

Loïc Le Marchand

List of Publications by Year in descending order

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Version: 2024-02-01

739
papers

62,230
citations

1713

107
h-index

2108

210
g-index

748
all docs

748
docs citations

748
times ranked

65998
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	13.7	3,823
2	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	13.7	2,165
3	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	9.4	1,818
4	Risk of COVID-19 among front-line health-care workers and the general community: a prospective cohort study. <i>Lancet Public Health</i> , The, 2020, 5, e475-e483.	4.7	1,595
5	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	13.7	1,328
6	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
7	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
8	Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptor-positive breast cancer. <i>Nature Genetics</i> , 2007, 39, 865-869.	9.4	774
9	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
10	Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , 2019, 570, 514-518.	13.7	679
11	Ethnic and Racial Differences in the Smoking-Related Risk of Lung Cancer. <i>New England Journal of Medicine</i> , 2006, 354, 333-342.	13.9	668
12	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	9.4	652
13	Multiple regions within 8q24 independently affect risk for prostate cancer. <i>Nature Genetics</i> , 2007, 39, 638-644.	9.4	621
14	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	3.0	596
15	<i>CCAT2</i> , a novel noncoding RNA mapping to 8q24, underlies metastatic progression and chromosomal instability in colon cancer. <i>Genome Research</i> , 2013, 23, 1446-1461.	2.4	526
16	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	9.4	519
17	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
18	Detectable clonal mosaicism from birth to old age and its relationship to cancer. <i>Nature Genetics</i> , 2012, 44, 642-650.	9.4	511

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19	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
20	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	9.4	492
21	Cancer preventive effects of flavonoids—a review. <i>Biomedicine and Pharmacotherapy</i> , 2002, 56, 296-301.	2.5	476
22	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	9.4	472
23	Identification of Lynch Syndrome Among Patients With Colorectal Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2012, 308, 1555.	3.8	443
24	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	13.9	414
25	Common variants on chromosome 5p12 confer susceptibility to estrogen receptor-positive breast cancer. <i>Nature Genetics</i> , 2008, 40, 703-706.	9.4	412
26	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014, 46, 1103-1109.	9.4	408
27	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. <i>Nature Genetics</i> , 2009, 41, 1116-1121.	9.4	389
28	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	9.4	377
29	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
30	Meta-analysis identifies common variants associated with body mass index in east Asians. <i>Nature Genetics</i> , 2012, 44, 307-311.	9.4	372
31	Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants: findings from the Prospective Lynch Syndrome Database. <i>Genetics in Medicine</i> , 2020, 22, 15-25.	1.1	365
32	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
33	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 404-412.	1.1	341
34	A common genetic risk factor for colorectal and prostate cancer. <i>Nature Genetics</i> , 2007, 39, 954-956.	9.4	336
35	Risks of Lynch Syndrome Cancers for MSH6 Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2010, 102, 193-201.	3.0	328
36	The landscape of recombination in African Americans. <i>Nature</i> , 2011, 476, 170-175.	13.7	319

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37	Colon Cancer Family Registry: An International Resource for Studies of the Genetic Epidemiology of Colon Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2331-2343.	1.1	315
38	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	315
39	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000.	9.4	294
40	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
41	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
42	Colorectal and Other Cancer Risks for Carriers and Noncarriers From Families With a DNA Mismatch Repair Gene Mutation: A Prospective Cohort Study. <i>Journal of Clinical Oncology</i> , 2012, 30, 958-964.	0.8	286
43	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. <i>JAMA Oncology</i> , 2016, 2, 1295.	3.4	285
44	Associations of key diet-quality indexes with mortality in the Multiethnic Cohort: the Dietary Patterns Methods Project. <i>American Journal of Clinical Nutrition</i> , 2015, 101, 587-597.	2.2	280
45	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptorâ€“negative breast cancer. <i>Nature Genetics</i> , 2011, 43, 1210-1214.	9.4	279
46	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	9.4	265
47	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
48	Metabolism and Biomarkers of Heterocyclic Aromatic Amines in Molecular Epidemiology Studies: Lessons Learned from Aromatic Amines. <i>Chemical Research in Toxicology</i> , 2011, 24, 1169-1214.	1.7	252
49	Meta- and pooled analyses of the effects of glutathione S-transferase M1 polymorphisms and smoking on lung cancer risk. <i>Carcinogenesis</i> , 2002, 23, 1343-1350.	1.3	250
50	Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. <i>Hepatology</i> , 2016, 64, 1969-1977.	3.6	237
51	Generalization and Dilution of Association Results from European GWAS in Populations of Non-European Ancestry: The PAGE Study. <i>PLoS Biology</i> , 2013, 11, e1001661.	2.6	235
52	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , 2013, 45, 690-696.	9.4	232
53	Association of a Low-Frequency Variant in <i>HNF1A</i> With Type 2 Diabetes in a Latino Population. <i>JAMA - Journal of the American Medical Association</i> , 2014, 311, 2305.	3.8	230
54	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers: the advantage of more extensive colon surgery. <i>Gut</i> , 2011, 60, 950-957.	6.1	227

