

# Gadi Rennert

## List of Publications by Year in descending order

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Version: 2024-02-01

287  
papers

20,490  
citations

9786

73  
h-index

14208

128  
g-index

306  
all docs

306  
docs citations

306  
times ranked

25152  
citing authors

#	ARTICLE	IF	CITATIONS
1	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	4.4	6
2	Genome-wide interaction analysis identified low-frequency variants with sex disparity in lung cancer risk. <i>Human Molecular Genetics</i> , 2022, 31, 2831-2843.	2.9	4
3	Gene-gene interaction of AhR with and within the Wnt cascade affects susceptibility to lung cancer. <i>European Journal of Medical Research</i> , 2022, 27, 14.	2.2	1
4	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	5.0	15
5	Risk Stratification for Early-Onset Colorectal Cancer Using a Combination of Genetic and Environmental Risk Scores: An International Multi-Center Study. <i>Journal of the National Cancer Institute</i> , 2022, , .	6.3	15
6	Selective serotonin reuptake inhibitors associated with increased mortality risk in breast cancer patients in Northern Israel. <i>International Journal of Epidemiology</i> , 2022, 51, 807-816.	1.9	6
7	A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , 2022, 2, 211-219.	1.7	6
8	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1077-1089.	2.5	6
9	OUP accepted manuscript. <i>Journal of the National Cancer Institute</i> , 2022, , .	6.3	0
10	A Large-Scale Genome-Wide Gene-Gene Interaction Study of Lung Cancer Susceptibility in Europeans With a Trans-Ethnic Validation in Asians. <i>Journal of Thoracic Oncology</i> , 2022, 17, 974-990.	1.1	18
11	Association of Body Mass Index With Colorectal Cancer Risk by Genome-Wide Variants. <i>Journal of the National Cancer Institute</i> , 2021, 113, 38-47.	6.3	14
12	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	6.3	45
13	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
14	Integration of multiomic annotation data to prioritize and characterize inflammation and immune-related risk variants in squamous cell lung cancer. <i>Genetic Epidemiology</i> , 2021, 45, 99-114.	1.3	7
15	Long term follow-up of EGFR mutated NSCLC cases. <i>Translational Oncology</i> , 2021, 14, 100934.	3.7	6
16	Causal relationships between body mass index, smoking and lung cancer: Univariable and multivariable Mendelian randomization. <i>International Journal of Cancer</i> , 2021, 148, 1077-1086.	5.1	73
17	Comprehensive functional annotation of susceptibility variants identifies genetic heterogeneity between lung adenocarcinoma and squamous cell carcinoma. <i>Frontiers of Medicine</i> , 2021, 15, 275-291.	3.4	21
18	CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	6.4	5

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19	Assessing Lung Cancer Absolute Risk Trajectory Based on a Polygenic Risk Model. <i>Cancer Research</i> , 2021, 81, 1607-1615.	0.9	50
20	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. <i>American Journal of Clinical Nutrition</i> , 2021, 113, 1490-1502.	4.7	27
21	Tumor immune infiltration estimated from gene expression profiles predicts colorectal cancer relapse. <i>Oncolmmunology</i> , 2021, 10, 1862529.	4.6	9
22	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	12.8	19
23	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334.	12.1	44
24	Rare Variants in the DNA Repair Pathway and the Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 895-903.	2.5	3
25	Genome-wide association meta-analysis identifies pleiotropic risk loci for aerodigestive squamous cell cancers. <i>PLoS Genetics</i> , 2021, 17, e1009254.	3.5	19
26	Ischemic stroke and myocardial ischemia in clopidogrel users and the association with CYP2C19 loss-of-function homozygosity: a real-world study. <i>Pharmacogenomics Journal</i> , 2021, 21, 402-408.	2.0	5
27	Nongenetic Determinants of Risk for Early-Onset Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab029.	2.9	39
28	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	2.4	16
29	Translational Research in Cancer Screening: Long-Term Population-Action Bridges to Diffuse Adherence. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 7883.	2.6	0
30	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	6.2	6
31	Abstract 2737: Clinical and epidemiologic predictors of clonal immune responses in colorectal cancer. , 2021, , .		0
32	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86.	5.0	7
33	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	6.4	9
34	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
35	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	2.5	19
36	A Combined Proteomics and Mendelian Randomization Approach to Investigate the Effects of Aspirin-Targeted Proteins on Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 564-575.	2.5	10

