

Ulrike Peters

List of Publications by Citations

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Version: 2024-04-27

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

245
papers

17,152
citations

57
h-index

127
g-index

268
ext. papers

22,296
ext. citations

8.8
avg. IF

5.25
L-index

#	Paper	IF	Citations
245	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
244	A reference panel of 64,976 haplotypes for genotype imputation. <i>Nature Genetics</i> , 2016 , 48, 1279-83	36.3	1447
243	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014 , 46, 1173-86	36.3	1339
242	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015 , 518, 187-196	50.4	920
241	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	36.3	516
240	Estimation of effect size distribution from genome-wide association studies and implications for future discoveries. <i>Nature Genetics</i> , 2010 , 42, 570-5	36.3	498
239	Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , 2019 , 570, 514-518	50.4	291
238	Dietary fibre and colorectal adenoma in a colorectal cancer early detection programme. <i>Lancet, The</i> , 2003 , 361, 1491-5	40	262
237	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , 2013 , 144, 799-807.e24	13.3	250
236	Genetic Mechanisms of Immune Evasion in Colorectal Cancer. <i>Cancer Discovery</i> , 2018 , 8, 730-749	24.4	235
235	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, 223-32	11	233
234	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	36.3	226
233	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014 , 46, 994-1000	36.3	226
232	Causes of death in patients with celiac disease in a population-based Swedish cohort. <i>Archives of Internal Medicine</i> , 2003 , 163, 1566-72		195
231	Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms. <i>Nature Genetics</i> , 2017 , 49, 1113-1119	36.3	184
230	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 126-135	4	183
229	Supplemental and dietary vitamin E, beta-carotene, and vitamin C intakes and prostate cancer risk. <i>Journal of the National Cancer Institute</i> , 2006 , 98, 245-54	9.7	177

228	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019 , 51, 76-83	6.3	177
227	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , 2012 , 131, 217-34	6.3	173
226	Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. <i>American Journal of Human Genetics</i> , 2014 , 94, 233-45	11	170
225	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. <i>Molecular Psychiatry</i> , 2015 , 20, 647-656	15.1	167
224	Genome-wide association analyses in East Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013 , 45, 191-6	36.3	155
223	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	9.7	155
222	Meta-analysis of gene-level tests for rare variant association. <i>Nature Genetics</i> , 2014 , 46, 200-4	36.3	142
221	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, 849-59	3.8	141
220	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, 1133-42	27.4	135
219	Genetic architecture of colorectal cancer. <i>Gut</i> , 2015 , 64, 1623-36	19.2	131
218	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	13.3	131
217	A prospective study of lycopene and tomato product intake and risk of prostate cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 92-8	4	127
216	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, 962-8	4	127
215	Characterization of gene-environment interactions for colorectal cancer susceptibility loci. <i>Cancer Research</i> , 2012 , 72, 2036-44	10.1	119
214	Methods for etiologic and early marker investigations in the PLCO trial. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2005 , 592, 147-54	3.3	113
213	Integrative analysis of exogenous, endogenous, tumour and immune factors for precision medicine. <i>Gut</i> , 2018 , 67, 1168-1180	19.2	111
212	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015 , 6, 7138	17.4	106
211	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	17.4	105

210	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018 , 9, 556	17.4	103
209	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017 , 13, e1006528	6	103
208	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	7.8	92
207	Rare and low-frequency coding variants in CXCR2 and other genes are associated with hematological traits. <i>Nature Genetics</i> , 2014 , 46, 629-34	36.3	92
206	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, 502-12	11	90
205	A model to determine colorectal cancer risk using common genetic susceptibility loci. <i>Gastroenterology</i> , 2015 , 148, 1330-9.e14	13.3	89
204	Variation in the selenoenzyme genes and risk of advanced distal colorectal adenoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008 , 17, 1144-54	4	88
203	Estimating the heritability of colorectal cancer. <i>Human Molecular Genetics</i> , 2014 , 23, 3898-905	5.6	85
202	Selenium and the prevention of prostate and colorectal cancer. <i>Molecular Nutrition and Food Research</i> , 2008 , 52, 1261-72	5.9	83
201	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , 2015 , 96, 487-97	11	77
200	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, 6616-33	5.6	77
199	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	2.8	74
198	Serum selenium and risk of prostate cancer-a nested case-control study. <i>American Journal of Clinical Nutrition</i> , 2007 , 85, 209-17	7	71
197	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 146-157	9.7	67
196	Genome-wide diet-gene interaction analyses for risk of colorectal cancer. <i>PLoS Genetics</i> , 2014 , 10, e1004228	4.2	66
195	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016 , 7, 66328-66343	3.3	66
194	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016 , 76, 5103-14	10.1	66
193	Fine Mapping and Identification of BMI Loci in African Americans. <i>American Journal of Human Genetics</i> , 2013 , 93, 661-71	11	63

