

# William B Isaacs

## List of Publications by Year in descending order

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351  
papers

38,458  
citations

2795

94  
h-index

3394

183  
g-index

359  
all docs

359  
docs citations

359  
times ranked

35517  
citing authors

#	ARTICLE	IF	CITATIONS
1	AR-V7 and Resistance to Enzalutamide and Abiraterone in Prostate Cancer. <i>New England Journal of Medicine</i> , 2014, 371, 1028-1038.	13.9	2,233
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	Inflammation in prostate carcinogenesis. <i>Nature Reviews Cancer</i> , 2007, 7, 256-269.	12.8	1,352
4	The evolutionary history of lethal metastatic prostate cancer. <i>Nature</i> , 2015, 520, 353-357.	13.7	1,185
5	Prostate Cancer. <i>New England Journal of Medicine</i> , 2003, 349, 366-381.	13.9	970
6	Ligand-Independent Androgen Receptor Variants Derived from Splicing of Cryptic Exons Signify Hormone-Refractory Prostate Cancer. <i>Cancer Research</i> , 2009, 69, 16-22.	0.4	939
7	Genome-wide association study identifies a second prostate cancer susceptibility variant at 8q24. <i>Nature Genetics</i> , 2007, 39, 631-637.	9.4	818
8	Evidence for a prostate cancer susceptibility locus on the X chromosome.. <i>Nature Genetics</i> , 1998, 20, 175-179.	9.4	641
9	Copy number analysis indicates monoclonal origin of lethal metastatic prostate cancer. <i>Nature Medicine</i> , 2009, 15, 559-565.	15.2	596
10	Frequency of homozygous deletion at p16/CDKN2 in primary human tumours. <i>Nature Genetics</i> , 1995, 11, 210-212.	9.4	593
11	Cumulative Association of Five Genetic Variants with Prostate Cancer. <i>New England Journal of Medicine</i> , 2008, 358, 910-919.	13.9	589
12	Germline Mutations in <i>HOXB13</i> and Prostate-Cancer Risk. <i>New England Journal of Medicine</i> , 2012, 366, 141-149.	13.9	566
13	Androgen-induced TOP2B-mediated double-strand breaks and prostate cancer gene rearrangements. <i>Nature Genetics</i> , 2010, 42, 668-675.	9.4	539
14	Hereditary Prostate Cancer: Epidemiologic and Clinical Features. <i>Journal of Urology</i> , 1993, 150, 797-802.	0.2	519
15	Distinct Transcriptional Programs Mediated by the Ligand-Dependent Full-Length Androgen Receptor and Its Splice Variants in Castration-Resistant Prostate Cancer. <i>Cancer Research</i> , 2012, 72, 3457-3462.	0.4	518
16	Hypermethylation of CpG Islands in Primary and Metastatic Human Prostate Cancer. <i>Cancer Research</i> , 2004, 64, 1975-1986.	0.4	467
17	Tracking the clonal origin of lethal prostate cancer. <i>Journal of Clinical Investigation</i> , 2013, 123, 4918-4922.	3.9	440
18	Pathological and molecular aspects of prostate cancer. <i>Lancet, The</i> , 2003, 361, 955-964.	6.3	421

