5.7	78
69	Estrogen and progesterone receptor status affect genome-wide DNA methylation profile in breast cancer. <i>Human Molecular Genetics</i> , 2010, 19, 4273-4277.	1.4	78
70	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
71	Genome-Wide Association Study of Ulcerative Colitis in Koreans Suggests Extensive Overlapping of Genetic Susceptibility With Caucasians. <i>Inflammatory Bowel Diseases</i> , 2013, 19, 954-966.	0.9	76
72	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.	2.6	76

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73	Meta- and Pooled Analyses of the Cytochrome P-450 1B1 Val432Leu Polymorphism and Breast Cancer: A HuGE-GSEC Review. <i>American Journal of Epidemiology</i> , 2006, 165, 115-125.	1.6	75
74	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
75	Epidemiology of Breast Cancer in Korea: Occurrence, High-Risk Groups, and Prevention. <i>Journal of Korean Medical Science</i> , 2002, 17, 1.	1.1	74
76	Association of paternal age at birth and the risk of breast cancer in offspring: a case control study. <i>BMC Cancer</i> , 2005, 5, 143.	1.1	74
77	Estrogen Receptor Alpha Gene Polymorphisms and Breast Cancer Risk. <i>Breast Cancer Research and Treatment</i> , 2003, 80, 127-131.	1.1	73
78	A Prospective Cohort Study on the Relationship of Sleep Duration With All-cause and Disease-specific Mortality in the Korean Multi-center Cancer Cohort Study. <i>Journal of Preventive Medicine and Public Health</i> , 2013, 46, 271-281.	0.7	73
79	Genome-wide association study of childhood acute lymphoblastic leukemia in Korea. <i>Leukemia Research</i> , 2010, 34, 1271-1274.	0.4	72
80	Genetic polymorphisms of cytochrome P450 19 and 1B1, alcohol use, and breast cancer risk in Korean women. <i>British Journal of Cancer</i> , 2003, 88, 675-678.	2.9	71
81	Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4693-4706.	1.4	71
82	Multiple HPV infection in cervical cancer screened by HPV DNA chip. <i>Cancer Letters</i> , 2003, 198, 187-192.	3.2	70
83	Fine scale mapping of the breast cancer 16q12 locus. <i>Human Molecular Genetics</i> , 2010, 19, 2507-2515.	1.4	68
84	Common Genetic Variants Associated with Breast Cancer in Korean Women and Differential Susceptibility According to Intrinsic Subtype. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 793-798.	1.1	68
85	Association between sleep duration and metabolic syndrome: a cross-sectional study. <i>BMC Public Health</i> , 2018, 18, 720.	1.2	68
86	Risk behaviours and benign prostatic hyperplasia. <i>BJU International</i> , 2004, 93, 1241-1245.	1.3	66
87	Genetic Polymorphisms of OPG, RANK, and ESR1 and Bone Mineral Density in Korean Postmenopausal Women. <i>Calcified Tissue International</i> , 2005, 77, 152-159.	1.5	65
88	Role of alcohol and genetic polymorphisms of CYP2E1 and ALDH2 in breast cancer development. <i>Pharmacogenetics and Genomics</i> , 2003, 13, 67-72.	5.7	64
89	Methylenetetrahydrofolate reductase polymorphism, diet, and breast cancer in Korean women. <i>Experimental and Molecular Medicine</i> , 2004, 36, 116-121.	3.2	64
90	Prevalence and epidemiologic characteristics of urolithiasis in Seoul, Korea. <i>Urology</i> , 2002, 59, 517-521.	0.5	63