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55	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. <i>Gastroenterology</i> , 2018, 154, 2152-2164.e19.	0.6	226
56	BODY SIZE AT DIFFERENT PERIODS OF LIFE AND BREAST CANCER RISK. <i>American Journal of Epidemiology</i> , 1988, 128, 137-152.	1.6	222
57	Genome-wide association study of glioma and meta-analysis. <i>Human Genetics</i> , 2012, 131, 1877-1888.	1.8	222
58	Sun exposure and melanoma risk at different latitudes: a pooled analysis of 5700 cases and 7216 controls. <i>International Journal of Epidemiology</i> , 2009, 38, 814-830.	0.9	219
59	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	9.4	218
60	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
61	Cancer Risks for <i>MLH1</i> and <i>MSH2</i> Mutation Carriers. <i>Human Mutation</i> , 2013, 34, 490-497.	1.1	201
62	Circulating Vitamin D and Colorectal Cancer Risk: An International Pooling Project of 17 Cohorts. <i>Journal of the National Cancer Institute</i> , 2019, 111, 158-169.	3.0	199
63	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011, 43, 570-573.	9.4	198
64	Risks of Primary Extracolonic Cancers Following Colorectal Cancer in Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1363-1372.	3.0	193
65	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	5.8	188
66	Smokers with the <i>CHRNA</i> Lung Cancer-Associated Variants Are Exposed to Higher Levels of Nicotine Equivalents and a Carcinogenic Tobacco-Specific Nitrosamine. <i>Cancer Research</i> , 2008, 68, 9137-9140.	0.4	186
67	<i>CYP1A1</i> and <i>GSTM1</i> genetic polymorphisms and lung cancer risk in Caucasian non-smokers: a pooled analysis. <i>Carcinogenesis</i> , 2003, 24, 875-882.	1.3	184
68	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	9.4	184
69	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , 2012, 131, 217-234.	1.8	183
70	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
71	Association of Coffee Consumption With Total and Cause-Specific Mortality Among Nonwhite Populations. <i>Annals of Internal Medicine</i> , 2017, 167, 228.	2.0	182
72	Circulating 25-Hydroxyvitamin D and Risk of Pancreatic Cancer: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. <i>American Journal of Epidemiology</i> , 2010, 172, 81-93.	1.6	181

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73	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
74	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
75	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	1.5	174
76	Phenome-Wide Association Study (PheWAS) for Detection of Pleiotropy within the Population Architecture using Genomics and Epidemiology (PAGE) Network. <i>PLoS Genetics</i> , 2013, 9, e1003087.	1.5	171
77	Association of Aspirin and NSAID Use With Risk of Colorectal Cancer According to Genetic Variants. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1133.	3.8	171
78	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017, 8, 14977.	5.8	169
79	Consistent Association of Type 2 Diabetes Risk Variants Found in Europeans in Diverse Racial and Ethnic Groups. <i>PLoS Genetics</i> , 2010, 6, e1001078.	1.5	168
80	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	1.4	168
81	Case-Control Study of Overweight, Obesity, and Colorectal Cancer Risk, Overall and by Tumor Microsatellite Instability Status. <i>Journal of the National Cancer Institute</i> , 2010, 102, 391-400.	3.0	162
82	B-vitamin intake, metabolic genes, and colorectal cancer risk (United States). <i>Cancer Causes and Control</i> , 2002, 13, 239-248.	0.8	161
83	The Next PAGE in Understanding Complex Traits: Design for the Analysis of Population Architecture Using Genetics and Epidemiology (PAGE) Study. <i>American Journal of Epidemiology</i> , 2011, 174, 849-859.	1.6	161
84	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011, 20, 3867-3875.	1.4	160
85	Previous Lung Diseases and Lung Cancer Risk: A Pooled Analysis From the International Lung Cancer Consortium. <i>American Journal of Epidemiology</i> , 2012, 176, 573-585.	1.6	160
86	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017, 13, e1006528.	1.5	158
87	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152
88	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
89	Interactions Between Genetic Variants and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. <i>Journal of the National Cancer Institute</i> , 2011, 103, 1252-1263.	3.0	147
90	Cancer Risks for <i>PMS2</i> -Associated Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2018, 36, 2961-2968.	0.8	147