192	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VT11A. <i>Nature Communications</i> , 2014 , 5, 4613	17.4	62
191	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, 835-46	8	60
190	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
189	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	3.7	59
188	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	6	56
187	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	2.6	55
186	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1024-31	4	54
185	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, 315-20	4	52
184	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
183	Characterization of the association between 8q24 and colon cancer: gene-environment exploration and meta-analysis. <i>BMC Cancer</i> , 2010 , 10, 670	4.8	50
182	Genotype imputation of Metabochip SNPs using a study-specific reference panel of ~4,000 haplotypes in African Americans from the Women's Health Initiative. <i>Genetic Epidemiology</i> , 2012 , 36, 107-17	2.6	49
181	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	11.6	49
180	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,	9.7	47
179	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, 1974-85	4	47
178	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	13.3	47
177	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
176	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	13.3	45
175	Genetic identification of a common collagen disease in puerto ricans via identity-by-descent mapping in a health system. <i>ELife</i> , 2017 , 6,	8.9	44

174	Mendelian randomization study of height and risk of colorectal cancer. <i>International Journal of Epidemiology</i> , 2015 , 44, 662-72	7.8	44
173	Calcium intake and colorectal adenoma in a US colorectal cancer early detection program. <i>American Journal of Clinical Nutrition</i> , 2004 , 80, 1358-65	7	43
172	Consistent directions of effect for established type 2 diabetes risk variants across populations: the population architecture using Genomics and Epidemiology (PAGE) Consortium. <i>Diabetes</i> , 2012 , 61, 1642-7 ⁹	7.9	42
171	Gene-environment interaction involving recently identified colorectal cancer susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 1824-33	4	40
170	Genetic variation in C-reactive protein in relation to colon and rectal cancer risk and survival. <i>International Journal of Cancer</i> , 2011 , 128, 2726-34	7.5	39
169	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. <i>PLoS ONE</i> , 2016 , 11, e0167758	3.7	39
168	Meta-analysis of 16 studies of the association of alcohol with colorectal cancer. <i>International Journal of Cancer</i> , 2020 , 146, 861-873	7.5	39
167	Rare and Coding Region Genetic Variants Associated With Risk of Ischemic Stroke: The NHLBI Exome Sequence Project. <i>JAMA Neurology</i> , 2015 , 72, 781-8	17.2	37
166	Association between colorectal cancer susceptibility loci and survival time after diagnosis with colorectal cancer. <i>Gastroenterology</i> , 2012 , 143, 51-4.e4	13.3	36
165	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. <i>Nature Communications</i> , 2020 , 11, 597	17.4	36
164	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	3.2	35
163	Circulating vitamin D metabolites, polymorphism in vitamin D receptor, and colorectal adenoma risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004 , 13, 546-52	4	34
162	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	4	32
161	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020 , 11, 3353	17.4	32
160	HNF1B and endometrial cancer risk: results from the PAGE study. <i>PLoS ONE</i> , 2012 , 7, e30390	3.7	32
159	Serum Tocopherol and Tocopherol concentrations and prostate cancer risk in the PLCO Screening Trial: a nested case-control study. <i>PLoS ONE</i> , 2012 , 7, e40204	3.7	32
158	Common genetic variation and survival after colorectal cancer diagnosis: a genome-wide analysis. <i>Carcinogenesis</i> , 2016 , 37, 87-95	4.6	31
157	Association of exome sequences with plasma C-reactive protein levels in >9000 participants. <i>Human Molecular Genetics</i> , 2015 , 24, 559-71	5.6	31

