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List of Publications by Citations

719 papers	45,875 citations	101 h-index	189 g-index
748 ext. papers	55,297 ext. citations	7.7 avg, IF	6.52 L-index

#	Paper	IF	Citations
719	Genetic studies of body mass index yield new insights for obesity biology. *Nature*, **2015**, 518, 197-206	50.4	2687
718	Genome-wide association study identifies novel breast cancer susceptibility loci. *Nature*, **2007**, 447, 1087-93	50.4	1957
717	Defining the role of common variation in the genomic and biological architecture of adult human height. *Nature Genetics*, **2014**, 46, 1173-86	36.3	1339
716	New genetic loci link adipose and insulin biology to body fat distribution. *Nature*, **2015**, 518, 187-196	50.4	920
715	Risk of COVID-19 among front-line health-care workers and the general community: a prospective cohort study. *Lancet Public Health, The*, **2020**, 5, e475-e483	22.4	899
714	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. *Nature Genetics*, **2013**, 45, 353-61, 361e1-2	36.3	813
713	Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptor-positive breast cancer. *Nature Genetics*, **2007**, 39, 865-9	36.3	715
712	Association analysis identifies 65 new breast cancer risk loci. *Nature*, **2017**, 551, 92-94	50.4	643
711	Multiple regions within 8q24 independently affect risk for prostate cancer. *Nature Genetics*, **2007**, 39, 638-44	36.3	563
710	Ethnic and racial differences in the smoking-related risk of lung cancer. *New England Journal of Medicine*, **2006**, 354, 333-42	59.2	538
709	Associations of breast cancer risk factors with tumor subtypes: a pooled analysis from the Breast Cancer Association Consortium studies. *Journal of the National Cancer Institute*, **2011**, 103, 250-63	9.7	513
708	CCAT2, a novel noncoding RNA mapping to 8q24, underlies metastatic progression and chromosomal instability in colon cancer. *Genome Research*, **2013**, 23, 1446-61	9.7	442
707	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. *Nature Genetics*, **2013**, 45, 371-84, 384e1-2	36.3	422
706	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. *Nature Genetics*, **2013**, 45, 385-91, 391e1-2	36.3	413
705	Detectable clonal mosaicism from birth to old age and its relationship to cancer. *Nature Genetics*, **2012**, 44, 642-50	36.3	409
704	Detectable clonal mosaicism and its relationship to aging and cancer. *Nature Genetics*, **2012**, 44, 651-8	36.3	409
703	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. *Nature Genetics*, **2015**, 47, 373-80	36.3	406

702	Cancer preventive effects of flavonoids--a review. *Biomedicine and Pharmacotherapy*, **2002**, 56, 296-301	7.5	403
701	Common variants on chromosome 5p12 confer susceptibility to estrogen receptor-positive breast cancer. *Nature Genetics*, **2008**, 40, 703-6	36.3	378
700	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. *American Journal of Human Genetics*, **2019**, 104, 21-34	11	363
699	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. *Nature Genetics*, **2009**, 41, 1116-21	36.3	360
698	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. *Nature Genetics*, **2018**, 50, 928-936	36.3	340
697	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. *Nature Genetics*, **2014**, 46, 1103-9	36.3	331
696	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. *Nature Genetics*, **2013**, 45, 392-8, 398e1-2	36.3	327
695	Identification of Lynch syndrome among patients with colorectal cancer. *JAMA - Journal of the American Medical Association*, **2012**, 308, 1555-65	27.4	323
694	A common genetic risk factor for colorectal and prostate cancer. *Nature Genetics*, **2007**, 39, 954-6	36.3	304
693	Meta-analysis identifies common variants associated with body mass index in east Asians. *Nature Genetics*, **2012**, 44, 307-11	36.3	301
692	Genetic analyses of diverse populations improves discovery for complex traits. *Nature*, **2019**, 570, 514-518	50.4	291
691	Heterogeneity of breast cancer associations with five susceptibility loci by clinical and pathological characteristics. *PLoS Genetics*, **2008**, 4, e1000054	6	280
690	Risks of Lynch syndrome cancers for MSH6 mutation carriers. *Journal of the National Cancer Institute*, **2010**, 102, 193-201	9.7	279
689	Colon Cancer Family Registry: an international resource for studies of the genetic epidemiology of colon cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2007**, 16, 2331-43	4	279
688	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. *Nature Genetics*, **2011**, 43, 1210-4	36.3	253
687	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. *Gastroenterology*, **2013**, 144, 799-807.e24	13.3	250
686	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. *Nature Genetics*, **2017**, 49, 1126-1132	36.3	246
685	Colorectal and other cancer risks for carriers and noncarriers from families with a DNA mismatch repair gene mutation: a prospective cohort study. *Journal of Clinical Oncology*, **2012**, 30, 958-64	2.2	245

