

Kathleen A Cooney

List of Publications by Year in descending order

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157
papers

14,992
citations

41323

49
h-index

19169

118
g-index

162
all docs

162
docs citations

162
times ranked

19277
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrative Clinical Genomics of Advanced Prostate Cancer. <i>Cell</i> , 2015, 161, 1215-1228.	13.5	2,660
2	REVEL: An Ensemble Method for Predicting the Pathogenicity of Rare Missense Variants. <i>American Journal of Human Genetics</i> , 2016, 99, 877-885.	2.6	1,555
3	A common variant associated with prostate cancer in European and African populations. <i>Nature Genetics</i> , 2006, 38, 652-658.	9.4	738
4	Multiple regions within 8q24 independently affect risk for prostate cancer. <i>Nature Genetics</i> , 2007, 39, 638-644.	9.4	621
5	Admixture mapping identifies 8q24 as a prostate cancer risk locus in African-American men. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 14068-14073.	3.3	575
6	Germline Mutations in <i>HOXB13</i> and Prostate-Cancer Risk. <i>New England Journal of Medicine</i> , 2012, 366, 141-149.	13.9	566
7	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	9.4	492
8	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. <i>Nature Genetics</i> , 2009, 41, 1116-1121.	9.4	389
9	Somatic mutations in the BRCA1 gene in sporadic ovarian tumours. <i>Nature Genetics</i> , 1995, 9, 439-443.	9.4	380
10	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	9.4	265
11	Germline Mutations in ATM and BRCA1/2 Distinguish Risk for Lethal and Indolent Prostate Cancer and are Associated with Early Age at Death. <i>European Urology</i> , 2017, 71, 740-747.	0.9	256
12	Development and Validation of a Scalable Next-Generation Sequencing System for Assessing Relevant Somatic Variants in Solid Tumors. <i>Neoplasia</i> , 2015, 17, 385-399.	2.3	212
13	Prostate cancer in young men: an important clinical entity. <i>Nature Reviews Urology</i> , 2014, 11, 317-323.	1.9	206
14	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011, 43, 570-573.	9.4	198
15	Prostate Cancer Susceptibility Locus on Chromosome 1q: a Confirmatory Study. <i>Journal of the National Cancer Institute</i> , 1997, 89, 955-959.	3.0	193
16	Somatic Single Hits Inactivate the X-Linked Tumor Suppressor FOXP3 in the Prostate. <i>Cancer Cell</i> , 2009, 16, 336-346.	7.7	190
17	Global Patterns of Prostate Cancer Incidence, Aggressiveness, and Mortality in Men of African Descent. <i>Prostate Cancer</i> , 2013, 2013, 1-12.	0.4	180
18	Risk Factors for Lower Urinary Tract Symptoms in a Population-based Sample of African-American Men. <i>American Journal of Epidemiology</i> , 2003, 157, 906-914.	1.6	174