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91	Genetic Determinants of Lipid Traits in Diverse Populations from the Population Architecture using Genomics and Epidemiology (PAGE) Study. <i>PLoS Genetics</i> , 2011, 7, e1002138.	1.5	146
92	Association of Coffee Intake With Reduced Incidence of Liver Cancer and Death From Chronic Liver Disease in the US Multiethnic Cohort. <i>Gastroenterology</i> , 2015, 148, 118-125.	0.6	145
93	CYP1A1 T3801 C polymorphism and lung cancer: A pooled analysis of 2,451 cases and 3,358 controls. <i>International Journal of Cancer</i> , 2003, 104, 650-657.	2.3	140
94	Genetic Variation at the CYP19A1 Locus Predicts Circulating Estrogen Levels but not Breast Cancer Risk in Postmenopausal Women. <i>Cancer Research</i> , 2007, 67, 1893-1897.	0.4	140
95	The contribution of rare variation to prostate cancer heritability. <i>Nature Genetics</i> , 2016, 48, 30-35.	9.4	139
96	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	5.8	138
97	Genome-Wide Association Study in East Asians Identifies Novel Susceptibility Loci for Breast Cancer. <i>PLoS Genetics</i> , 2012, 8, e1002532.	1.5	137
98	A Common 8q24 Variant in Prostate and Breast Cancer from a Large Nested Case-Control Study. <i>Cancer Research</i> , 2007, 67, 2951-2956.	0.4	136
99	Testing the Predictive Validity of the Healthy Eating Index-2015 in the Multiethnic Cohort: Is the Score Associated with a Reduced Risk of All-Cause and Cause-Specific Mortality?. <i>Nutrients</i> , 2018, 10, 452.	1.7	132
100	Quality Assessment and Correlation of Microsatellite Instability and Immunohistochemical Markers among Population- and Clinic-Based Colorectal Tumors. <i>Journal of Molecular Diagnostics</i> , 2011, 13, 271-281.	1.2	131
101	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	3.0	129
102	Genetic determinants of telomere length and risk of common cancers: a Mendelian randomization study. <i>Human Molecular Genetics</i> , 2015, 24, 5356-5366.	1.4	128
103	Exposure of nonsmoking women to environmental tobacco smoke: a 10-country collaborative study. <i>Cancer Causes and Control</i> , 1990, 1, 243-252.	0.8	127
104	Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. <i>Carcinogenesis</i> , 2014, 35, 2526-2533.	1.3	124
105	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444.	2.6	124
106	A Review of the Application of Inflammatory Biomarkers in Epidemiologic Cancer Research. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1729-1751.	1.1	123
107	Obesity in youth and middle age and risk of colorectal cancer in men. <i>Cancer Causes and Control</i> , 1992, 3, 349-354.	0.8	119
108	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. <i>Human Molecular Genetics</i> , 2013, 22, 408-415.	1.4	118