684	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. *Nature Genetics*, **2011**, 43, 785-91	36.3	243
683	The landscape of recombination in African Americans. *Nature*, **2011**, 476, 170-5	50.4	243
682	Meta- and pooled analyses of the effects of glutathione S-transferase M1 polymorphisms and smoking on lung cancer risk. *Carcinogenesis*, **2002**, 23, 1343-50	4.6	228
681	Associations of key diet-quality indexes with mortality in the Multiethnic Cohort: the Dietary Patterns Methods Project. *American Journal of Clinical Nutrition*, **2015**, 101, 587-97	7	227
680	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. *Nature Genetics*, **2014**, 46, 994-1000	36.3	226
679	Metabolism and biomarkers of heterocyclic aromatic amines in molecular epidemiology studies: lessons learned from aromatic amines. *Chemical Research in Toxicology*, **2011**, 24, 1169-214	4	209
678	Body size at different periods of life and breast cancer risk. *American Journal of Epidemiology*, **1988**, 128, 137-52	3.8	208
677	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. *Nature Genetics*, **2009**, 41, 1055-7	36.3	201
676	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. *Nature Genetics*, **2013**, 45, 690-6	36.3	192
675	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers: the advantage of more extensive colon surgery. *Gut*, **2011**, 60, 950-7	19.2	192
674	Genome-wide association study of glioma and meta-analysis. *Human Genetics*, **2012**, 131, 1877-88	6.3	191
673	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. *Nature Genetics*, **2017**, 49, 680-691	36.3	190
672	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. *JAMA Oncology*, **2016**, 2, 1295-1302	13.4	189
671	Sun exposure and melanoma risk at different latitudes: a pooled analysis of 5700 cases and 7216 controls. *International Journal of Epidemiology*, **2009**, 38, 814-30	7.8	187
670	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. *Nature Genetics*, **2017**, 49, 1767-1778	36.3	186
669	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 404-412	4	185
668	Discovery of common and rare genetic risk variants for colorectal cancer. *Nature Genetics*, **2019**, 51, 76-87	36.3	177
667	Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. *Hepatology*, **2016**, 64, 1969-1977	11.2	175

666	Meta-analysis of new genome-wide association studies of colorectal cancer risk. *Human Genetics*, **2012**, 131, 217-34	6.3	173
665	Cancer risks for MLH1 and MSH2 mutation carriers. *Human Mutation*, **2013**, 34, 490-7	4.7	171
664	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. *Nature Genetics*, **2011**, 43, 570-3	36.3	171
663	Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. *American Journal of Human Genetics*, **2013**, 92, 489-503	11	167
662	Smokers with the CHRNA lung cancer-associated variants are exposed to higher levels of nicotine equivalents and a carcinogenic tobacco-specific nitrosamine. *Cancer Research*, **2008**, 68, 9137-40	10.1	166
661	Association of a low-frequency variant in HNF1A with type 2 diabetes in a Latino population. *JAMA - Journal of the American Medical Association*, **2014**, 311, 2305-14	27.4	164
660	Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants: findings from the Prospective Lynch Syndrome Database. *Genetics in Medicine*, **2020**, 22, 15-25	8.1	164
659	Risks of primary extracolonic cancers following colorectal cancer in lynch syndrome. *Journal of the National Cancer Institute*, **2012**, 104, 1363-72	9.7	158
658	Generalization and dilution of association results from European GWAS in populations of non-European ancestry: the PAGE study. *PLoS Biology*, **2013**, 11, e1001661	9.7	155
657	Circulating 25-hydroxyvitamin D and risk of pancreatic cancer: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. *American Journal of Epidemiology*, **2010**, 172, 81-93	3.8	155
656	Replication of lung cancer susceptibility loci at chromosomes 15q25, 5p15, and 6p21: a pooled analysis from the International Lung Cancer Consortium. *Journal of the National Cancer Institute*, **2010**, 102, 959-71	9.7	153
655	CYP1A1 and GSTM1 genetic polymorphisms and lung cancer risk in Caucasian non-smokers: a pooled analysis. *Carcinogenesis*, **2003**, 24, 875-82	4.6	152
654	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. *Human Molecular Genetics*, **2012**, 21, 5373-84	5.6	143
653	Genome-wide association study identifies new prostate cancer susceptibility loci. *Human Molecular Genetics*, **2011**, 20, 3867-75	5.6	143
652	Consistent association of type 2 diabetes risk variants found in europeans in diverse racial and ethnic groups. *PLoS Genetics*, **2010**, 6, e1001078	6	142
651	B-vitamin intake, metabolic genes, and colorectal cancer risk (United States). *Cancer Causes and Control*, **2002**, 13, 239-48	2.8	142
650	The Next PAGE in understanding complex traits: design for the analysis of Population Architecture Using Genetics and Epidemiology (PAGE) Study. *American Journal of Epidemiology*, **2011**, 174, 849-59	3.8	141
649	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2011**, 20, 3289-303	5.6	140