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19	Implementation of Germline Testing for Prostate Cancer: Philadelphia Prostate Cancer Consensus Conference 2019. <i>Journal of Clinical Oncology</i> , 2020, 38, 2798-2811.	0.8	170
20	Targeting Androgen Receptor and DNA Repair in Metastatic Castration-Resistant Prostate Cancer: Results From NCI 9012. <i>Journal of Clinical Oncology</i> , 2018, 36, 991-999.	0.8	169
21	HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). <i>Human Genetics</i> , 2013, 132, 5-14.	1.8	166
22	Androgen deprivation therapy for prostate cancer results in significant loss of bone density. <i>Urology</i> , 1999, 54, 607-611.	0.5	156
23	Role of Genetic Testing for Inherited Prostate Cancer Risk: Philadelphia Prostate Cancer Consensus Conference 2017. <i>Journal of Clinical Oncology</i> , 2018, 36, 414-424.	0.8	155
24	Elevated Risk of Prostate Cancer Among Men With Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2013, 31, 1713-1718.	0.8	144
25	A Combined Genomewide Linkage Scan of 1,233 Families for Prostate Cancer—Susceptibility Genes Conducted by the International Consortium for Prostate Cancer Genetics. <i>American Journal of Human Genetics</i> , 2005, 77, 219-229.	2.6	138
26	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. <i>Human Molecular Genetics</i> , 2013, 22, 408-415.	1.4	118
27	Risk of second primary tumors in men diagnosed with prostate cancer: A population-based cohort study. <i>Cancer</i> , 2014, 120, 2735-2741.	2.0	105
28	Comparison of lower urinary tract symptom severity and associated bother between community-dwelling black and white men: the Olmsted County Study of Urinary Symptoms and Health Status and the Flint Men's Health Study. <i>Urology</i> , 2003, 61, 1086-1091.	0.5	91
29	Genome-wide scan for prostate cancer susceptibility genes using families from the University of Michigan prostate cancer genetics project finds evidence for linkage on chromosome 17 nearBRCA1. <i>Prostate</i> , 2003, 57, 326-334.	1.2	90
30	Validation of Genome-Wide Prostate Cancer Associations in Men of African Descent. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 23-32.	1.1	88
31	Features of the metabolic syndrome and prostate cancer in African-American men. <i>Cancer</i> , 2007, 109, 875-881.	2.0	83
32	Sexual Behavior, Sexually Transmitted Diseases and Prostatitis: The Risk of Prostate Cancer in Black Men. <i>Journal of Urology</i> , 2006, 176, 1108-1113.	0.2	77
33	Phase II Evaluations of Cilengitide in Asymptomatic Patients with Androgen-Independent Prostate Cancer: Scientific Rationale and Study Design. <i>Clinical Genitourinary Cancer</i> , 2006, 4, 299-302.	0.9	73
34	Combined Genome-Wide Scan for Prostate Cancer Susceptibility Genes. <i>Journal of the National Cancer Institute</i> , 2004, 96, 1240-1247.	3.0	72
35	Phase II study of Cilengitide (EMD 121974, NSC 707544) in patients with non-metastatic castration resistant prostate cancer, NCI-6735. A study by the DOD/PCF prostate cancer clinical trials consortium. <i>Investigational New Drugs</i> , 2012, 30, 749-757.	1.2	72
36	Insulin-like growth factor-1, insulin-like growth factor binding protein-3, and body mass index: clinical correlates of prostate volume among Black men. <i>Urology</i> , 2002, 59, 362-367.	0.5	69

