Mark E Jenkins

List of Publications by Year
in descending order

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Version: 2024-02-01

		6486	11946
421	25,821	82	139
papers	citations	h-index	g-index

446	446	446	28058
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Informed choice and attitudes regarding a genomic test to predict risk of colorectal cancer in general practice. Patient Education and Counseling, 2022, 105, 987-995.	1.0	7
2	The SMARTscreen Trial: a randomised controlled trial investigating the efficacy of a GP-endorsed narrative SMS to increase participation in the Australian National Bowel Cancer Screening Program. Trials, 2022, 23, 31.	0.7	0
3	Genome-wide association study identifies tumor anatomical site-specific risk variants for colorectal cancer survival. Scientific Reports, 2022, 12, 127.	1.6	6
4	Genetic variants associated with circulating Câ€reactive protein levels and colorectal cancer survival: Sexâ€specific and lifestyle factors specific associations. International Journal of Cancer, 2022, 150, 1447-1454.	2.3	2
5	Risk Stratification for Early-Onset Colorectal Cancer Using a Combination of Genetic and Environmental Risk Scores: An International Multi-Center Study. Journal of the National Cancer Institute, 2022, , .	3.0	15
6	Genetically proxied therapeutic inhibition of antihypertensive drug targets and risk of common cancers: A mendelian randomization analysis. PLoS Medicine, 2022, 19, e1003897.	3.9	30
7	Worldwide prevalence of Lynch syndrome in patients with colorectal cancer: Systematic review and meta-analysis. Genetics in Medicine, 2022, 24, 971-985.	1.1	18
8	Familial Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 1483.	1.7	6
9	Diabetes mellitus in relation to colorectal tumor molecular subtypes â€•a pooled analysis of more than 9,000 cases. International Journal of Cancer, 2022, , .	2.3	2
10	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1077-1089.	1.1	6
11	Germline MBD4 deficiency causes a multi-tumor predisposition syndrome. American Journal of Human Genetics, 2022, 109, 953-960.	2.6	23
12	OUP accepted manuscript. Journal of the National Cancer Institute, 2022, , .	3.0	0
13	Exploring a novel method for optimising the implementation of a colorectal cancer risk prediction tool into primary care: a qualitative study. Implementation Science, 2022, 17, 31.	2.5	5
14	Cancer Risk C (CR-C), a functional genomics test is a sensitive and rapid test for germline mismatch repair deficiency. Genetics in Medicine, 2022, 24, 1821-1830.	1.1	2
15	Genetic Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 2767.	1.7	5
16	Identifying colorectal cancer caused by biallelic MUTYH pathogenic variants using tumor mutational signatures. Nature Communications, 2022, 13, .	5.8	15
17	Association of Body Mass Index With Colorectal Cancer Risk by Genome-Wide Variants. Journal of the National Cancer Institute, 2021, 113, 38-47.	3.0	14
18	Analysis in the Prospective Lynch Syndrome Database identifies sarcoma as part of the Lynch syndrome tumor spectrum. International Journal of Cancer, 2021, 148, 512-513.	2.3	9