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91	Serum High-Density Lipoprotein Cholesterol and Breast Cancer Risk by Menopausal Status, Body Mass Index, and Hormonal Receptor in Korea. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 508-515.	1.1	62
92	Intake of Soy Products and Other Foods and Gastric Cancer Risk: A Prospective Study. <i>Journal of Epidemiology</i> , 2013, 23, 337-343.	1.1	61
93	CYP2E1 and NQO1 genotypes, smoking and bladder cancer. <i>Pharmacogenetics and Genomics</i> , 2003, 13, 349-355.	5.7	60
94	ImmunoChip Analysis Identification of 6 Additional Susceptibility Loci for Crohn's Disease in Koreans. <i>Inflammatory Bowel Diseases</i> , 2015, 21, 1-7.	0.9	60
95	Expression of cyclooxygenase-1 and -2 associated with expression of VEGF in primary cervical cancer and at metastatic lymph nodes. <i>Gynecologic Oncology</i> , 2003, 90, 83-90.	0.6	59
96	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , 2016, 99, 903-911.	2.6	59
97	Expression of cyclooxygenase-2 in association with clinicopathological prognostic factors and molecular markers in epithelial ovarian cancer. <i>Gynecologic Oncology</i> , 2004, 92, 927-935.	0.6	58
98	Cancer incidence among pesticide applicators exposed to trifluralin in the Agricultural Health Study. <i>Environmental Research</i> , 2008, 107, 271-276.	3.7	58
99	Genome-wide association studies in East Asians identify new loci for waist-hip ratio and waist circumference. <i>Scientific Reports</i> , 2016, 6, 17958.	1.6	58
100	Association of Sleep Duration With All- and Major-Cause Mortality Among Adults in Japan, China, Singapore, and Korea. <i>JAMA Network Open</i> , 2021, 4, e2122837.	2.8	58
101	CagA-producing <i>Helicobacter pylori</i> and increased risk of gastric cancer: a nested case-control study in Korea. <i>British Journal of Cancer</i> , 2006, 95, 639-641.	2.9	57
102	Candidate gene approach evaluates association between innate immunity genes and breast cancer risk in Korean women. <i>Carcinogenesis</i> , 2009, 30, 1528-1531.	1.3	57
103	Five Polymorphisms and Breast Cancer Risk: Results from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1610-1616.	1.1	57
104	Postmenopausal obesity as a breast cancer risk factor according to estrogen and progesterone receptor status (Japan). <i>Cancer Letters</i> , 2001, 167, 57-63.	3.2	55
105	Egg Consumption and Risk of Metabolic Syndrome in Korean Adults: Results from the Health Examinees Study. <i>Nutrients</i> , 2017, 9, 687.	1.7	55
106	Korean Multi-center Cancer Cohort Study including a Biological Materials Bank (KMCC-I). <i>Asian Pacific Journal of Cancer Prevention</i> , 2002, 3, 85-92.	0.5	55
107	N-acetyltransferase (NAT1, NAT2) and glutathione S-transferase (GSTM1, GSTT1) polymorphisms in breast cancer. <i>Cancer Letters</i> , 2003, 196, 179-186.	3.2	54
108	CYP1A1, GSTM1, and GSTT1 Polymorphisms, Smoking, and Lung Cancer Risk in a Pooled Analysis among Asian Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 1120-1126.	1.1	54

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109	Common genetic polymorphisms of microRNA biogenesis pathway genes and breast cancer survival. <i>BMC Cancer</i> , 2012, 12, 195.	1.1	54
110	The Effect of Breastfeeding Duration and Parity on the Risk of Epithelial Ovarian Cancer: A Systematic Review and Meta-analysis. <i>Journal of Preventive Medicine and Public Health</i> , 2016, 49, 349-366.	0.7	54
111	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	1.4	53
112	What Are the Major Determinants in the Success of Smoking Cessation: Results from the Health Examinees Study. <i>PLoS ONE</i> , 2015, 10, e0143303.	1.1	53
113	Genetic Polymorphisms of SULT1A1 and SULT1E1 and the Risk and Survival of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1090-1095.	1.1	52
114	Polymorphism in the nuclear factor kappa-B binding promoter region of cyclooxygenase-2 is associated with an increased risk of bladder cancer. <i>Cancer Letters</i> , 2005, 217, 11-16.	3.2	52
115	Age and sex interactions in gastric cancer incidence and mortality trends in Korea. <i>Gastric Cancer</i> , 2015, 18, 580-589.	2.7	52
116	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. <i>Breast Cancer Research</i> , 2016, 18, 124.	2.2	52
117	High urine 1-hydroxypyrene glucuronide concentrations in Linxian, China, an area of high risk for squamous oesophageal cancer. <i>Biomarkers</i> , 2001, 6, 381-386.	0.9	51
118	Genetic polymorphisms of TGF- β 1 & TNF- β and breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2005, 90, 149-155.	1.1	51
119	A nested case-control study of the association of <i>Helicobacter pylori</i> infection with gastric adenocarcinoma in Korea. <i>British Journal of Cancer</i> , 2005, 92, 1273-1275.	2.9	51
120	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	2.3	51
121	Comparison of 6q25 Breast Cancer Hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC). <i>PLoS ONE</i> , 2012, 7, e42380.	1.1	51
122	hOGG1 Ser 326 Cys Polymorphism and Breast Cancer Risk among Asian Women. <i>Breast Cancer Research and Treatment</i> , 2003, 79, 59-62.	1.1	49
123	Combined effect of glutathione S-transferase M1 and T1 genotypes on bladder cancer risk. <i>Cancer Letters</i> , 2002, 177, 173-179.	3.2	48
124	Genetic Polymorphisms of Ataxia Telangiectasia Mutated and Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 821-825.	1.1	48
125	Associations between Breast Cancer Susceptibility Gene Polymorphisms and Clinicopathological Features. <i>Clinical Cancer Research</i> , 2004, 10, 124-130.	3.2	47
126	Dose-dependent protective effect of breast-feeding against breast cancer among ever-lactated women in Korea. <i>European Journal of Cancer Prevention</i> , 2007, 16, 124-129.	0.6	47