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19	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014, 46, 1103-1109.	9.4	408
20	Alpha-methylacyl-CoA racemase: a new molecular marker for prostate cancer. <i>Cancer Research</i> , 2002, 62, 2220-6.	0.4	384
21	Cyclooxygenases in cancer: progress and perspective. <i>Cancer Letters</i> , 2004, 215, 1-20.	3.2	368
22	Establishment and characterization of seven dunning rat prostatic cancer cell lines and their use in developing methods for predicting metastatic abilities of prostatic cancers. <i>Prostate</i> , 1986, 9, 261-281.	1.2	367
23	Phenotypic Analysis of Prostate-Infiltrating Lymphocytes Reveals TH17 and Treg Skewing. <i>Clinical Cancer Research</i> , 2008, 14, 3254-3261.	3.2	367
24	Nuclear MYC protein overexpression is an early alteration in human prostate carcinogenesis. <i>Modern Pathology</i> , 2008, 21, 1156-1167.	2.9	363
25	Common sequence variants on 2p15 and Xp11.22 confer susceptibility to prostate cancer. <i>Nature Genetics</i> , 2008, 40, 281-283.	9.4	357
26	Extensive transduction of nonrepetitive DNA mediated by L1 retrotransposition in cancer genomes. <i>Science</i> , 2014, 345, 1251-1343.	6.0	348
27	Prostate carcinogenesis and inflammation: emerging insights. <i>Carcinogenesis</i> , 2005, 26, 1170-1181.	1.3	330
28	The landscape of recombination in African Americans. <i>Nature</i> , 2011, 476, 170-175.	18.7	319
29	Germline mutations and sequence variants of the macrophage scavenger receptor 1 gene are associated with prostate cancer risk. <i>Nature Genetics</i> , 2002, 32, 321-325.	9.4	318
30	PTEN Protein Loss by Immunostaining: Analytic Validation and Prognostic Indicator for a High Risk Surgical Cohort of Prostate Cancer Patients. <i>Clinical Cancer Research</i> , 2011, 17, 6563-6573.	3.2	309
31	Rb Loss Is Characteristic of Prostatic Small Cell Neuroendocrine Carcinoma. <i>Clinical Cancer Research</i> , 2014, 20, 890-903.	3.2	275
32	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
33	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
34	DNA Hypomethylation Arises Later in Prostate Cancer Progression than CpG Island Hypermethylation and Contributes to Metastatic Tumor Heterogeneity. <i>Cancer Research</i> , 2008, 68, 8954-8967.	0.4	255
35	Androgen receptor outwits prostate cancer drugs. <i>Nature Medicine</i> , 2004, 10, 26-27.	15.2	242
36	Two Genome-wide Association Studies of Aggressive Prostate Cancer Implicate Putative Prostate Tumor Suppressor Gene DAB2IP. <i>Journal of the National Cancer Institute</i> , 2007, 99, 1836-1844.	3.0	235

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37	DNA Methylation Alterations Exhibit Intraindividual Stability and Interindividual Heterogeneity in Prostate Cancer Metastases. <i>Science Translational Medicine</i> , 2013, 5, 169ra10.	5.8	231
38	Human prostate cancer precursors and pathobiology. <i>Urology</i> , 2003, 62, 55-62.	0.5	229
39	Understanding the Mechanisms of Androgen Deprivation Resistance in Prostate Cancer at the Molecular Level. <i>European Urology</i> , 2015, 67, 470-479.	0.9	225
40	GSTP1 CpG Island Hypermethylation Is Responsible for the Absence of GSTP1 Expression in Human Prostate Cancer Cells. <i>American Journal of Pathology</i> , 2001, 159, 1815-1826.	1.9	219
41	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	9.4	218
42	Sequence Variants of Toll-Like Receptor 4 Are Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2004, 64, 2918-2922.	0.4	214
43	Human prostate-infiltrating CD8 <sup>+</sup> T lymphocytes are oligoclonal and PD-1 <sup>+</sup> . <i>Prostate</i> , 2009, 69, 1694-1703.	1.2	206
44	A Germline DNA Polymorphism Enhances Alternative Splicing of the KLF6 Tumor Suppressor Gene and Is Associated with Increased Prostate Cancer Risk. <i>Cancer Research</i> , 2005, 65, 1213-1222.	0.4	202
45	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
46	Sequencing of prostate cancers identifies new cancer genes, routes of progression and drug targets. <i>Nature Genetics</i> , 2018, 50, 682-692.	9.4	182
47	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
48	A snapshot of the expression signature of androgen receptor splicing variants and their distinctive transcriptional activities. <i>Prostate</i> , 2011, 71, 1656-1667.	1.2	177
49	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
50	Deletional, mutational, and methylation analyses of CDKN2 (p16/MTS1) in primary and metastatic prostate cancer. <i>Genes Chromosomes and Cancer</i> , 1997, 19, 90-96.	1.5	169
51	Sequence Variants in Toll-Like Receptor Gene Cluster (TLR6-TLR1-TLR10) and Prostate Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2005, 97, 525-532.	3.0	169
52	Detection and analysis of $\beta$ -catenin mutations in prostate cancer. <i>Prostate</i> , 2000, 45, 323-334.	1.2	167
53	A molecular analysis of prokaryotic and viral DNA sequences in prostate tissue from patients with prostate cancer indicates the presence of multiple and diverse microorganisms. <i>Prostate</i> , 2008, 68, 306-320.	1.2	167
54	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