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19	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. Gastroenterology, 2021, 160, 1164-1178.e6.	0.6	36
20	Risk-reducing hysterectomy and bilateral salpingo-oophorectomy in female heterozygotes of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. Genetics in Medicine, 2021, 23, 705-712.	1.1	28
21	Novel mammogramâ€based measures improve breast cancer risk prediction beyond an established mammographic density measure. International Journal of Cancer, 2021, 148, 2193-2202.	2.3	18
22	Associations of Height With the Risks of Colorectal and Endometrial Cancer in Persons With Lynch Syndrome. American Journal of Epidemiology, 2021, 190, 230-238.	1.6	2
23	Evaluating the utility of tumour mutational signatures for identifying hereditary colorectal cancer and polyposis syndrome carriers. Gut, 2021, 70, 2138-2149.	6.1	27
24	Lack of an association between gallstone disease and bilirubin levels with risk of colorectal cancer: a Mendelian randomisation analysis. British Journal of Cancer, 2021, 124, 1169-1174.	2.9	6
25	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.	2.2	27
26	An inverse stageâ€shift model to estimate the excess mortality and health economic impact of delayed access to cancer services due to the COVIDâ€19 pandemic. Asia-Pacific Journal of Clinical Oncology, 2021, 17, 359-367.	0.7	59
27	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	6.1	44
28	Rare Variants in the DNA Repair Pathway and the Risk of Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 895-903.	1.1	3
29	Germline and Tumor Sequencing as a Diagnostic Tool To Resolve Suspected Lynch Syndrome. Journal of Molecular Diagnostics, 2021, 23, 358-371.	1.2	12
30	Response to Li and Hopper. American Journal of Human Genetics, 2021, 108, 527-529.	2.6	5
31	Assessment of a Polygenic Risk Score for Colorectal Cancer to Predict Risk of Lynch Syndrome Colorectal Cancer. JNCI Cancer Spectrum, 2021, 5, pkab022.	1.4	15
32	Uptake of hysterectomy and bilateral salpingo-oophorectomy in carriers of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. European Journal of Cancer, 2021, 148, 124-133.	1.3	11
33	Nongenetic Determinants of Risk forÂEarly-Onset Colorectal Cancer. JNCI Cancer Spectrum, 2021, 5, pkab029.	1.4	39
34	DNA Methylation Signatures and the Contribution of Age-Associated Methylomic Drift to Carcinogenesis in Early-Onset Colorectal Cancer. Cancers, 2021, 13, 2589.	1.7	18
35	A Meta-Analysis of Obesity and Risk of Colorectal Cancer in Patients with Lynch Syndrome: The Impact of Sex and Genetics. Nutrients, 2021, 13, 1736.	1.7	10
36	Genetically Predicted Circulating C-Reactive Protein Concentration and Colorectal Cancer Survival: A Mendelian Randomization Consortium Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1349-1358.	1.1	6

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37	Association between Smoking and Molecular Subtypes of Colorectal Cancer. JNCI Cancer Spectrum, 2021, 5, pkab056.	1.4	8
38	No Difference in Penetrance between Truncating and Missense/Aberrant Splicing Pathogenic Variants in MLH1 and MSH2: A Prospective Lynch Syndrome Database Study. Journal of Clinical Medicine, 2021, 10, 2856.	1.0	11
39	Abstract 817: Probing the diabetes - colorectal cancer link using gene - environment interaction analyses. , 2021, , .		0
40	Tetranucleotide and Low Microsatellite Instability Are Inversely Associated with the CpG Island Methylator Phenotype in Colorectal Cancer. Cancers, 2021, 13, 3529.	1.7	3
41	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. Lancet Oncology, The, 2021, 22, 1014-1022.	5.1	58
42	32Do the risks of Lynch syndrome-related cancers depend on the parent-of-origin of the mutation?. International Journal of Epidemiology, 2021, 50, .	0.9	0
43	Rare germline variants in the AXIN2 gene in families with colonic polyposis and colorectal cancer. Familial Cancer, 2021, , 1.	0.9	5
44	DNA repair and cancer in colon and rectum: Novel players in genetic susceptibility. International Journal of Cancer, 2020, 146, 363-372.	2.3	40
45	Interval breast cancer risk associations with breast density, family history and breast tissue aging. International Journal of Cancer, 2020, 147, 375-382.	2.3	22
46	Development and external validation study of a melanoma risk prediction model incorporating clinically assessed naevi and solar lentigines. British Journal of Dermatology, 2020, 182, 1262-1268.	1.4	12
47	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.	1.1	365
48	Modifiable pathways for colorectal cancer: a mendelian randomisation analysis. The Lancet Gastroenterology and Hepatology, 2020, 5, 55-62.	3.7	79
49	Cumulative Burden of Colorectal Cancerâ€"Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	0.6	110
50	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. Gastroenterology, 2020, 158, 1300-1312.e20.	0.6	90
51	Cost-Effectiveness of Personalized Screening for Colorectal Cancer Based on Polygenic Risk and Family History. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 10-21.	1.1	22
52	Postmenopausal Hormone Therapy and Colorectal Cancer Risk by Molecularly Defined Subtypes and Tumor Location. JNCI Cancer Spectrum, 2020, 4, pkaa042.	1.4	8
53	Landscape of somatic single nucleotide variants and indels in colorectal cancer and impact on survival. Nature Communications, 2020, 11, 3644.	5.8	55
54	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1800-1808.	1.1	1