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37	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787.	3.3	2
38	A prospective prostate cancer screening programme for men with pathogenic variants in mismatch repair genes (IMPACT): initial results from an international prospective study. <i>Lancet Oncology</i> , The, 2021, 22, 1618-1631.	10.7	48
39	Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. <i>Nutrients</i> , 2021, 13, 4164.	4.1	3
40	Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2130-2138.	5.1	13
41	Transcriptome-wide association study reveals candidate causal genes for lung cancer. <i>International Journal of Cancer</i> , 2020, 146, 1862-1878.	5.1	33
42	Genome-wide association study of INDELs identified four novel susceptibility loci associated with lung cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2855-2864.	5.1	7
43	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
44	Immune-mediated genetic pathways resulting in pulmonary function impairment increase lung cancer susceptibility. <i>Nature Communications</i> , 2020, 11, 27.	12.8	23
45	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
46	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
47	Cancer Risk After Radioactive Iodine Treatment for Hyperthyroidism: A Cohort Study. <i>Thyroid</i> , 2020, 30, 243-250.	4.5	39
48	Hormonal and reproductive factors and reduction in the risk of colorectal cancer. <i>European Journal of Cancer Prevention</i> , 2020, 29, 229-237.	1.3	3
49	Lymphocytic infiltration in stage II microsatellite stable colorectal tumors: A retrospective prognosis biomarker analysis. <i>PLoS Medicine</i> , 2020, 17, e1003292.	8.4	25
50	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1800-1808.	2.5	1
51	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	2.4	82
52	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
53	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. <i>BMC Medicine</i> , 2020, 18, 229.	5.5	28
54	Inherited Rare, Deleterious Variants in ATM Increase Lung Adenocarcinoma Risk. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1871-1879.	1.1	24

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55	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. BMC Medicine, 2020, 18, 396.	5.5	76
56	Hemochromatosis risk genotype is not associated with colorectal cancer or age at its diagnosis. Human Genetics and Genomics Advances, 2020, 1, 100010.	1.7	3
57	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. Nature Genetics, 2020, 52, 572-581.	21.4	265
58	Protein-altering germline mutations implicate novel genes related to lung cancer development. Nature Communications, 2020, 11, 2220.	12.8	31
59	Stroke and Bleeding Risks in Patients with Atrial Fibrillation Treated with Reduced Apixaban Dose: A Real-Life Study. Clinical Pharmacology and Therapeutics, 2020, 108, 1265-1273.	4.7	15
60	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	2.5	26
61	Association between myelofibrosis and risk of non-hematologic malignancies: a population-based retrospective cohort study. Annals of Hematology, 2020, 99, 1007-1016.	1.8	0
62	Functional informed genome-wide interaction analysis of body mass index, diabetes and colorectal cancer risk. Cancer Medicine, 2020, 9, 3563-3573.	2.8	7
63	Association between myelofibrosis and thromboembolism: A population-based retrospective cohort study. Journal of Thrombosis and Haemostasis, 2020, 18, 916-925.	3.8	12
64	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. Genetic Epidemiology, 2020, 44, 442-468.	1.3	32
65	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. Nature Communications, 2020, 11, 312.	12.8	30
66	Preoperative direct oral anticoagulants treatment and all-cause mortality in elderly patients with hip fracture: A retrospective cohort study. Thrombosis Research, 2020, 189, 48-54.	1.7	6
67	Association Analysis of Driver Gene-Related Genetic Variants Identified Novel Lung Cancer Susceptibility Loci with 20,871 Lung Cancer Cases and 15,971 Controls. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1423-1429.	2.5	6
68	Use of metformin and risk of breast and colorectal cancer. Diabetes Research and Clinical Practice, 2020, 165, 108232.	2.8	19
69	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	12.8	193
70	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	6.3	129
71	Lung Cancer Risk in Never-Smokers of European Descent is Associated With Genetic Variation in the 5p15.33 TERT-CLPTM1L Region. Journal of Thoracic Oncology, 2019, 14, 1360-1369.	1.1	27
72	Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524.	3.3	5

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73	Interim Results from the IMPACT Study: Evidence for Prostate-specific Antigen Screening in BRCA2 Mutation Carriers. <i>European Urology</i> , 2019, 76, 831-842.	1.9	148
74	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
75	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 935-942.	2.5	21
76	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	6.4	19
77	Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. <i>Oncotarget</i> , 2019, 10, 1760-1774.	1.8	25
78	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
79	Body mass index and the association between low-density lipoprotein cholesterol as predicted by HMGR genetic variants and breast cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 1727-1730.	1.9	3
80	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	6.4	52
81	Association of Diabetes and Glycated Hemoglobin With the Risk of Intracerebral Hemorrhage: A Population-Based Cohort Study. <i>Diabetes Care</i> , 2019, 42, 682-688.	8.6	39
82	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
83	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , 2019, 156, 1455-1466.	1.3	111
84	Red meat and processed meat intake and risk of colorectal cancer: a population-based case-control study. <i>European Journal of Cancer Prevention</i> , 2019, 28, 287-293.	1.3	14
85	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	6.2	711
86	Alcohol consumption and lung cancer risk: A pooled analysis from the International Lung Cancer Consortium and the SYNERGY study. <i>Cancer Epidemiology</i> , 2019, 58, 25-32.	1.9	22
87	Mendelian randomization analysis of C-reactive protein on colorectal cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 767-780.	1.9	35
88	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2019, 40, 432-440.	2.8	5
89	Reply to: From beta-blockers to Parkinson's disease in respect of essential tremor. <i>Movement Disorders</i> , 2019, 34, 154-154.	3.9	3
90	Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. <i>International Journal of Epidemiology</i> , 2019, 48, 751-766.	1.9	32

