

Ulrike Peters

List of Publications by Year in descending order

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Version: 2024-02-01

255
papers

25,731
citations

16451

64
h-index

9345

143
g-index

268
all docs

268
docs citations

268
times ranked

35973
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.	27.8	3,823
2	A reference panel of 64,976 haplotypes for genotype imputation. <i>Nature Genetics</i> , 2016, 48, 1279-1283.	21.4	2,421
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.	21.4	1,818
4	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	27.8	1,328
5	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. <i>Nature Genetics</i> , 2019, 51, 237-244.	21.4	1,307
6	Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , 2019, 570, 514-518.	27.8	679
7	Estimation of effect size distribution from genome-wide association studies and implications for future discoveries. <i>Nature Genetics</i> , 2010, 42, 570-575.	21.4	609
8	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	21.4	377
9	Genetic Mechanisms of Immune Evasion in Colorectal Cancer. <i>Cancer Discovery</i> , 2018, 8, 730-749.	9.4	367
10	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	21.4	357
11	Dietary fibre and colorectal adenoma in a colorectal cancer early detection programme. <i>Lancet</i> , The, 2003, 361, 1491-1495.	13.7	302
12	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000.	21.4	294
13	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , 2013, 144, 799-807.e24.	1.3	292
14	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. <i>American Journal of Human Genetics</i> , 2014, 94, 223-232.	6.2	287
15	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	2.5	278
16	Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms. <i>Nature Genetics</i> , 2017, 49, 1113-1119.	21.4	260
17	Causes of Death in Patients With Celiac Disease in a Population-Based Swedish Cohort. <i>Archives of Internal Medicine</i> , 2003, 163, 1566.	3.8	239
18	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.	5.6	235

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19	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. <i>Molecular Psychiatry</i> , 2015, 20, 647-656.	7.9	235
20	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.	1.3	226
21	Supplemental and Dietary Vitamin E, β -Carotene, and Vitamin C Intakes and Prostate Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2006, 98, 245-254.	6.3	206
22	Whole-Exome Sequencing Identifies Rare and Low-Frequency Coding Variants Associated with LDL Cholesterol. <i>American Journal of Human Genetics</i> , 2014, 94, 233-245.	6.2	193
23	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. <i>Nature Communications</i> , 2020, 11, 597.	12.8	193
24	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	12.8	188
25	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , 2012, 131, 217-234.	3.8	183
26	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
27	Meta-analysis of gene-level tests for rare variant association. <i>Nature Genetics</i> , 2014, 46, 200-204.	21.4	178
28	Genome-wide association analyses in east Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013, 45, 191-196.	21.4	173
29	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.	7.4	171
30	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.	12.8	169
31	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.	3.4	161
32	Genome-wide physical activity interactions in adiposity $\hat{\epsilon}$. A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017, 13, e1006528.	3.5	158
33	A Prospective Study of Lycopene and Tomato Product Intake and Risk of Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 92-98.	2.5	153
34	Genetic architecture of colorectal cancer. <i>Gut</i> , 2015, 64, 1623-1636.	12.1	152
35	Serum Lycopene, Other Carotenoids, and Prostate Cancer Risk: a Nested Case-Control Study in the Prostate, Lung, Colorectal, and Ovarian Cancer Screening Trial. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 962-968.	2.5	147
36	Characterization of Gene-Environment Interactions for Colorectal Cancer Susceptibility Loci. <i>Cancer Research</i> , 2012, 72, 2036-2044.	0.9	140