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55	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	2.6	124
56	A Genomic Test for Colorectal Cancer Risk: Is This Acceptable and Feasible in Primary Care?. Public Health Genomics, 2020, 23, 110-121.	0.6	12
57	Pathways to a cancer-free future: a protocol for modelled evaluations to minimise the future burden of colorectal cancer in Australia. BMJ Open, 2020, 10, e036475.	0.8	1
58	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. BMC Medicine, 2020, 18, 229.	2.3	28
59	The Impact of a Comprehensive Risk Prediction Model for Colorectal Cancer on a Population Screening Program. JNCI Cancer Spectrum, 2020, 4, pkaa062.	1.4	5
60	A general framework for functionally informed set-based analysis: Application to a large-scale colorectal cancer study. PLoS Genetics, 2020, 16, e1008947.	1.5	6
61	Monoallelic NTHL1 Loss-of-Function Variants and Risk of Polyposis and Colorectal Cancer. Gastroenterology, 2020, 159, 2241-2243.e6.	0.6	20
62	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.	0.4	26
63	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. BMC Medicine, 2020, 18, 396.	2.3	76
64	Quantifying the Effect of Location Matching on Accuracy of Multiparametric Magnetic Resonance Imaging Prior to Prostate Biopsyâ€"A Multicentre Study. European Urology Open Science, 2020, 20, 28-36.	0.2	1
65	Hemochromatosis risk genotype is not associated with colorectal cancer or age at its diagnosis. Human Genetics and Genomics Advances, 2020, 1, 100010.	1.0	3
66	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	1.1	26
67	Functional informed genomeâ€wide interaction analysis of body mass index, diabetes and colorectal cancer risk. Cancer Medicine, 2020, 9, 3563-3573.	1.3	7
68	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. Nature Communications, 2020, 11, 3353.	5.8	75
69	Do the risks of Lynch syndrome-related cancers depend on the parent of origin of the mutation?. Familial Cancer, 2020, 19, 215-222.	0.9	1
70	Association Between Molecular Subtypes of Colorectal Tumors and Patient Survival, Based on Pooled Analysis of 7 International Studies. Gastroenterology, 2020, 158, 2158-2168.e4.	0.6	34
71	A New Comprehensive Colorectal Cancer Risk Prediction Model Incorporating Family History, Personal Characteristics, and Environmental Factors. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 549-557.	1.1	25
72	Utility of the methylated SEPT9 test for the early detection of colorectal cancer: a systematic review and meta-analysis of diagnostic test accuracy. BMJ Open Gastroenterology, 2020, 7, e000355.	1.1	28

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73	Systematic meta-analyses, field synopsis and global assessment of the evidence of genetic association studies in colorectal cancer. Gut, 2020, 69, 1460-1471.	6.1	27
74	Potential impact of family historyâ€“based screening guidelines on the detection of earlyâ€onset colorectal cancer. Cancer, 2020, 126, 3013-3020.	2.0	45
75	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	5.8	193
76	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	3.0	129
77	Assessment of a European Bladder Cancer Predictive Model for Non-Muscle Invasive Bladder Cancer in an Australian Cohort. Bladder Cancer, 2019, 5, 31-38.	0.2	3
78	Type 2 diabetes mellitus, blood cholesterol, triglyceride and colorectal cancer risk in Lynch syndrome. British Journal of Cancer, 2019, 121, 869-876.	2.9	10
79	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
80	Ability of known susceptibility SNPs to predict colorectal cancer risk for persons with and without a family history. Familial Cancer, 2019, 18, 389-397.	0.9	23
81	Tumor mutational signatures in sebaceous skin lesions from individuals with Lynch syndrome. Molecular Genetics & Genomic Medicine, 2019, 7, e00781.	0.6	8
82	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. Nature Communications, 2019, 10, 2154.	5.8	172
83	Meeting report from the joint IARCâ€“NCI international cancer seminar series: a focus on colorectal cancer. Annals of Oncology, 2019, 30, 510-519.	0.6	42
84	Genetic variant predictors of gene expression provide new insight into risk of colorectal cancer. Human Genetics, 2019, 138, 307-326.	1.8	44
85	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. Gastroenterology, 2019, 156, 1455-1466.	0.6	111
86	Trends in Colon and Rectal Cancer Incidence in Australia from 1982 to 2014: Analysis of Data on Over 375,000 Cases. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 83-90.	1.1	81
87	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. International Journal of Epidemiology, 2019, 48, 767-780.	0.9	35
88	Risks of Colorectal Cancer and Cancer-Related Mortality in Familial Colorectal Cancer Type X and Lynch Syndrome Families. Journal of the National Cancer Institute, 2019, 111, 675-683.	3.0	12
89	Global trends in colorectal cancer mortality: projections to the year 2035. International Journal of Cancer, 2019, 144, 2992-3000.	2.3	348
90	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377