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91	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	21.4	377
92	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	1.9	81
93	Better Ask Than Tell: Responses to mHealth Interrogative Reminders and Associations With Colorectal Cancer Screening Subsequent Uptake in a Prospective Cohort Intervention. <i>JMIR MHealth and UHealth</i> , 2019, 7, e9351.	3.7	5
94	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
95	Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. <i>Carcinogenesis</i> , 2018, 39, 336-346.	2.8	29
96	Prostate-specific antigen velocity in a prospective prostate cancer screening study of men with genetic predisposition. <i>British Journal of Cancer</i> , 2018, 118, 266-276.	6.4	12
97	Outcomes of Chemotherapy for Microsatellite Instable "High Metastatic Colorectal Cancers. <i>JCO Precision Oncology</i> , 2018, 2, 1-10.	3.0	15
98	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. <i>Nature Communications</i> , 2018, 9, 3927.	12.8	43
99	Adrenoceptor agonists and antagonists and risk of Parkinson's disease. <i>Movement Disorders</i> , 2018, 33, 1465-1471.	3.9	76
100	Cancer-driving H3G34V/R/D mutations block H3K36 methylation and H3K36me3 "MutS" interaction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 9598-9603.	7.1	87
101	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. <i>British Journal of Cancer</i> , 2018, 118, 1639-1647.	6.4	16
102	Genetic modifiers of radon-induced lung cancer risk: a genome-wide interaction study in former uranium miners. <i>International Archives of Occupational and Environmental Health</i> , 2018, 91, 937-950.	2.3	27
103	Association of statin use with spontaneous intracerebral hemorrhage. <i>Neurology</i> , 2018, 91, e400-e409.	1.1	33
104	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018, 78, 5419-5430.	0.9	54
105	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018, 9, 3221.	12.8	60
106	Association of factor V activity with risk of venous thromboembolism and atherothrombotic cardiovascular events: A retrospective population-based cohort study. <i>Thrombosis Research</i> , 2018, 168, 14-19.	1.7	2
107	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
108	Association of atrial fibrillation and cancer: Analysis from two large population-based case-control studies. <i>PLoS ONE</i> , 2018, 13, e0190324.	2.5	30

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109	Alcohol and lung cancer risk among never smokers: A pooled analysis from the international lung cancer consortium and the SYNERGY study. <i>International Journal of Cancer</i> , 2017, 140, 1976-1984.	5.1	35
110	Factor XI deficiency is associated with lower risk for cardiovascular and venous thromboembolism events. <i>Blood</i> , 2017, 129, 1210-1215.	1.4	149
111	Catheter ablation of atrial fibrillation is associated with reduced risk of stroke and mortality: A propensity score-matched analysis. <i>Heart Rhythm</i> , 2017, 14, 635-642.	0.7	74
112	Tyrosine kinase-targeting drugs-associated heart failure. <i>British Journal of Cancer</i> , 2017, 116, 1366-1373.	6.4	28
113	Reproductive factors, hormones and colorectal cancer still unresolved. <i>British Journal of Cancer</i> , 2017, 116, 1-3.	6.4	16
114	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	21.4	472
115	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	21.4	356
116	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
117	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
118	Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 117-134.	2.5	18
119	Oral Bisphosphonates and Improved Survival of Breast Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 1684-1689.	7.0	48
120	Red cell distribution width and all-cause mortality in patients with atrial fibrillation: A cohort study. <i>Journal of Arrhythmia</i> , 2017, 33, 56-62.	1.2	14
121	Pleiotropy of genetic variants on obesity and smoking phenotypes: Results from the Oncoarray Project of The International Lung Cancer Consortium. <i>PLoS ONE</i> , 2017, 12, e0185660.	2.5	11
122	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017, 12, e0177875.	2.5	79
123	Fine-Scale Mapping at 9p22.2 Identifies Candidate Causal Variants That Modify Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS ONE</i> , 2016, 11, e0158801.	2.5	10
124	Self-rated health is prospectively associated with uptake of screening for the early detection of colorectal cancer, not vice versa. <i>European Journal of Cancer Prevention</i> , 2016, 25, 282-287.	1.3	3
125	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	5.0	31
126	Coffee Consumption and the Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 634-639.	2.5	68