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37	Tissue Microarray Assessment of Prostate Cancer Tumor Proliferation in African- American and White Men. <i>Journal of the National Cancer Institute</i> , 2000, 92, 937-939.	3.0	68
38	Age-specific distribution of serum prostate-specific antigen in a community-based study of African-American men. <i>Urology</i> , 2001, 57, 91-96.	0.5	67
39	Early onset prostate cancer has a significant genetic component. <i>Prostate</i> , 2012, 72, 147-156.	1.2	65
40	Genetic polymorphisms in CYP17, CYP3A4, CYP19A1, SRD5A2, IGF-1, and IGFBP-3 and prostate cancer risk in African-American men: The Flint Men's Health Study. <i>Prostate</i> , 2008, 68, 296-305.	1.2	60
41	Evidence for association of SNPs in <i>ABCB1</i> and <i>CBR3</i> , but not <i>RAC2</i> , <i>NCF4</i> , <i>SLC28A3</i> or <i>TOP2B</i> , with chronic cardiotoxicity in a cohort of breast cancer patients treated with anthracyclines. <i>Pharmacogenomics</i> , 2016, 17, 231-240.	0.6	59
42	Pooled genome linkage scan of aggressive prostate cancer: results from the International Consortium for Prostate Cancer Genetics. <i>Human Genetics</i> , 2006, 120, 471-485.	1.8	57
43	A comprehensive evaluation of <i>CHEK2</i> germline mutations in men with prostate cancer. <i>Prostate</i> , 2018, 78, 607-615.	1.2	57
44	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1121-1129.	1.1	56
45	Genetic variation in Glutathione S-Transferase Omega-1, Arsenic Methyltransferase and Methylene-tetrahydrofolate Reductase, arsenic exposure and bladder cancer: a case-control study. <i>Environmental Health</i> , 2012, 11, 43.	1.7	55
46	Relationship of serum sex-steroid hormones and prostate volume in African American men. <i>Prostate</i> , 2002, 53, 322-329.	1.2	53
47	The androgen receptor CAG and GGN repeat polymorphisms and prostate cancer susceptibility in African-American men: results from the Flint Men's Health Study. <i>Journal of Human Genetics</i> , 2008, 53, 220-226.	1.1	52
48	Analysis of the Prostate Cancer Susceptibility Locus HPC20 in 172 Families Affected by Prostate Cancer. <i>American Journal of Human Genetics</i> , 2001, 68, 795-801.	2.6	51
49	Hereditary prostate cancer as a feature of Lynch Syndrome. <i>Familial Cancer</i> , 2011, 10, 37-42.	0.9	51
50	Role of the Nijmegen Breakage Syndrome 1 Gene in Familial and Sporadic Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 935-938.	1.1	49
51	Cilengitide (EMD 121974, NSC 707544) in asymptomatic metastatic castration resistant prostate cancer patients: a randomized phase II trial by the prostate cancer clinical trials consortium. <i>Investigational New Drugs</i> , 2011, 29, 1432-1440.	1.2	49
52	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.	5.1	48
53	THE NATURAL HISTORY OF LOWER URINARY TRACT SYMPTOMS IN BLACK AMERICAN MEN: RELATIONSHIPS WITH AGING, PROSTATE SIZE, FLOW RATE AND BOTHERSOMENESS. <i>Journal of Urology</i> , 2001, 165, 1521-1525.	0.2	47
54	The <i>HOXB13</i> G84E Mutation Is Associated with an Increased Risk for Prostate Cancer and Other Malignancies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1366-1372.	1.1	47

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55	Associations of prostate cancer risk variants with disease aggressiveness: results of the NCI-SPORE Genetics Working Group analysis of 18,343 cases. <i>Human Genetics</i> , 2015, 134, 439-450.	1.8	45
56	Germline genetic variants in men with prostate cancer and one or more additional cancers. <i>Cancer</i> , 2017, 123, 3925-3932.	2.0	45
57	Genome-wide linkage scan for prostate cancer aggressiveness loci using families from the University of Michigan Prostate Cancer Genetics Project. <i>Prostate</i> , 2006, 66, 173-179.	1.2	42
58	EZH2 regulates the transcription of estrogen-responsive genes through association with REA, an estrogen receptor corepressor. <i>Breast Cancer Research and Treatment</i> , 2008, 107, 235-242.	1.1	41
59	HOXB13 G84E-related Familial Prostate Cancers. <i>American Journal of Surgical Pathology</i> , 2014, 38, 615-626.	2.1	41
60	Rare Germline Pathogenic Mutations of DNA Repair Genes Are Most Strongly Associated with Grade Group 5 Prostate Cancer. <i>European Urology Oncology</i> , 2020, 3, 224-230.	2.6	41
61	Chromosome 17q12 Variants Contribute to Risk of Early-Onset Prostate Cancer. <i>Cancer Research</i> , 2008, 68, 6492-6495.	0.4	40
62	Common Variation in the <i>BRCA1</i> Gene and Prostate Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1510-1516.	1.1	37
63	Genome-wide association of familial prostate cancer cases identifies evidence for a rare segregating haplotype at 8q24.21. <i>Human Genetics</i> , 2016, 135, 923-938.	1.8	37
64	Identifying Susceptibility Genes for Prostate Cancer--A Family-Based Association Study of Polymorphisms in CYP17, CYP19, CYP11A1, and LH- β . <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 2035-2039.	1.1	36
65	Analysis of the gene coding for the BRCA2-interacting protein PALB2 in hereditary prostate cancer. <i>Prostate</i> , 2008, 68, 675-678.	1.2	36
66	Performance of Three Inherited Risk Measures for Predicting Prostate Cancer Incidence and Mortality: A Population-based Prospective Analysis. <i>European Urology</i> , 2021, 79, 419-426.	0.9	36
67	Association between Agent Orange and prostate cancer: a pilot case-control study. <i>Urology</i> , 2004, 63, 757-760.	0.5	35
68	Truncating BRCA1 Mutations Are Uncommon in a Cohort of Hereditary Prostate Cancer Families with Evidence of Linkage to 17q Markers. <i>Clinical Cancer Research</i> , 2004, 10, 5975-5980.	3.2	34
69	Risk perception and concern among brothers of men with prostate carcinoma. <i>Cancer</i> , 2004, 100, 1537-1544.	2.0	34
70	Body Composition and Serum Prostate-Specific Antigen: Review and Findings from Flint Men's Health Study. <i>Urology</i> , 2008, 71, 554-560.	0.5	33
71	Sequence variation in the mitochondrial gene cytochrome <i>c</i> oxidase subunit I and prostate cancer in African American men. <i>Prostate</i> , 2009, 69, 956-960.	1.2	32
72	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2020, 78, 316-320.	0.9	32