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37	Integrative analysis of exogenous, endogenous, tumour and immune factors for precision medicine. <i>Gut</i> , 2018, 67, 1168-1180.	12.1	139
38	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	12.8	138
39	A Model to Determine Colorectal Cancer Risk Using Common Genetic Susceptibility Loci. <i>Gastroenterology</i> , 2015, 148, 1330-1339.e14.	1.3	129
40	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	6.3	129
41	Methods for etiologic and early marker investigations in the PLCO trial. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2005, 592, 147-154.	1.0	125
42	Mendelian randomization study of adiposity-related traits and risk of breast, ovarian, prostate, lung and colorectal cancer. <i>International Journal of Epidemiology</i> , 2016, 45, 896-908.	1.9	124
43	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444.	6.2	124
44	Estimating the heritability of colorectal cancer. <i>Human Molecular Genetics</i> , 2014, 23, 3898-3905.	2.9	114
45	Rare and low-frequency coding variants in CXCR2 and other genes are associated with hematological traits. <i>Nature Genetics</i> , 2014, 46, 629-634.	21.4	113
46	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.	1.3	110
47	Genome-wide Association and Population Genetic Analysis of C-Reactive Protein in African American and Hispanic American Women. <i>American Journal of Human Genetics</i> , 2012, 91, 502-512.	6.2	107
48	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	12.8	106
49	Variation in the Selenoenzyme Genes and Risk of Advanced Distal Colorectal Adenoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 1144-1154.	2.5	101
50	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
51	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.	0.9	100
52	Selenium and the prevention of prostate and colorectal cancer. <i>Molecular Nutrition and Food Research</i> , 2008, 52, 1261-1272.	3.3	93
53	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.	2.9	90
54	Circulating Levels of Insulin-like Growth Factor 1 and Insulin-like Growth Factor Binding Protein 3 Associate With Risk of Colorectal Cancer Based on Serologic and Mendelian Randomization Analyses. <i>Gastroenterology</i> , 2020, 158, 1300-1312.e20.	1.3	90

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55	Meta-analysis of 16 studies of the association of alcohol with colorectal cancer. <i>International Journal of Cancer</i> , 2020, 146, 861-873.	5.1	89
56	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016, 7, 66328-66343.	1.8	88
57	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
58	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	12.8	86
59	Vitamin E and selenium supplementation and risk of prostate cancer in the Vitamins and lifestyle (VITAL) study cohort. <i>Cancer Causes and Control</i> , 2008, 19, 75-87.	1.8	85
60	Genome-Wide Diet-Gene Interaction Analyses for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , 2014, 10, e1004228.	3.5	81
61	Fine Mapping and Identification of BMI Loci in African Americans. <i>American Journal of Human Genetics</i> , 2013, 93, 661-671.	6.2	77
62	Serum selenium and risk of prostate cancer—a nested case-control study. <i>American Journal of Clinical Nutrition</i> , 2007, 85, 209-217.	4.7	76
63	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. <i>BMC Medicine</i> , 2020, 18, 396.	5.5	76
64	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353.	12.8	75
65	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VT11A. <i>Nature Communications</i> , 2014, 5, 4613.	12.8	72
66	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. <i>PLoS ONE</i> , 2016, 11, e0167758.	2.5	72
67	Evaluation of the MetaboChip Genotyping Array in African Americans and Implications for Fine Mapping of GWAS-Identified Loci: The PAGE Study. <i>PLoS ONE</i> , 2012, 7, e35651.	2.5	71
68	Association between Adult Height and Risk of Colorectal, Lung, and Prostate Cancer: Results from Meta-analyses of Prospective Studies and Mendelian Randomization Analyses. <i>PLoS Medicine</i> , 2016, 13, e1002118.	8.4	69
69	Genetic risk factors for BMI and obesity in an ethnically diverse population: Results from the population architecture using genomics and epidemiology (PAGE) study. <i>Obesity</i> , 2013, 21, 835-846.	3.0	68
70	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1024-1031.	2.5	67
71	Genetic identification of a common collagen disease in Puerto Ricans via identity-by-descent mapping in a health system. <i>ELife</i> , 2017, 6, .	6.0	65
72	A Systematic Mapping Approach of 16q12.2/FTO and BMI in More Than 20,000 African Americans Narrows in on the Underlying Functional Variation: Results from the Population Architecture using Genomics and Epidemiology (PAGE) Study. <i>PLoS Genetics</i> , 2013, 9, e1003171.	3.5	63