156	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020 , 107, 432-444	11	31
155	Genetic variation in calcium-sensing receptor and risk for colon cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008 , 17, 2755-65	4	30
154	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , 2016 , 12, e1006296	6	30
153	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , 2016 , 139, 2655-2670	7.5	30
152	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, 3131-9	4	29
151	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, 1227-31	4	29
150	Genome-wide search for gene-gene interactions in colorectal cancer. <i>PLoS ONE</i> , 2012 , 7, e52535	3.7	29
149	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	9.7	28
148	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, 1822-30	4	28
147	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, 686-94	4.4	28
146	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	5.4	28
145	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , 2014 , 63, 800-7	19.2	27
144	SBERIA: set-based gene-environment interaction test for rare and common variants in complex diseases. <i>Genetic Epidemiology</i> , 2013 , 37, 452-64	2.6	27
143	Genetic predictors of circulating 25-hydroxyvitamin d and risk of colorectal cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 2037-46	4	26
142	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. <i>Human Genetics</i> , 2015 , 134, 1249-1262	6.3	25
141	The influence of obesity-related single nucleotide polymorphisms on BMI across the life course: the PAGE study. <i>Diabetes</i> , 2013 , 62, 1763-7	0.9	25
140	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	9.7	25
139	Selenium, selenoenzymes, oxidative stress and risk of neoplastic progression from Barrett's esophagus: results from biomarkers and genetic variants. <i>PLoS ONE</i> , 2012 , 7, e38612	3.7	24

138	Association of serum phospholipid fatty acids with breast cancer risk among postmenopausal cigarette smokers. <i>Cancer Causes and Control</i> , 2009 , 20, 497-504	2.8	24
137	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	6.3	23
136	Common single-nucleotide polymorphisms in the estrogen receptor β promoter are associated with colorectal cancer survival in postmenopausal women. <i>Cancer Research</i> , 2013 , 73, 767-75	10.1	23
135	Analyses of 7,635 Patients with Colorectal Cancer Using Independent Training and Validation Cohorts Show That rs9929218 in CDH1 Is a Prognostic Marker of Survival. <i>Clinical Cancer Research</i> , 2015 , 21, 3453-61	12.9	22
134	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	5.6	22
133	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. <i>BMC Medical Genetics</i> , 2013 , 14, 98	2.1	22
132	The use of imputed values in the meta-analysis of genome-wide association studies. <i>Genetic Epidemiology</i> , 2011 , 35, 597-605	2.6	22
131	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. <i>Carcinogenesis</i> , 2015 , 36, 999-1007	4.6	21
130	MiRNA-Related SNPs and Risk of Esophageal Adenocarcinoma and Barrett's Esophagus: Post Genome-Wide Association Analysis in the BEACON Consortium. <i>PLoS ONE</i> , 2015 , 10, e0128617	3.7	21
129	Genetic variation predicts serum lycopene concentrations in a multiethnic population of postmenopausal women. <i>Journal of Nutrition</i> , 2015 , 145, 187-92	4.1	21
128	Association of cancer susceptibility variants with risk of multiple primary cancers: The population architecture using genomics and epidemiology study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 2568-78	4	21
127	Variation in selenoenzyme genes and prostate cancer risk and survival. <i>Prostate</i> , 2013 , 73, 734-42	4.2	21
126	Relationship of prediagnostic body mass index with survival after colorectal cancer: Stage-specific associations. <i>International Journal of Cancer</i> , 2016 , 139, 1065-72	7.5	21
125	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. <i>American Journal of Human Genetics</i> , 2018 , 102, 904-919	11	20
124	Effects of smoking on the genetic risk of obesity: the population architecture using genomics and epidemiology study. <i>BMC Medical Genetics</i> , 2013 , 14, 6	2.1	20
123	Phenotypic and tumor molecular characterization of colorectal cancer in relation to a susceptibility SMAD7 variant associated with survival. <i>Carcinogenesis</i> , 2013 , 34, 292-8	4.6	20
122	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. <i>Journal of Nutrition</i> , 2012 , 142, 419-26	4.1	20
121	Sex-specific gene and pathway modeling of inherited glioma risk. <i>Neuro-Oncology</i> , 2019 , 21, 71-82	1	19