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73	Compelling evidence for a prostate cancer gene at 22q12.3 by the International Consortium for Prostate Cancer Genetics. <i>Human Molecular Genetics</i> , 2007, 16, 1271-1278.	1.4	31
74	Fine-mapping the putative chromosome 17q21-22 prostate cancer susceptibility gene to a 10cM region based on linkage analysis. <i>Human Genetics</i> , 2007, 121, 49-55.	1.8	30
75	Use of complementary and alternative medicine in men with family history of prostate cancer: a pilot study. <i>Urology</i> , 2004, 63, 282-287.	0.5	28
76	Two-stage Study of Familial Prostate Cancer by Whole-exome Sequencing and Custom Capture Identifies 10 Novel Genes Associated with the Risk of Prostate Cancer. <i>European Urology</i> , 2021, 79, 353-361.	0.9	28
77	Genome-wide linkage scan for prostate cancer susceptibility from the university of michigan prostate cancer genetics project: Suggestive evidence for linkage at 16q23. <i>Prostate</i> , 2009, 69, 385-391.	1.2	27
78	Risk of Prostate Cancer Associated With Familial and Hereditary Cancer Syndromes. <i>Journal of Clinical Oncology</i> , 2020, 38, 1807-1813.	0.8	27
79	Potential selection bias in a community-based study of PSA levels in African-American men. <i>Journal of Clinical Epidemiology</i> , 2001, 54, 142-148.	2.4	26
80	Genome-Wide Association Scan for Variants Associated with Early-Onset Prostate Cancer. <i>PLoS ONE</i> , 2014, 9, e93436.	1.1	25
81	Association analysis of 9,560 prostate cancer cases from the International Consortium of Prostate Cancer Genetics confirms the role of reported prostate cancer associated SNPs for familial disease. <i>Human Genetics</i> , 2014, 133, 347-356.	1.8	24
82	Chromosome 8q24 markers: Risk of early-onset and familial prostate cancer. <i>International Journal of Cancer</i> , 2008, 122, 2876-2879.	2.3	23
83	Doxorubicin-induced cardiac dysfunction in unselected patients with a history of early-stage breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 163-172.	1.1	23
84	Genome-wide linkage analysis of 1,233 prostate cancer pedigrees from the International Consortium for prostate cancer Genetics using novel sumLINK and sumLOD analyses. <i>Prostate</i> , 2010, 70, 735-744.	1.2	22
85	Comprehensive serial molecular profiling of an N of 1 exceptional non-responder with metastatic prostate cancer progressing to small cell carcinoma on treatment. <i>Journal of Hematology and Oncology</i> , 2015, 8, 109.	6.9	22
86	Genome-wide linkage scan for prostate cancer susceptibility genes in men with aggressive disease: significant evidence for linkage at chromosome 15q12. <i>Human Genetics</i> , 2006, 119, 400-407.	1.8	21
87	Validation of prostate cancer risk-related loci identified from genome-wide association studies using family-based association analysis: evidence from the International Consortium for Prostate Cancer Genetics (ICPCG). <i>Human Genetics</i> , 2012, 131, 1095-1103.	1.8	21
88	Prevalence of the HOXB13 G84E prostate cancer risk allele in men treated with radical prostatectomy. <i>BJU International</i> , 2014, 113, 830-835.	1.3	21
89	CD38 in Advanced Prostate Cancers. <i>European Urology</i> , 2021, 79, 736-746.	0.9	21
90	Clonality of sarcomatous and carcinomatous elements in sarcomatoid carcinoma of the prostate. <i>Urology</i> , 2006, 67, 423.e5-423.e8.	0.5	20