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91	Earlyâ€life exposure to sibling modifies the relationship between CD14 polymorphisms and allergic sensitization. Clinical and Experimental Allergy, 2019, 49, 331-340.	1.4	2
92	Assessing the ProMCol classifier as a prognostic marker for non-metastatic colorectal cancer within the Melbourne Collaborative Cohort Study. Gut, 2019, 68, 761-762.	6.1	2
93	Cancer screening in Australia: future directions in melanoma, Lynch syndrome, and liver, lung and prostate cancers. Public Health Research and Practice, 2019, 29, .	0.7	5
94	Discussions about predictive genetic testing for Lynch syndrome: the role of health professionals and families in decisions to decline. Familial Cancer, 2018, 17, 547-555.	0.9	8
95	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). International Journal of Epidemiology, 2018, 47, 387-388i.	0.9	40
96	Costs and outcomes of Lynch syndrome screening in the Australian colorectal cancer population. Journal of Gastroenterology and Hepatology (Australia), 2018, 33, 1737-1744.	1.4	11
97	Evaluation of the benefits, harms and costâ€effectiveness of potential alternatives to iFOBT testing for colorectal cancer screening in Australia. International Journal of Cancer, 2018, 143, 269-282.	2.3	28
98	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. American Journal of Human Genetics, 2018, 102, 904-919.	2.6	30
99	Somatic mutations of the coding microsatellites within the beta-2-microglobulin gene in mismatch repair-deficient colorectal cancers and adenomas. Familial Cancer, 2018, 17, 91-100.	0.9	21
100	Cancer risk and survival in path_MMR carriers by gene and gender up to 75 years of age: a report from the Prospective Lynch Syndrome Database. Gut, 2018, 67, 1306-1316.	6.1	410
101	Breast Cancer Risk Associations with Digital Mammographic Density by Pixel Brightness Threshold and Mammographic System. Radiology, 2018, 286, 433-442.	3.6	29
102	Genomeâ€wide association study and metaâ€analysis in Northern European populations replicate multiple colorectal cancer risk loci. International Journal of Cancer, 2018, 142, 540-546.	2.3	26
103	Associations of alcohol intake, smoking, physical activity and obesity with survival following colorectal cancer diagnosis by stage, anatomic site and tumor molecular subtype. International Journal of Cancer, 2018, 142, 238-250.	2.3	83
104	Risk of colorectal cancer for carriers of a germ-line mutation in POLE or POLD1. Genetics in Medicine, 2018, 20, 890-895.	1.1	49
105	Genomic Characterization of Upper-Tract Urothelial Carcinoma in Patients With Lynch Syndrome. JCO Precision Oncology, 2018, 2018, 1-13.	1.5	29
106	Cancer Risks for PMS2-Associated Lynch Syndrome. Journal of Clinical Oncology, 2018, 36, 2961-2968.	0.8	147
107	Revised Australian national guidelines for colorectal cancer screening: family history. Medical Journal of Australia, 2018, 209, 455-460.	0.8	35
108	Predicting interval and screen-detected breast cancers from mammographic density defined by different brightness thresholds. Breast Cancer Research, 2018, 20, 152.	2.2	24