648	Association of aspirin and NSAID use with risk of colorectal cancer according to genetic variants. *JAMA - Journal of the American Medical Association*, **2015**, 313, 1133-42	27.4	135
647	Interactions between genetic variants and breast cancer risk factors in the breast and prostate cancer cohort consortium. *Journal of the National Cancer Institute*, **2011**, 103, 1252-63	9.7	134
646	Genetic variation at the CYP19A1 locus predicts circulating estrogen levels but not breast cancer risk in postmenopausal women. *Cancer Research*, **2007**, 67, 1893-7	10.1	134
645	Case-control study of overweight, obesity, and colorectal cancer risk, overall and by tumor microsatellite instability status. *Journal of the National Cancer Institute*, **2010**, 102, 391-400	9.7	133
644	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. *Gastroenterology*, **2018**, 154, 2152-2164.e19	13.3	131
643	Circulating Vitamin D and Colorectal Cancer Risk: An International Pooling Project of 17 Cohorts. *Journal of the National Cancer Institute*, **2019**, 111, 158-169	9.7	131
642	Genetic determinants of lipid traits in diverse populations from the population architecture using genomics and epidemiology (PAGE) study. *PLoS Genetics*, **2011**, 7, e1002138	6	128
641	A common 8q24 variant in prostate and breast cancer from a large nested case-control study. *Cancer Research*, **2007**, 67, 2951-6	10.1	127
640	Phenome-wide association study (PheWAS) for detection of pleiotropy within the Population Architecture using Genomics and Epidemiology (PAGE) Network. *PLoS Genetics*, **2013**, 9, e1003087	6	126
639	Previous lung diseases and lung cancer risk: a pooled analysis from the International Lung Cancer Consortium. *American Journal of Epidemiology*, **2012**, 176, 573-85	3.8	123
638	Association of Coffee Consumption With Total and Cause-Specific Mortality Among Nonwhite Populations. *Annals of Internal Medicine*, **2017**, 167, 228-235	8	121
637	CYP1A1 T3801 C polymorphism and lung cancer: a pooled analysis of 2451 cases and 3358 controls. *International Journal of Cancer*, **2003**, 104, 650-7	7.5	121
636	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. *Journal of Medical Genetics*, **2016**, 53, 800-811	5.8	121
635	Genome-wide association study in east Asians identifies novel susceptibility loci for breast cancer. *PLoS Genetics*, **2012**, 8, e1002532	6	118
634	A Population-Based Study of Genes Previously Implicated in Breast Cancer. *New England Journal of Medicine*, **2021**, 384, 440-451	59.2	115
633	Association of coffee intake with reduced incidence of liver cancer and death from chronic liver disease in the US multiethnic cohort. *Gastroenterology*, **2015**, 148, 118-25; quiz e15	13.3	109
632	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. *Human Molecular Genetics*, **2013**, 22, 408-15	5.6	109
631	Obesity in youth and middle age and risk of colorectal cancer in men. *Cancer Causes and Control*, **1992**, 3, 349-54	2.8	109