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127	Age at menarche and age at natural menopause in East Asian women: a genome-wide association study. <i>Age</i> , 2016, 38, 513-523.	3.0	47
128	Whole-Exome Sequencing Identifies Mutations of KIF22 in Spondyloepimetaphyseal Dysplasia with Joint Laxity, Leptodactylic Type. <i>American Journal of Human Genetics</i> , 2011, 89, 760-766.	2.6	46
129	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	5.8	46
130	Association of exposure to polycyclic aromatic hydrocarbons (estimated from job category) with concentration of 1-hydroxypyrene glucuronide in urine from workers at a steel plant.. <i>Occupational and Environmental Medicine</i> , 1995, 52, 593-599.	1.3	45
131	Polymorphisms in the estrogen receptor-alpha gene and breast cancer risk. <i>Cancer Letters</i> , 2002, 178, 175-180.	3.2	45
132	Soybean Product Intake Modifies the Association between Interleukin-10 Genetic Polymorphisms and Gastric Cancer Risk. <i>Journal of Nutrition</i> , 2009, 139, 1008-1012.	1.3	45
133	Common genetic polymorphisms of microRNA biogenesis pathway genes and risk of breast cancer: a case-control study in Korea. <i>Breast Cancer Research and Treatment</i> , 2011, 130, 939-951.	1.1	45
134	Combined COMT and GST genotypes and hormone replacement therapy associated breast cancer risk. <i>Pharmacogenetics and Genomics</i> , 2002, 12, 67-72.	5.7	44
135	A Haplotype Analysis of HER-2 Gene Polymorphisms: Association with Breast Cancer Risk, HER-2 Protein Expression in the Tumor, and Disease Recurrence in Korea. <i>Clinical Cancer Research</i> , 2005, 11, 4775-4778.	3.2	44
136	The role of TNF genetic variants and the interaction with cigarette smoking for gastric cancer risk: a nested case-control study. <i>BMC Cancer</i> , 2009, 9, 238.	1.1	44
137	Korean Risk Assessment Model for Breast Cancer Risk Prediction. <i>PLoS ONE</i> , 2013, 8, e76736.	1.1	44
138	Genetic polymorphisms of GSTM1, p21, p53 and HPV infection with cervical cancer in Korean women. <i>Gynecologic Oncology</i> , 2004, 93, 14-18.	0.6	43
139	Genetic polymorphisms of interleukin-1 beta (IL-1B) and IL-1 receptor antagonist (IL-1RN) and breast cancer risk in Korean women. <i>Breast Cancer Research and Treatment</i> , 2006, 96, 197-202.	1.1	43
140	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017, 19, 119.	2.2	43
141	Uteroglobin gene polymorphisms affect the progression of immunoglobulin A nephropathy by modulating the level of uteroglobin expression. <i>Pharmacogenetics and Genomics</i> , 2001, 11, 299-305.	5.7	42
142	Rare variant of hypoxia-inducible factor-1 β (HIF-1A) and breast cancer risk in Korean women. <i>Clinica Chimica Acta</i> , 2008, 389, 167-170.	0.5	42
143	Breast cancer prevention based on gene-environment interaction. <i>Molecular Carcinogenesis</i> , 2011, 50, 280-290.	1.3	42
144	Hormone-related factors and post-menopausal onset depression: Results from KNHANES (2010-2012). <i>Journal of Affective Disorders</i> , 2015, 175, 176-183.	2.0	42

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145	Associations of Body Mass Index, Smoking, and Alcohol Consumption With Prostate Cancer Mortality in the Asia Cohort Consortium. <i>American Journal of Epidemiology</i> , 2015, 182, 381-389.	1.6	42
146	A longitudinal study of atrazine and 2,4-DE exposure and oxidative stress markers among iowa corn farmers. <i>Environmental and Molecular Mutagenesis</i> , 2017, 58, 30-38.	0.9	42
147	One-carbon metabolism gene polymorphisms and risk of non-Hodgkin lymphoma in Australia. <i>Human Genetics</i> , 2007, 122, 525-533.	1.8	41
148	Breast Cancer and Urinary Biomarkers of Polycyclic Aromatic Hydrocarbon and Oxidative Stress in the Shanghai Women's Health Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 877-883.	1.1	41
149	Menarche age, menopause age and other reproductive factors in association with post-menopausal onset depression: Results from Health Examinees Study (HEXA). <i>Journal of Affective Disorders</i> , 2015, 187, 127-135.	2.0	41
150	Variations in urinary 1-hydroxypyrene glucuronide in relation to smoking and the modification effects of GSTM1 and GSTT1. <i>Toxicology Letters</i> , 1999, 108, 217-223.	0.4	40
151	Association of serum lipids and glucose with the risk of colorectal adenomatous polyp in men: a case-control study in Korea. <i>Journal of Korean Medical Science</i> , 2000, 15, 690.	1.1	40
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