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73	Cross Cancer Genomic Investigation of Inflammation Pathway for Five Common Cancers: Lung, Ovary, Prostate, Breast, and Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv246.	6.3	63
74	Common genetic variation and survival after colorectal cancer diagnosis: a genome-wide analysis. <i>Carcinogenesis</i> , 2016, 37, 87-95.	2.8	62
75	Polymorphism analysis of six selenoprotein genes: support for a selective sweep at the glutathione peroxidase 1 locus (3p21) in Asian populations. <i>BMC Genetics</i> , 2006, 7, 56.	2.7	59
76	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1003-1012.	6.3	59
77	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. <i>Nature Communications</i> , 2020, 11, 2542.	12.8	59
78	High Serum Selenium and Reduced Risk of Advanced Colorectal Adenoma in a Colorectal Cancer Early Detection Program. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 315-320.	2.5	58
79	Genotype Imputation of M -etabochip SNPs Using a Study-Specific Reference Panel of $\sim 44,000$ Haplotypes in African Americans From the Women's Health Initiative. <i>Genetic Epidemiology</i> , 2012, 36, 107-117.	1.3	57
80	Mendelian randomization study of height and risk of colorectal cancer. <i>International Journal of Epidemiology</i> , 2015, 44, 662-672.	1.9	55
81	Landscape of somatic single nucleotide variants and indels in colorectal cancer and impact on survival. <i>Nature Communications</i> , 2020, 11, 3644.	12.8	55
82	Characterization of the association between 8q24 and colon cancer: gene-environment exploration and meta-analysis. <i>BMC Cancer</i> , 2010, 10, 670.	2.6	54
83	A Pooled Analysis of Smoking and Colorectal Cancer: Timing of Exposure and Interactions with Environmental Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1974-1985.	2.5	54
84	Sex-specific gene and pathway modeling of inherited glioma risk. <i>Neuro-Oncology</i> , 2019, 21, 71-82.	1.2	52
85	Calcium intake and colorectal adenoma in a US colorectal cancer early detection program. <i>American Journal of Clinical Nutrition</i> , 2004, 80, 1358-1365.	4.7	51
86	Consistent Directions of Effect for Established Type 2 Diabetes Risk Variants Across Populations. <i>Diabetes</i> , 2012, 61, 1642-1647.	0.6	49
87	Rare and Coding Region Genetic Variants Associated With Risk of Ischemic Stroke. <i>JAMA Neurology</i> , 2015, 72, 781.	9.0	49
88	Gene-Environment Interaction Involving Recently Identified Colorectal Cancer Susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1824-1833.	2.5	48
89	Genetic variation in C-reactive protein in relation to colon and rectal cancer risk and survival. <i>International Journal of Cancer</i> , 2011, 128, 2726-2734.	5.1	47
90	Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. <i>Human Genetics</i> , 2019, 138, 307-326.	3.8	44

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91	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334.	12.1	44
92	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , 2016, 139, 2655-2670.	5.1	43
93	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. <i>International Journal of Cancer</i> , 2020, 146, 363-372.	5.1	40
94	Association Between Colorectal Cancer Susceptibility Loci and Survival Time After Diagnosis With Colorectal Cancer. <i>Gastroenterology</i> , 2012, 143, 51-54.e4.	1.3	39
95	SBERIA: Set-Based Gene-Environment Interaction Test for Rare and Common Variants in Complex Diseases. <i>Genetic Epidemiology</i> , 2013, 37, 452-464.	1.3	39
96	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab029.	2.9	39
97	Urinary mutagenesis and fried red meat intake: Influence of cooking temperature, phenotype, and genotype of metabolizing enzymes in a controlled feeding study. <i>Environmental and Molecular Mutagenesis</i> , 2004, 43, 53-74.	2.2	38
98	Red Meat Intake, NAT2, and Risk of Colorectal Cancer: A Pooled Analysis of 11 Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 198-205.	2.5	38
99	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , 2016, 12, e1006296.	3.5	38
100	Association of exome sequences with plasma C-reactive protein levels in >9000 participants. <i>Human Molecular Genetics</i> , 2015, 24, 559-571.	2.9	36
101	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , 2021, 160, 1164-1178.e6.	1.3	36
102	Association of <i>Fusobacterium nucleatum</i> with Specific T-cell Subsets in the Colorectal Carcinoma Microenvironment. <i>Clinical Cancer Research</i> , 2021, 27, 2816-2826.	7.0	36
103	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , 2014, 63, 800-807.	12.1	35
104	Pleiotropic Associations of Risk Variants Identified for Other Cancers With Lung Cancer Risk: The PAGE and TRICL Consortia. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju061.	6.3	35
105	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 767-780.	1.9	35
106	The Risk of Ovarian Cancer Increases with an Increase in the Lifetime Number of Ovulatory Cycles: An Analysis from the Ovarian Cancer Cohort Consortium (OC3). <i>Cancer Research</i> , 2020, 80, 1210-1218.	0.9	35
107	Genome-Wide Search for Gene-Gene Interactions in Colorectal Cancer. <i>PLoS ONE</i> , 2012, 7, e52535.	2.5	35
108	Associations Between Glycemic Traits and Colorectal Cancer: A Mendelian Randomization Analysis. <i>Journal of the National Cancer Institute</i> , 2022, 114, 740-752.	6.3	35