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109	Benefits, Harms, and Cost-Effectiveness of Potential Age Extensions to the National Bowel Cancer Screening Program in Australia. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 1450-1461.	1.1	26
110	Diagnostics for Pleiotropy in Mendelian Randomization Studies: Global and Individual Tests for Direct Effects. American Journal of Epidemiology, 2018, 187, 2672-2680.	1.6	18
111	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. PLoS ONE, 2018, 13, e0196245.	1.1	9
112	The International Mismatch Repair Consortium. , 2018, , 479-495.		0
113	The melanoma genomics managing your risk study: A protocol for a randomized controlled trial evaluating the impact of personal genomic risk information on skin cancer prevention behaviors. Contemporary Clinical Trials, 2018, 70, 106-116.	0.8	19
114	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. British Journal of Cancer, 2018, 118, 1639-1647.	2.9	16
115	The use of a risk assessment and decision support tool (CRISP) compared with usual care in general practice to increase risk-stratified colorectal cancer screening: study protocol for a randomised controlled trial. Trials, 2018, 19, 397.	0.7	13
116	Sunscreen Use and Melanoma Risk Among Young Australian Adults. JAMA Dermatology, 2018, 154, 1001.	2.0	40
117	Family historyâ€“based colorectal cancer screening in Australia: A modelling study of the costs, benefits, and harms of different participation scenarios. PLoS Medicine, 2018, 15, e1002630.	3.9	6
118	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. Journal of Investigative Dermatology, 2018, 138, 2617-2624.	0.3	52
119	Interaction between polymorphisms in aspirin metabolic pathways, regular aspirin use and colorectal cancer risk: A case-control study in unselected white European populations. PLoS ONE, 2018, 13, e0192223.	1.1	5
120	Physical activity and the risk of colorectal cancer in Lynch syndrome. International Journal of Cancer, 2018, 143, 2250-2260.	2.3	23
121	DNA mismatch repair protein deficient non-neoplastic colonic crypts: a novel indicator of Lynch syndrome. Modern Pathology, 2018, 31, 1608-1618.	2.9	32
122	The Colon Cancer Family Registry Cohort. , 2018, , 427-459.		3
123	The CRISP-Q study: Communicating the risks and benefits of colorectal cancer screening. , 2018, 47, 139-144.		12
124	â€˜Why donâ€™t I need a colonoscopy?â€™ A novel approach to communicating risks and benefits of colorectal cancer screening. , 2018, 47, 343-349.		9
125	Towards personalised risk assessment and clinical management: A worldwide study of age-, sex-, geographic region-, gene- and cancer-specific risks for Lynch syndrome.. Journal of Clinical Oncology, 2018, 36, 1526-1526.	0.8	0
126	Abstract 229: Genome-wide association study by colorectal carcinoma subtype. , 2018, , .		0

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127	Abstract 245: Genetic variation related to innate immunity and colorectal cancer risk. , 2018, , .		0
128	Abstract 2965: Functionally informed genome-wide interaction analysis of nonsteroidal anti-inflammatory drugs on colorectal cancer risk. , 2018, , .		0
129	Abstract 1238: International comparison of cancer risks for Lynch syndrome. , 2018, , .		0
130	Abstract 5268: Interactions between genetic predictors of gene expression and dietary factors associated with risk of colorectal cancer. , 2018, , .		0
131	Cohort Profile: The Tasmanian Longitudinal Health STUDY (TAHS). International Journal of Epidemiology, 2017, 46, dyw028.	0.9	26
132	Mammographic density defined by higher than conventional brightness thresholds better predicts breast cancer risk. International Journal of Epidemiology, 2017, 46, dyw212.	0.9	24
133	Tumor testing to identify lynch syndrome in two Australian colorectal cancer cohorts. Journal of Gastroenterology and Hepatology (Australia), 2017, 32, 427-438.	1.4	47
134	Modeling of Successive Cancer Risks in Lynch Syndrome Families in the Presence of Competing Risks Using Copulas. Biometrics, 2017, 73, 271-282.	0.8	5
135	Comparison of the efficiency of colorectal cancer screening programs based on age and genetic risk for reduction of colorectal cancer mortality. European Journal of Human Genetics, 2017, 25, 832-838.	1.4	19
136	Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. Gut, 2017, 66, 1657-1664.	6.1	127
137	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. International Journal of Cancer, 2017, 140, 2701-2708.	2.3	76
138	Lack of evidence for germline RNF43 mutations in patients with serrated polyposis syndrome from a large multinational study. Gut, 2017, 66, 1170-1172.	6.1	42
139	Lifetime alcohol intake is associated with an increased risk of KRAS+ and BRAFâ€‘KRASâ€•but not BRAF+ colorectal cancer. International Journal of Cancer, 2017, 140, 1485-1493.	2.3	27
140	Risk factors for metachronous colorectal cancer or polyp: A systematic review and metaâ€analysis. Journal of Gastroenterology and Hepatology (Australia), 2017, 32, 301-326.	1.4	13
141	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. European Journal of Cancer, 2017, 84, 228-238.	1.3	81
142	Inherited variation in circadian rhythm genes and risks of prostate cancer and three other cancer sites in combined cancer consortia. International Journal of Cancer, 2017, 141, 1794-1802.	2.3	28
143	Choosing not to undergo predictive genetic testing for hereditary colorectal cancer syndromes: expanding our understanding of decliners and declining. Journal of Behavioral Medicine, 2017, 40, 583-594.	1.1	35
144	Testing for Gene-Environment Interactions Using a Prospective Family Cohort Design: Body Mass Index in Early and Later Adulthood and Risk of Breast Cancer. American Journal of Epidemiology, 2017, 185, 487-500.	1.6	5