630	Exposure of nonsmoking women to environmental tobacco smoke: a 10-country collaborative study. *Cancer Causes and Control*, **1990**, 1, 243-52	2.8	108
629	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. *Journal of the National Cancer Institute*, **2015**, 107, djv279	9.7	107
628	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. *Nature Communications*, **2015**, 6, 7138	17.4	106
627	The contribution of rare variation to prostate cancer heritability. *Nature Genetics*, **2016**, 48, 30-5	36.3	106
626	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. *Nature Communications*, **2017**, 8, 14977	17.4	105
625	Correlates of circulating 25-hydroxyvitamin D: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. *American Journal of Epidemiology*, **2010**, 172, 21-35	3.8	105
624	Genetic determinants of telomere length and risk of common cancers: a Mendelian randomization study. *Human Molecular Genetics*, **2015**, 24, 5356-66	5.6	104
623	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. *Nature Communications*, **2018**, 9, 556	17.4	103
622	Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. *Carcinogenesis*, **2014**, 35, 2526-33	4.6	103
621	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. *PLoS Genetics*, **2017**, 13, e1006528	6	103
620	Vegetable and fruit consumption in relation to prostate cancer risk in Hawaii: a reevaluation of the effect of dietary beta-carotene. *American Journal of Epidemiology*, **1991**, 133, 215-9	3.8	102
619	Cancer Risks for PMS2-Associated Lynch Syndrome. *Journal of Clinical Oncology*, **2018**, 36, 2961-2968	2.2	102
618	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. *Nature Genetics*, **2018**, 50, 968-978	36.3	101
617	FGFR2 variants and breast cancer risk: fine-scale mapping using African American studies and analysis of chromatin conformation. *Human Molecular Genetics*, **2009**, 18, 1692-703	5.6	100
616	Characterizing genetic risk at known prostate cancer susceptibility loci in African Americans. *PLoS Genetics*, **2011**, 7, e1001387	6	98
615	Associations between smoking, alcohol consumption, and colorectal cancer, overall and by tumor microsatellite instability status. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 2745-50	4	98
614	Common genetic variation in IGF1 and prostate cancer risk in the Multiethnic Cohort. *Journal of the National Cancer Institute*, **2006**, 98, 123-34	9.7	97
613	Quality assessment and correlation of microsatellite instability and immunohistochemical markers among population- and clinic-based colorectal tumors results from the Colon Cancer Family Registry. *Journal of Molecular Diagnostics*, **2011**, 13, 271-81	5.1	95

612	Generalizability of associations from prostate cancer genome-wide association studies in multiple populations. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 1285-9	4	95
611	Trans-ethnic fine-mapping of lipid loci identifies population-specific signals and allelic heterogeneity that increases the trait variance explained. *PLoS Genetics*, **2013**, 9, e1003379	6	94
610	Prediagnostic leptin, adiponectin, C-reactive protein, and the risk of postmenopausal breast cancer. *Cancer Prevention Research*, **2013**, 6, 188-95	3.2	94
609	Identification of a functional genetic variant at 16q12.1 for breast cancer risk: results from the Asia Breast Cancer Consortium. *PLoS Genetics*, **2010**, 6, e1001002	6	93
608	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). *PLoS ONE*, **2008**, 3, e2578	3.7	93
607	Common breast cancer susceptibility variants in LSP1 and RAD51L1 are associated with mammographic density measures that predict breast cancer risk. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 1156-66	4	92
606	Carotenoids, retinol, tocopherols, and prostate cancer risk: pooled analysis of 15 studies. *American Journal of Clinical Nutrition*, **2015**, 102, 1142-57	7	89
605	Plasma 25-hydroxyvitamin D levels and the risk of colorectal cancer: the multiethnic cohort study. *Cancer Epidemiology Biomarkers and Prevention*, **2010**, 19, 130-4	4	89
604	A review of the application of inflammatory biomarkers in epidemiologic cancer research. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 1729-51	4	88
603	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. *Human Molecular Genetics*, **2013**, 22, 2520-8	5.6	88
602	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. *Nature Communications*, **2014**, 4, 4999	17.4	87
601	Risk of metachronous colon cancer following surgery for rectal cancer in mismatch repair gene mutation carriers. *Annals of Surgical Oncology*, **2013**, 20, 1829-36	3.1	87
600	A Meta-analysis of Individual Participant Data Reveals an Association between Circulating Levels of IGF-I and Prostate Cancer Risk. *Cancer Research*, **2016**, 76, 2288-2300	10.1	85
599	Estimating the heritability of colorectal cancer. *Human Molecular Genetics*, **2014**, 23, 3898-905	5.6	85
598	Common variants at 11q12, 10q26 and 3p11.2 are associated with prostate cancer susceptibility in Japanese. *Nature Genetics*, **2012**, 44, 426-9, S1	36.3	84
597	Pooled analysis of the CYP1A1 exon 7 polymorphism and lung cancer (United States). *Cancer Causes and Control*, **2003**, 14, 339-46	2.8	83
596	Variants on 9p24 and 8q24 are associated with risk of colorectal cancer: results from the Colon Cancer Family Registry. *Cancer Research*, **2007**, 67, 11128-32	10.1	82
595	Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. *American Journal of Human Genetics*, **2013**, 93, 1046-60	11	80