120	Rare coding variation in paraoxonase-1 is associated with ischemic stroke in the NHLBI Exome Sequencing Project. <i>Journal of Lipid Research</i> , 2014 , 55, 1173-8	6.3	19
119	Influence of Smoking, Body Mass Index, and Other Factors on the Preventive Effect of Nonsteroidal Anti-Inflammatory Drugs on Colorectal Cancer Risk. <i>Cancer Research</i> , 2018 , 78, 4790-4799	10.1	19
118	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. <i>International Journal of Epidemiology</i> , 2019 , 48, 767-780	7.8	18
117	Association of genetic variants in the calcium-sensing receptor with risk of colorectal adenoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004 , 13, 2181-6	4	18
116	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	10.1	17
115	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	13.3	17
114	Cross-cancer pleiotropic analysis of endometrial cancer: PAGE and E2C2 consortia. <i>Carcinogenesis</i> , 2014 , 35, 2068-73	4.6	17
113	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. <i>BMC Medicine</i> , 2020 , 18, 396	11.4	17
112	Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. <i>Human Genetics</i> , 2019 , 138, 307-326	6.3	17
111	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. <i>Nature Communications</i> , 2020 , 11, 2542	17.4	16
110	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. <i>British Journal of Cancer</i> , 2016 , 114, 221-9	8.7	16
109	Transethnic insight into the genetics of glycaemic traits: fine-mapping results from the Population Architecture using Genomics and Epidemiology (PAGE) consortium. <i>Diabetologia</i> , 2017 , 60, 2384-2398	10.3	16
108	Replication of associations between GWAS SNPs and melanoma risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. <i>Journal of Investigative Dermatology</i> , 2014 , 134, 2049-2052	4.3	16
107	Landscape of somatic single nucleotide variants and indels in colorectal cancer and impact on survival. <i>Nature Communications</i> , 2020 , 11, 3644	17.4	16
106	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 557-567	9.7	16
105	Fine mapping of QT interval regions in global populations refines previously identified QT interval loci and identifies signals unique to African and Hispanic descent populations. <i>Heart Rhythm</i> , 2017 , 14, 572-580	6.7	15
104	Genetic variants of adiponectin and risk of colorectal cancer. <i>International Journal of Cancer</i> , 2015 , 137, 154-64	7.5	15
103	Glutathione peroxidase tagSNPs: associations with rectal cancer but not with colon cancer. <i>Genes Chromosomes and Cancer</i> , 2012 , 51, 598-605	5	15

102	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021 , 5, pkbab009	15	
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	13.3	15
100	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. <i>PLoS ONE</i> , 2018 , 13, e0200486	3.7	14
99	Pleiotropic and sex-specific effects of cancer GWAS SNPs on melanoma risk in the population architecture using genomics and epidemiology (PAGE) study. <i>PLoS ONE</i> , 2015 , 10, e0120491	3.7	14
98	Pleiotropy of cancer susceptibility variants on the risk of non-Hodgkin lymphoma: the PAGE consortium. <i>PLoS ONE</i> , 2014 , 9, e89791	3.7	14
97	Variation in the association between colorectal cancer susceptibility loci and colorectal polyps by polyp type. <i>American Journal of Epidemiology</i> , 2014 , 180, 223-32	3.8	13
96	Fine-mapping and initial characterization of QT interval loci in African Americans. <i>PLoS Genetics</i> , 2012 , 8, e1002870	6	13
95	A haplotype-based test of association using data from cohort and nested case-control epidemiologic studies. <i>Human Heredity</i> , 2004 , 58, 18-29	1.1	13
94	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. <i>International Journal of Cancer</i> , 2020 , 146, 363-372	7.5	13
93	Anthropometric Risk Factors for Cancers of the Biliary Tract in the Biliary Tract Cancers Pooling Project. <i>Cancer Research</i> , 2019 , 79, 3973-3982	10.1	12
92	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 860-870	4	12
91	Modeling the effect of prolonged ethanol exposure on global gene expression and chromatin accessibility in normal 3D colon organoids. <i>PLoS ONE</i> , 2020 , 15, e0227116	3.7	12
90	Association of with Specific T-cell Subsets in the Colorectal Carcinoma Microenvironment. <i>Clinical Cancer Research</i> , 2021 , 27, 2816-2826	12.9	12
89	Genome-wide association study of heart rate and its variability in Hispanic/Latino cohorts. <i>Heart Rhythm</i> , 2017 , 14, 1675-1684	6.7	11
88	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	9.7	11
87	Powerful Set-Based Gene-Environment Interaction Testing Framework for Complex Diseases. <i>Genetic Epidemiology</i> , 2015 , 39, 609-18	2.6	11
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