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109	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	3.9	118
110	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. <i>PLoS Genetics</i> , 2011, 7, e1001387.	1.5	117
111	A Meta-analysis of Individual Participant Data Reveals an Association between Circulating Levels of IGF-I and Prostate Cancer Risk. <i>Cancer Research</i> , 2016, 76, 2288-2300.	0.4	117
112	Vegetable and Fruit Consumption in Relation to Prostate Cancer Risk in Hawaii: A Reevaluation of the Effect of Dietary Beta-Carotene. <i>American Journal of Epidemiology</i> , 1991, 133, 215-219.	1.6	115
113	Correlates of Circulating 25-Hydroxyvitamin D: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. <i>American Journal of Epidemiology</i> , 2010, 172, 21-35.	1.6	114
114	Estimating the heritability of colorectal cancer. <i>Human Molecular Genetics</i> , 2014, 23, 3898-3905.	1.4	114
115	Trans-Ethnic Fine-Mapping of Lipid Loci Identifies Population-Specific Signals and Allelic Heterogeneity That Increases the Trait Variance Explained. <i>PLoS Genetics</i> , 2013, 9, e1003379.	1.5	112
116	Prediagnostic Leptin, Adiponectin, C-Reactive Protein, and the Risk of Postmenopausal Breast Cancer. <i>Cancer Prevention Research</i> , 2013, 6, 188-195.	0.7	112
117	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
118	Cumulative Burden of Colorectal Cancer-associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12.	0.6	110
119	Associations between Smoking, Alcohol Consumption, and Colorectal Cancer, Overall and by Tumor Microsatellite Instability Status. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2745-2750.	1.1	109
120	Common Genetic Variation in IGF1 and Prostate Cancer Risk in the Multiethnic Cohort. <i>Journal of the National Cancer Institute</i> , 2006, 98, 123-134.	3.0	107
121	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
122	Carotenoids, retinol, tocopherols, and prostate cancer risk: pooled analysis of 15 studies. <i>American Journal of Clinical Nutrition</i> , 2015, 102, 1142-1157.	2.2	107
123	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in <i>MUTYH</i> . <i>International Journal of Cancer</i> , 2016, 139, 1557-1563.	2.3	107
124	IGF-1, IGFBP-1, and IGFBP-3 Polymorphisms Predict Circulating IGF Levels but Not Breast Cancer Risk: Findings from the Breast and Prostate Cancer Cohort Consortium (BPC3). <i>PLoS ONE</i> , 2008, 3, e2578.	1.1	106
125	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
126	Risk of Metachronous Colon Cancer Following Surgery for Rectal Cancer in Mismatch Repair Gene Mutation Carriers. <i>Annals of Surgical Oncology</i> , 2013, 20, 1829-1836.	0.7	103

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127	Generalizability of Associations from Prostate Cancer Genome-Wide Association Studies in Multiple Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1285-1289.	1.1	102
128	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	1.1	101
129	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
130	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. <i>Human Molecular Genetics</i> , 2013, 22, 2520-2528.	1.4	100
131	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
132	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
133	Pooled analysis of the CYP1A1 exon 7 polymorphism and lung cancer (United States). <i>Cancer Causes and Control</i> , 2003, 14, 339-346.	0.8	98
134	Common variants at 11q12, 10q26 and 3p11.2 are associated with prostate cancer susceptibility in Japanese. <i>Nature Genetics</i> , 2012, 44, 426-429.	9.4	98
135	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
136	High-Quality Diets Associate With Reduced Risk of Colorectal Cancer: Analyses of Diet Quality Indexes in the Multiethnic Cohort. <i>Gastroenterology</i> , 2017, 153, 386-394.e2.	0.6	98
137	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	0.6	97
138	Plasma 25-Hydroxyvitamin D Levels and the Risk of Colorectal Cancer: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 130-134.	1.1	96
139	Associations of plasma trimethylamine N-oxide, choline, carnitine, and betaine with inflammatory and cardiometabolic risk biomarkers and the fecal microbiome in the Multiethnic Cohort Adiposity Phenotype Study. <i>American Journal of Clinical Nutrition</i> , 2020, 111, 1226-1234.	2.2	96
140	A priori-defined diet quality indexes and risk of type 2 diabetes: the Multiethnic Cohort. <i>Diabetologia</i> , 2015, 58, 98-112.	2.9	94
141	Dietary fiber and colorectal cancer risk: the multiethnic cohort study. <i>Cancer Causes and Control</i> , 2007, 18, 753-764.	0.8	93
142	Cancer risks for monoallelic <i>MUTYH</i> mutation carriers with a family history of colorectal cancer. <i>International Journal of Cancer</i> , 2011, 129, 2256-2262.	2.3	93
143	Identification, Replication, and Fine-Mapping of Loci Associated with Adult Height in Individuals of African Ancestry. <i>PLoS Genetics</i> , 2011, 7, e1002298.	1.5	93
144	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

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145	Lynch Syndromeâ€Associated Breast Cancers: Clinicopathologic Characteristics of a Case Series from the Colon Cancer Family Registry. <i>Clinical Cancer Research</i> , 2010, 16, 2214-2224.	3.2	91
146	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
147	Meta-analysis of 16 studies of the association of alcohol with colorectal cancer. <i>International Journal of Cancer</i> , 2020, 146, 861-873.	2.3	89
148	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , 2015, 6, 6889.	5.8	88
149	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016, 7, 66328-66343.	0.8	88
150	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	5.8	88
151	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
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739	Neighborhood Obesogenic Environment and Risk of Prostate Cancer: The Multiethnic Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 972-981.	1.1	0