594	Lynch syndrome-associated breast cancers: clinicopathologic characteristics of a case series from the colon cancer family registry. *Clinical Cancer Research*, **2010**, 16, 2214-24	12.9	80
593	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. *PLoS Medicine*, **2016**, 13, e1002105	11.6	80
592	Genome-wide association study identifies breast cancer risk variant at 10q21.2: results from the Asia Breast Cancer Consortium. *Human Molecular Genetics*, **2011**, 20, 4991-9	5.6	79
591	Characterization of large structural genetic mosaicism in human autosomes. *American Journal of Human Genetics*, **2015**, 96, 487-97	11	77
590	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. *Human Molecular Genetics*, **2014**, 23, 6616-33	5.6	77
589	Meat and heterocyclic amine intake, smoking, NAT1 and NAT2 polymorphisms, and colorectal cancer risk in the multiethnic cohort study. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 2098-106	4	77
588	Identification, replication, and fine-mapping of Loci associated with adult height in individuals of african ancestry. *PLoS Genetics*, **2011**, 7, e1002298	6	77
587	Nicotine metabolism in three ethnic/racial groups with different risks of lung cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2008**, 17, 3526-35	4	77
586	Dietary fiber and colorectal cancer risk: the multiethnic cohort study. *Cancer Causes and Control*, **2007**, 18, 753-64	2.8	77
585	Two susceptibility loci identified for prostate cancer aggressiveness. *Nature Communications*, **2015**, 6, 6889	17.4	75
584	Association of ESR1 gene tagging SNPs with breast cancer risk. *Human Molecular Genetics*, **2009**, 18, 1131-9	5.6	75
583	Sun exposure, diet, and melanoma in Hawaii Caucasians. *American Journal of Epidemiology*, **2006**, 164, 232-45	3.8	75
582	Association of the cyclin D1 A870G polymorphism with advanced colorectal cancer. *JAMA - Journal of the American Medical Association*, **2003**, 290, 2843-8	27.4	73
581	Fine mapping of the association with obesity at the FTO locus in African-derived populations. *Human Molecular Genetics*, **2010**, 19, 2907-16	5.6	72
580	Cyclin D1 splice variants: polymorphism, risk, and isoform-specific regulation in prostate cancer. *Clinical Cancer Research*, **2009**, 15, 5338-49	12.9	72
579	A pooled analysis of melanocytic nevus phenotype and the risk of cutaneous melanoma at different latitudes. *International Journal of Cancer*, **2009**, 124, 420-8	7.5	72
578	Frequency of deletions of EPCAM (TACSTD1) in MSH2-associated Lynch syndrome cases. *Journal of Molecular Diagnostics*, **2011**, 13, 93-9	5.1	71
577	Identification of nine new susceptibility loci for endometrial cancer. *Nature Communications*, **2018**, 9, 3166	17.4	70