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91	Evidence for an association between prostate cancer and chromosome 8q24 and 10q11 genetic variants in African American men: The flint men's health study. <i>Prostate</i> , 2011, 71, 225-231.	1.2	20
92	Rare germline mutations in African American men diagnosed with early-onset prostate cancer. <i>Prostate</i> , 2018, 78, 321-326.	1.2	20
93	R726L androgen receptor mutation is uncommon in prostate cancer families in the united states. <i>Prostate</i> , 2003, 54, 306-309.	1.2	19
94	Identification and characterization of novel SNPs in CHEK2 in Ashkenazi Jewish men with prostate cancer. <i>Cancer Letters</i> , 2008, 270, 173-180.	3.2	19
95	Sequence variation in Î±-methylacyl-CoA racemase and risk of early-onset and familial prostate cancer. <i>Prostate</i> , 2007, 67, 1507-1513.	1.2	18
96	Bayesian inference for the stereotype regression model: Application to a case-control study of prostate cancer. <i>Statistics in Medicine</i> , 2009, 28, 3139-3157.	0.8	18
97	Identification of a novel germline SPOP mutation in a family with hereditary prostate cancer. <i>Prostate</i> , 2014, 74, 983-990.	1.2	18
98	Longitudinal changes in lower urinary tract symptoms among a cohort of black American men: The Flint Men's Health Study. <i>Urology</i> , 2004, 64, 959-965.	0.5	16
99	Hyperglycemia, Obesity, and Cancer Risks on the Horizon. <i>JAMA - Journal of the American Medical Association</i> , 2005, 293, 235.	3.8	16
100	Two-locus genome-wide linkage scan for prostate cancer susceptibility genes with an interaction effect. <i>Human Genetics</i> , 2006, 118, 716-724.	1.8	16
101	Mutational landscape of candidate genes in familial prostate cancer. <i>Prostate</i> , 2014, 74, 1371-1378.	1.2	16
102	Prostate cancer early detection practices among men with a family history of disease. <i>Urology</i> , 2003, 62, 470-475.	0.5	15
103	Identification of a novel NBN truncating mutation in a family with hereditary prostate cancer. <i>Familial Cancer</i> , 2012, 11, 595-600.	0.9	15
104	Prevalence and Correlates of Vitamin and Supplement Usage Among Men With a Family History of Prostate Cancer. <i>Integrative Cancer Therapies</i> , 2012, 11, 83-89.	0.8	14
105	Chromosomes 4 and 8 implicated in a genome wide SNP linkage scan of 762 prostate cancer families collected by the ICPCG. <i>Prostate</i> , 2012, 72, 410-426.	1.2	14
106	Association between family history of prostate and breast cancer among African-American men with prostate cancer. <i>Urology</i> , 2006, 68, 1072-1076.	0.5	13
107	Statin Use and the Risk of Recurrence After Radical Prostatectomy in a Cohort of Men With Inherited and/or Early-onset Forms of Prostate Cancer. <i>Urology</i> , 2014, 83, 1356-1361.	0.5	13
108	HOXB13 and other high penetrant genes for prostate cancer. <i>Asian Journal of Andrology</i> , 2016, 18, 530.	0.8	13