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109	Circulating vitamin D metabolites, polymorphism in vitamin D receptor, and colorectal adenoma risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 546-52.	2.5	35
110	HNF1B and Endometrial Cancer Risk: Results from the PAGE study. <i>PLoS ONE</i> , 2012, 7, e30390.	2.5	34
111	Serum α -Tocopherol and β -Tocopherol Concentrations and Prostate Cancer Risk in the PLCO Screening Trial: A Nested Case-Control Study. <i>PLoS ONE</i> , 2012, 7, e40204.	2.5	34
112	Association Between Molecular Subtypes of Colorectal Tumors and Patient Survival, Based on Pooled Analysis of 7 International Studies. <i>Gastroenterology</i> , 2020, 158, 2158-2168.e4.	1.3	34
113	Serum Retinol and Prostate Cancer Risk: a Nested Case-Control Study in the Prostate, Lung, Colorectal, and Ovarian Cancer Screening Trial. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1227-1231.	2.5	33
114	Serum Selenium, Genetic Variation in Selenoenzymes, and Risk of Colorectal Cancer: Primary Analysis from the Women's Health Initiative Observational Study and Meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1822-1830.	2.5	33
115	The Future of Genomic Studies Must Be Globally Representative: Perspectives from PAGE. <i>Annual Review of Genomics and Human Genetics</i> , 2019, 20, 181-200.	6.2	33
116	Development and application of a simple routine method for the determination of selenium in serum by octopole reaction system ICPMS. <i>Analytical and Bioanalytical Chemistry</i> , 2005, 381, 686-694.	3.7	32
117	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. <i>Human Genetics</i> , 2017, 136, 771-800.	3.8	31
118	Anthropometric Risk Factors for Cancers of the Biliary Tract in the Biliary Tract Cancers Pooling Project. <i>Cancer Research</i> , 2019, 79, 3973-3982.	0.9	31
119	Genetic Variation in Calcium-Sensing Receptor and Risk for Colon Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2755-2765.	2.5	30
120	Characterization of 9p24 Risk Locus and Colorectal Adenoma and Cancer: Gene-Environment Interaction and Meta-Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 3131-3139.	2.5	30
121	Genetic Predictors of Circulating 25-Hydroxyvitamin D and Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 2037-2046.	2.5	30
122	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. <i>American Journal of Human Genetics</i> , 2018, 102, 904-919.	6.2	30
123	The Influence of Obesity-Related Single Nucleotide Polymorphisms on BMI Across the Life Course: The PAGE Study. <i>Diabetes</i> , 2013, 62, 1763-1767.	0.6	29
124	Fine-mapping of lipid regions in global populations discovers ethnic-specific signals and refines previously identified lipid loci. <i>Human Molecular Genetics</i> , 2016, 25, 5500-5512.	2.9	29
125	Circulating adipokine concentrations and risk of five obesity-related cancers: A Mendelian randomization study. <i>International Journal of Cancer</i> , 2021, 148, 1625-1636.	5.1	29
126	Association of serum phospholipid fatty acids with breast cancer risk among postmenopausal cigarette smokers. <i>Cancer Causes and Control</i> , 2009, 20, 497-504.	1.8	28