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145	How does genetic risk information for Lynch syndrome translate to risk management behaviours?. Hereditary Cancer in Clinical Practice, 2017, 15, 1.	0.6	6
146	The CRISP colorectal cancer risk prediction tool: an exploratory study using simulated consultations in Australian primary care. BMC Medical Informatics and Decision Making, 2017, 17, 13.	1.5	28
147	Alcohol Consumption and the Risk of Colorectal Cancer for Mismatch Repair Gene Mutation Carriers. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 366-375.	1.1	37
148	Integrating personalised genomics into risk stratification models of population screening for colorectal cancer. Australian and New Zealand Journal of Public Health, 2017, 41, 3-4.	0.8	3
149	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 404-412.	1.1	341
150	Germline miRNA DNA variants and the risk of colorectal cancer by subtype. Genes Chromosomes and Cancer, 2017, 56, 177-184.	1.5	7
151	Enrichment of colorectal cancer associations in functional regions: Insight for using epigenomics data in the analysis of whole genome sequence-imputed GWAS data. PLoS ONE, 2017, 12, e0186518.	1.1	8
152	Colorectal cancer incidence in path_MLH1 carriers subjected to different follow-up protocols: a Prospective Lynch Syndrome Database report. Hereditary Cancer in Clinical Practice, 2017, 15, 18.	0.6	49
153	Abstract PR05: Does a comprehensive family history of colorectal cancer improve risk prediction?. , 2017, , .		3
154	Abstract B04: Development of a comprehensive colorectal cancer risk prediction tool (CRiPT) incorporating known and unknown major genes and polygenes. , 2017, , .		0
155	Abstract 4266: Double somatic mutations as a cause of tumor mismatch repair-deficiency in population-based colorectal and endometrial cancer with Lynch-like syndrome. , 2017, , .		1
156	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.	3.9	118
157	Fine-Mapping of Common Genetic Variants Associated with Colorectal Tumor Risk Identified Potential Functional Variants. PLoS ONE, 2016, 11, e0157521.	1.1	8
158	P-236 Screening practices of Australians at population and familial risk following the partial roll-out of the National Bowel Cancer Screening Program, 2009-2012. Annals of Oncology, 2016, 27, ii67.	0.6	0
159	Risk factors for metachronous colorectal cancer following a primary colorectal cancer: A prospective cohort study. International Journal of Cancer, 2016, 139, 1081-1090.	2.3	32
160	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. British Journal of Cancer, 2016, 115, 266-272.	2.9	57
161	Cholecystectomy and the risk of colorectal cancer by tumor mismatch repair deficiency status. International Journal of Colorectal Disease, 2016, 31, 1451-1457.	1.0	6
162	Colorectal cancer screening is cost-effective in the elderly who have had less intense prior screening, high baseline risk of colorectal cancer and less comorbidities. Evidence-Based Medicine, 2016, 21, 182-182.	0.6	5

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163	Multivitamin, calcium and folic acid supplements and the risk of colorectal cancer in Lynch syndrome. International Journal of Epidemiology, 2016, 45, 940-953.	0.9	27
164	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. Cancer Research, 2016, 76, 5103-5114.	0.4	100
165	Common variants in the obesity-associated genes FTO and MC4R are not associated with risk of colorectal cancer. Cancer Epidemiology, 2016, 44, 1-4.	0.8	12
166	SNP rs16906252C>T Is an Expression and Methylation Quantitative Trait Locus Associated with an Increased Risk of Developing MGMT-Methylated Colorectal Cancer. Clinical Cancer Research, 2016, 22, 6266-6277.	3.2	22
167	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in MUTYH. International Journal of Cancer, 2016, 139, 1557-1563.	2.3	107
168	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. British Journal of Cancer, 2016, 114, 221-229.	2.9	18
169	Determining the familial risk distribution of colorectal cancer: a data mining approach. Familial Cancer, 2016, 15, 241-251.	0.9	6
170	Associations of 5HTTLPR polymorphism with major depressive disorder and alcohol dependence: A systematic review and meta-analysis. Australian and New Zealand Journal of Psychiatry, 2016, 50, 842-857.	1.3	37
171	Variation at 2q35 (PNKDandTMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. Human Molecular Genetics, 2016, 25, 2349-2359.	1.4	37
172	Quantifying the utility of single nucleotide polymorphisms to guide colorectal cancer screening. Future Oncology, 2016, 12, 503-513.	1.1	42
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