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109	The HOXB13 variant X285K is associated with clinical significance and early age at diagnosis in African American prostate cancer patients. <i>British Journal of Cancer</i> , 2022, 126, 791-796.	2.9	13
110	Testing for the Recurrent HOXB13 G84E Germline Mutation in Men with Clinical Indications for Prostate Biopsy. <i>Journal of Urology</i> , 2013, 189, 849-853.	0.2	12
111	Germline mutations in PPFIBP2 are associated with lethal prostate cancer. <i>Prostate</i> , 2018, 78, 1222-1228.	1.2	12
112	Prostate Cancer Predisposition. <i>Urologic Clinics of North America</i> , 2021, 48, 283-296.	0.8	12
113	Limitations of Prostate-specific Antigen Testing After a Prostate Cancer Diagnosis. <i>European Urology</i> , 2016, 70, 209-210.	0.9	11
114	Genetic factors influencing prostate cancer risk in Norwegian men. <i>Prostate</i> , 2018, 78, 186-192.	1.2	11
115	Germline HOXB13 G84E mutation carriers and risk to twenty common types of cancer: results from the UK Biobank. <i>British Journal of Cancer</i> , 2020, 123, 1356-1359.	2.9	11
116	Inherited Predisposition to Prostate Cancer: From Gene Discovery to Clinical Impact. <i>Transactions of the American Clinical and Climatological Association</i> , 2017, 128, 14-23.	0.9	11
117	RE: OSTEOPOROSIS AFTER ORCHIECTOMY FOR PROSTATE CANCER. <i>Journal of Urology</i> , 1998, 160, 1809-1809.	0.2	10
118	Observed evidence for guideline-recommended genes in predicting prostate cancer risk from a large population-based cohort. <i>Prostate</i> , 2021, 81, 1002-1008.	1.2	10
119	Truncating Variants in p53AIP1 Disrupting DNA Damage-Induced Apoptosis Are Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2006, 66, 10302-10307.	0.4	9
120	gsSKAT: Rapid gene set analysis and multiple testing correction for rare variant association studies using weighted linear kernels. <i>Genetic Epidemiology</i> , 2017, 41, 297-308.	0.6	9
121	Defining low-value PSA testing in a large retrospective cohort: Finding common ground between discordant guidelines. <i>Cancer Epidemiology</i> , 2018, 56, 112-117.	0.8	9
122	Family history of prostate cancer and relapse after definitive external beam radiation therapy. <i>International Journal of Radiation Oncology Biology Physics</i> , 2003, 57, 371-376.	0.4	8
123	APC I1307K and the Risk of Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 468-473.	1.1	8
124	Phase II Evaluation of Oral Estramustine, Oral Etoposide, and Intravenous Paclitaxel in Patients with Hormone-Sensitive Prostate Adenocarcinoma. <i>Clinical Genitourinary Cancer</i> , 2007, 5, 318-322.	0.9	8
125	What men want: Qualitative analysis of what men with prostate cancer (PCa) want to learn regarding genetic referral, counseling, and testing. <i>Prostate</i> , 2020, 80, 441-450.	1.2	8
126	Metastatic Dedifferentiated Chordoma With Elevated β -hCG. <i>American Journal of Clinical Oncology: Cancer Clinical Trials</i> , 2002, 25, 274-276.	0.6	7