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127	Selenium, Selenoenzymes, Oxidative Stress and Risk of Neoplastic Progression from Barrett's Esophagus: Results from Biomarkers and Genetic Variants. PLoS ONE, 2012, 7, e38612.	2.5	28
128	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. Human Genetics, 2015, 134, 1249-1262.	3.8	28
129	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. Carcinogenesis, 2015, 36, 999-1007.	2.8	28
130	Combined effect of modifiable and non-modifiable risk factors for colorectal cancer risk in a pooled analysis of 11 population-based studies. BMJ Open Gastroenterology, 2019, 6, e000339.	2.7	28
131	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. BMC Medicine, 2020, 18, 229.	5.5	28
132	The use of imputed values in the meta-analysis of genome-wide association studies. Genetic Epidemiology, 2011, 35, 597-605.	1.3	27
133	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. American Journal of Clinical Nutrition, 2021, 113, 1490-1502.	4.7	27
134	Common Single-Nucleotide Polymorphisms in the Estrogen Receptor β Promoter Are Associated with Colorectal Cancer Survival in Postmenopausal Women. Cancer Research, 2013, 73, 767-775.	0.9	26
135	Variation in selenoenzyme genes and prostate cancer risk and survival. Prostate, 2013, 73, 734-742.	2.3	26
136	Genetic Variation Predicts Serum Lycopene Concentrations in a Multiethnic Population of Postmenopausal Women. Journal of Nutrition, 2015, 145, 187-192.	2.9	26
137	Relationship of prediagnostic body mass index with survival after colorectal cancer: Stage-specific associations. International Journal of Cancer, 2016, 139, 1065-1072.	5.1	26
138	Intake of Dietary Fruit, Vegetables, and Fiber and Risk of Colorectal Cancer According to Molecular Subtypes: A Pooled Analysis of 9 Studies. Cancer Research, 2020, 80, 4578-4590.	0.9	26
139	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	2.5	26
140	Influence of Smoking, Body Mass Index, and Other Factors on the Preventive Effect of Nonsteroidal Anti-Inflammatory Drugs on Colorectal Cancer Risk. Cancer Research, 2018, 78, 4790-4799.	0.9	26
141	The genetic underpinnings of variation in ages at menarche and natural menopause among women from the multi-ethnic Population Architecture using Genomics and Epidemiology (PAGE) Study: A trans-ethnic meta-analysis. PLoS ONE, 2018, 13, e0200486.	2.5	25
142	Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 477-486.	2.5	25
143	Genetic variants associated with fasting glucose and insulin concentrations in an ethnically diverse population: results from the Population Architecture using Genomics and Epidemiology (PAGE) study. BMC Medical Genetics, 2013, 14, 98.	2.1	24
144	Analyses of 7,635 Patients with Colorectal Cancer Using Independent Training and Validation Cohorts Show That rs9929218 in <i>CDH1</i> Is a Prognostic Marker of Survival. Clinical Cancer Research, 2015, 21, 3453-3461.	7.0	24

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145	Genetic Variation in GPX1 Is Associated with GPX1 Activity in a Comprehensive Analysis of Genetic Variations in Selenoenzyme Genes and Their Activity and Oxidative Stress in Humans,. Journal of Nutrition, 2012, 142, 419-426.	2.9	23
146	Association of Cancer Susceptibility Variants with Risk of Multiple Primary Cancers: The Population Architecture using Genomics and Epidemiology Study. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2568-2578.	2.5	23
147	Rare coding variation in paraoxonase-1 is associated with ischemic stroke in the NHLBI Exome Sequencing Project. Journal of Lipid Research, 2014, 55, 1173-1178.	4.2	23
148	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. Human Genetics and Genomics Advances, 2021, 2, 100029.	1.7	23
149	Phenotypic and tumor molecular characterization of colorectal cancer in relation to a susceptibility SMAD7 variant associated with survival. Carcinogenesis, 2013, 34, 292-298.	2.8	22
150	Modeling the effect of prolonged ethanol exposure on global gene expression and chromatin accessibility in normal 3D colon organoids. PLoS ONE, 2020, 15, e0227116.	2.5	22
151	Effects of smoking on the genetic risk of obesity: the population architecture using genomics and epidemiology study. BMC Medical Genetics, 2013, 14, 6.	2.1	21
152	Replication of Associations between GWAS SNPs and Melanoma Risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. Journal of Investigative Dermatology, 2014, 134, 2049-2052.	0.7	21
153	MiRNA-Related SNPs and Risk of Esophageal Adenocarcinoma and Barrett's Esophagus: Post Genome-Wide Association Analysis in the BEACON Consortium. PLoS ONE, 2015, 10, e0128617.	2.5	21
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