576	Circulating fatty acids and prostate cancer risk in a nested case-control study: the Multiethnic Cohort. *Cancer Causes and Control*, **2009**, 20, 211-23	2.8	70
575	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?. *Nutrients*, **2018**, 10,	6.7	69
574	Exposure to secondhand tobacco smoke and lung cancer by histological type: a pooled analysis of the International Lung Cancer Consortium (ILCCO). *International Journal of Cancer*, **2014**, 135, 1918-30	7.5	69
573	Lifetime occupational physical activity and prostate cancer risk. *American Journal of Epidemiology*, **1991**, 133, 103-11	3.8	69
572	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in MUTYH. *International Journal of Cancer*, **2016**, 139, 1557-63	7.5	67
571	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. *Journal of the National Cancer Institute*, **2019**, 111, 146-157	9.7	67
570	Meat consumption, heterocyclic amines and colorectal cancer risk: the Multiethnic Cohort Study. *International Journal of Cancer*, **2012**, 131, E1125-33	7.5	67
569	High-Quality Diets Associate With Reduced Risk of Colorectal Cancer: Analyses of Diet Quality Indexes in the Multiethnic Cohort. *Gastroenterology*, **2017**, 153, 386-394.e2	13.3	66
568	A priori-defined diet quality indexes and risk of type 2 diabetes: the Multiethnic Cohort. *Diabetologia*, **2015**, 58, 98-112	10.3	66
567	Genome-wide diet-gene interaction analyses for risk of colorectal cancer. *PLoS Genetics*, **2014**, 10, e1004228	8	66
566	Generalizability and epidemiologic characterization of eleven colorectal cancer GWAS hits in multiple populations. *Cancer Epidemiology Biomarkers and Prevention*, **2011**, 20, 70-81	4	66
565	Cancer risks for monoallelic MUTYH mutation carriers with a family history of colorectal cancer. *International Journal of Cancer*, **2011**, 129, 2256-62	7.5	66
564	Polymorphisms in base excision repair genes as colorectal cancer risk factors and modifiers of the effect of diets high in red meat. *Cancer Epidemiology Biomarkers and Prevention*, **2010**, 19, 3167-73	4	66
563	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. *Oncotarget*, **2016**, 7, 66328-66343	3.3	66
562	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. *Cancer Research*, **2016**, 76, 5103-14	10.1	66
561	Heterocyclic amines content of meat and fish cooked by Brazilian methods. *Journal of Food Composition and Analysis*, **2010**, 23, 61-69	4.1	65
560	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. *Gastroenterology*, **2016**, 150, 1633-1645	13.3	64
559	Circulating 25-hydroxyvitamin D and risk of esophageal and gastric cancer: Cohort Consortium Vitamin D Pooling Project of Rarer Cancers. *American Journal of Epidemiology*, **2010**, 172, 94-106	3.8	64

558	Fine Mapping and Identification of BMI Loci in African Americans. *American Journal of Human Genetics*, **2013**, 93, 661-71	11	63
557	A genome-wide association study of breast cancer in women of African ancestry. *Human Genetics*, **2013**, 132, 39-48	6.3	63
556	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTI1A. *Nature Communications*, **2014**, 5, 4613	17.4	62
555	PTGS2 and IL6 genetic variation and risk of breast and prostate cancer: results from the Breast and Prostate Cancer Cohort Consortium (BPC3). *Carcinogenesis*, **2010**, 31, 455-61	4.6	62
554	Association of selenium, tocopherols, carotenoids, retinol, and 15-isoprostane F(2t) in serum or urine with prostate cancer risk: the multiethnic cohort. *Cancer Causes and Control*, **2009**, 20, 1161-71	2.8	62
553	Genetic and functional analysis of CHEK2 (CHK2) variants in multiethnic cohorts. *International Journal of Cancer*, **2007**, 121, 2661-7	7.5	62
552	Association of a common polymorphism in the human GH1 gene with colorectal neoplasia. *Journal of the National Cancer Institute*, **2002**, 94, 454-60	9.7	62
551	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. *Nature Genetics*, **2021**, 53, 65-75	36.3	62
550	Replication and functional genomic analyses of the breast cancer susceptibility locus at 6q25.1 generalize its importance in women of chinese, Japanese, and European ancestry. *Cancer Research*, **2011**, 71, 1344-55	10.1	60
549	Common genetic variation in the sex steroid hormone-binding globulin (SHBG) gene and circulating shbg levels among postmenopausal women: the Multiethnic Cohort. *Journal of Clinical Endocrinology and Metabolism*, **2005**, 90, 2198-204	5.6	60
548	Serologic Response to Helicobacter pylori Proteins Associated With Risk of Colorectal Cancer Among Diverse Populations in the United States. *Gastroenterology*, **2019**, 156, 175-186.e2	13.3	60
547	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. *Nature Communications*, **2016**, 7, 11843	17.4	59
546	Association of the colorectal CpG island methylator phenotype with molecular features, risk factors, and family history. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 512-519	4	59
545	Fine-scale mapping of the 5q11.2 breast cancer locus reveals at least three independent risk variants regulating MAP3K1. *American Journal of Human Genetics*, **2015**, 96, 5-20	11	59
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