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127	INSPstI polymorphism and prostate cancer in African-American men. <i>Prostate</i> , 2005, 65, 83-87.	1.2	7
128	Pathogenic Germline DNA Repair Gene and <i>HOXB13</i> Mutations in Men With Metastatic Prostate Cancer. <i>JCO Precision Oncology</i> , 2020, 4, 139-151.	1.5	7
129	A phase 2 trial of salvage radiation and concurrent weekly docetaxel after a rising prostate-specific antigen level after radical prostatectomy. <i>Advances in Radiation Oncology</i> , 2016, 1, 59-66.	0.6	6
130	Low-Cost Intervention to Increase Influenza Vaccination Rate at a Comprehensive Cancer Center. <i>Journal of Cancer Education</i> , 2017, 32, 871-877.	0.6	6
131	Mitochondrial alterations may underlie race-specific differences in cancer risk and outcome. <i>Journal of Clinical Investigation</i> , 2019, 129, 2187-2188.	3.9	6
132	Analysis of Xq27-28 linkage in the international consortium for prostate cancer genetics (ICPCG) families. <i>BMC Medical Genetics</i> , 2012, 13, 46.	2.1	5
133	Post hoc Analysis for Detecting Individual Rare Variant Risk Associations Using Probit Regression Bayesian Variable Selection Methods in Case-Control Sequencing Studies. <i>Genetic Epidemiology</i> , 2016, 40, 461-469.	0.6	5
134	Provider Practice Competition and Adoption of Medicare's Oncology Care Model. <i>Medical Care</i> , 2020, 58, 154-160.	1.1	5
135	Efficacy and Effect of Cabozantinib on Bone Metastases in Treatment-naive Castration-resistant Prostate Cancer. <i>Clinical Genitourinary Cancer</i> , 2020, 18, 332-339.e2.	0.9	5
136	Decreasing age at prostate cancer diagnosis over successive generations in prostate cancer families. <i>Prostate</i> , 2005, 64, 60-66.	1.2	4
137	Dose escalation of oral vinorelbine in combination with estramustine in hormone-refractory adenocarcinoma of the prostate. <i>Cancer</i> , 2006, 106, 2617-2623.	2.0	4
138	Association between Germline Variation in the FHIT Gene and Prostate Cancer in Caucasians and African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1294-1297.	1.1	4
139	Association of germline rare pathogenic mutations in guideline-recommended genes with prostate cancer progression: A meta-analysis. <i>Prostate</i> , 2022, 82, 107-119.	1.2	4
140	Assessing the Cumulative Contribution of New and Established Common Genetic Risk Factors to Early-Onset Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 766-772.	1.1	3
141	Abstract 5569: The association between statin use and risk of biochemical recurrence in men treated with radical prostatectomy in a cohort of men with inherited forms of prostate cancer. <i>Cancer Research</i> , 2011, 71, 5569-5569.	0.4	3
142	KLK3 germline mutation I179T complements DNA repair genes for predicting prostate cancer progression. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, , .	2.0	3
143	<i>HOXB13</i> mutations and prostate cancer risk. <i>BJU International</i> , 2016, 118, 496-497.	1.3	2
144	Factors associated with appropriate and low-value PSA testing. <i>Cancer Epidemiology</i> , 2020, 66, 101724.	0.8	2

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145	Semiparametric Bayesian modeling of random genetic effects in family-based association studies. <i>Statistics in Medicine</i> , 2009, 28, 113-139.	0.8	1
146	Advances in inherited cancers: Introduction. <i>Seminars in Oncology</i> , 2016, 43, 527.	0.8	1
147	Finding a Needle in the Haystack: The Search for Germline Variants Associated with Prostate Cancer Clinical Outcomes. <i>European Urology</i> , 2018, 74, 720-721.	0.9	1
148	THE NATURAL HISTORY OF LOWER URINARY TRACT SYMPTOMS IN BLACK AMERICAN MEN: RELATIONSHIPS WITH AGING, PROSTATE SIZE, FLOW RATE AND BOTHERSOMENESS. <i>Journal of Urology</i> , 2001, , 1521-1525.	0.2	1
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