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127	Usefulness of CHADS2 and CHA2DS2-VASc Scores in the Prediction of New-Onset Atrial Fibrillation: A Population-Based Study. <i>American Journal of Medicine</i> , 2016, 129, 843-849.	1.5	111
128	A Recurrent <i>ERCC3</i> Truncating Mutation Confers Moderate Risk for Breast Cancer. <i>Cancer Discovery</i> , 2016, 6, 1267-1275.	9.4	41
129	Harnessing the Questionâ€Behavior Effect to Enhance Colorectal Cancer Screening in an mHealth Experiment. <i>American Journal of Public Health</i> , 2016, 106, 1998-2004.	2.7	21
130	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
131	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
132	The role of CHADS2 and CHA2DS2â€VASc scores in the prediction of stroke in individuals without atrial fibrillation: a populationâ€based study. <i>Journal of Thrombosis and Haemostasis</i> , 2016, 14, 1155-1162.	3.8	23
133	Tumor-Infiltrating Lymphocytes, Crohnâ€™s-Like Lymphoid Reaction, and Survival From Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2016, 108, .	6.3	162
134	Treatment of infertility does not increase the risk of ovarian cancer among women with a BRCA1 or BRCA2 mutation. <i>Fertility and Sterility</i> , 2016, 105, 781-785.	1.0	38
135	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	21.4	125
136	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.	6.3	77
137	The EUROMED CANCER network: state-of-art of cancer screening programmes in non-EU Mediterranean countries. <i>European Journal of Public Health</i> , 2016, 26, 83-89.	0.3	21
138	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	1.4	18
139	A Germline Variant on Chromosome 4q31.1 Associates with Susceptibility to Developing Colon Cancer Metastasis. <i>PLoS ONE</i> , 2016, 11, e0146435.	2.5	2
140	Risk-Reducing (Chemopreventive) Agents in Breast Cancer Prevention. , 2016, , 183-212.		0
141	Handling Individuals with High Cancer Risk: One Size Doesn't Fit All. <i>Israel Medical Association Journal</i> , 2016, 18, 567-568.	0.1	0
142	Laypersons' views of material incentives for enhancing colorectal cancer screening. <i>Health Expectations</i> , 2015, 18, 1194-1203.	2.6	5
143	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2015, 17, 61.	5.0	26
144	Lower lung cancer rates in Jewish smokers in Israel and the USA. <i>International Journal of Cancer</i> , 2015, 137, 2155-2162.	5.1	2

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145	Assessing Associations between the AURKA-HMMR-TPX2-TUBG1 Functional Module and Breast Cancer Risk in BRCA1/2 Mutation Carriers. PLoS ONE, 2015, 10, e0120020.	2.5	34
146	Global implementation of genomic medicine: We are not alone. Science Translational Medicine, 2015, 7, 290ps13.	12.4	146
147	Glycated hemoglobin and risk of first episode stroke in diabetic patients with atrial fibrillation: A cohort study. Heart Rhythm, 2015, 12, 886-892.	0.7	40
148	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	21.4	221
149	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. Nature Communications, 2015, 6, 7138.	12.8	138
150	Hypothyroidism Is a Risk Factor for New-Onset Diabetes: A Cohort Study. Diabetes Care, 2015, 38, 1657-1664.	8.6	93
151	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. Human Genetics, 2015, 134, 1249-1262.	3.8	28
152	Associated Links Among Smoking, Chronic Obstructive Pulmonary Disease, and Small Cell Lung Cancer: A Pooled Analysis in the International Lung Cancer Consortium. EBioMedicine, 2015, 2, 1677-1685.	6.1	49
153	Development of APE1 enzymatic DNA repair assays: low APE1 activity is associated with increase lung cancer risk. Carcinogenesis, 2015, 36, 982-991.	2.8	24
154	Neutrophil to lymphocyte ratio and risk of a first episode of stroke in patients with atrial fibrillation: a cohort study. Journal of Thrombosis and Haemostasis, 2015, 13, 1971-1979.	3.8	42
155	Factors influencing ovulation and the risk of ovarian cancer in <sc><i>BRCA1</i></sc> and <sc><i>BRCA2</i></sc> mutation carriers. International Journal of Cancer, 2015, 137, 1136-1146.	5.1	56
156	MicroRNA Polymorphisms and Risk of Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 65-72.	2.5	11
157	The Association Between Red Cell Distribution Width and Stroke in Patients with Atrial Fibrillation. American Journal of Medicine, 2015, 128, 192.e11-192.e18.	1.5	43
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