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55	Pathological and molecular mechanisms of prostate carcinogenesis: Implications for diagnosis, detection, prevention, and treatment. <i>Journal of Cellular Biochemistry</i> , 2004, 91, 459-477.	1.2	164
56	In vitro evidence for complex modes of nuclear $\beta$ -catenin signaling during prostate growth and tumorigenesis. <i>Oncogene</i> , 2002, 21, 2679-2694.	2.6	160
57	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011, 20, 3867-3875.	1.4	160
58	Evidence for two independent prostate cancer risk-associated loci in the HNF1B gene at 17q12. <i>Nature Genetics</i> , 2008, 40, 1153-1155.	9.4	158
59	Peroxisomal branched chain fatty acid $\beta$ -oxidation pathway is upregulated in prostate cancer. <i>Prostate</i> , 2005, 63, 316-323.	1.2	155
60	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
61	CYP3A4-V and prostate cancer in African Americans: causal or confounding association because of population stratification?. <i>Human Genetics</i> , 2002, 110, 553-560.	1.8	152
62	Ligand-dependent inhibition of $\beta$ -catenin/TCF signaling by androgen receptor. <i>Oncogene</i> , 2002, 21, 8453-8469.	2.6	144
63	DNA copy number alterations in prostate cancers: A combined analysis of published CGH studies. <i>Prostate</i> , 2007, 67, 692-700.	1.2	141
64	Carbohydrate restriction, prostate cancer growth, and the insulin-like growth factor axis. <i>Prostate</i> , 2008, 68, 11-19.	1.2	140
65	Germline DNA-repair Gene Mutations and Outcomes in Men with Metastatic Castration-resistant Prostate Cancer Receiving First-line Abiraterone and Enzalutamide. <i>European Urology</i> , 2018, 74, 218-225.	0.9	140
66	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
67	Germline Mutations in ATM and BRCA1/2 Are Associated with Grade Reclassification in Men on Active Surveillance for Prostate Cancer. <i>European Urology</i> , 2019, 75, 743-749.	0.9	138
68	Linkage and Association Studies of Prostate Cancer Susceptibility: Evidence for Linkage at 8p22-23. <i>American Journal of Human Genetics</i> , 2001, 69, 341-350.	2.6	137
69	A Novel Role of Myosin VI in Human Prostate Cancer. <i>American Journal of Pathology</i> , 2006, 169, 1843-1854.	1.9	133
70	Effects of RNase L mutations associated with prostate cancer on apoptosis induced by 2',5'-oligoadenylates. <i>Cancer Research</i> , 2003, 63, 6795-801.	0.4	133
71	Human polymorphisms at long non-coding RNAs (lncRNAs) and association with prostate cancer risk. <i>Carcinogenesis</i> , 2011, 32, 1655-1659.	1.3	132
72	Macrophage Inhibitory Cytokine 1: A New Prognostic Marker in Prostate Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 6658-6664.	3.2	129

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73	Association Between Two Unlinked Loci at 8q24 and Prostate Cancer Risk Among European Americans. <i>Journal of the National Cancer Institute</i> , 2007, 99, 1525-1533.	3.0	126
74	Acute inflammatory proteins constitute the organic matrix of prostatic corpora amylacea and calculi in men with prostate cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3443-3448.	3.3	124
75	Allelic loss of the retinoblastoma gene in primary human prostatic adenocarcinomas. <i>Prostate</i> , 1995, 26, 35-39.	1.2	123
76	MSH2 Loss in Primary Prostate Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 6863-6874.	3.2	122
77	Loss of PTEN Is Associated with Aggressive Behavior in ERG-Positive Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 2333-2344.	1.1	121
78	Î±-Methylacyl-CoA Racemase. <i>American Journal of Surgical Pathology</i> , 2003, 27, 1128-1133.	2.1	120
79	Gene expression signature of benign prostatic hyperplasia revealed by cDNA microarray analysis. <i>Prostate</i> , 2002, 51, 189-200.	1.2	119
80	Associations between hOGG1 sequence variants and prostate cancer susceptibility. <i>Cancer Research</i> , 2002, 62, 2253-7.	0.4	119
81	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
82	Explaining racial differences in prostate cancer in the United States: Sociology or biology?. <i>Prostate</i> , 2005, 62, 243-252.	1.2	117
83	Association of <i>IL10</i> and Other immune response and obesity-related genes with prostate cancer in CLUE II. <i>Prostate</i> , 2009, 69, 874-885.	1.2	117
84	Polygenic Risk Score Improves Prostate Cancer Risk Prediction: Results from the Stockholm-1 Cohort Study. <i>European Urology</i> , 2011, 60, 21-28.	0.9	117
85	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. <i>PLoS Genetics</i> , 2011, 7, e1001387.	1.5	117
86	Structure and Methylation-Associated Silencing of a Gene within a Homozygously Deleted Region of Human Chromosome Band 8p22. <i>Genomics</i> , 1996, 35, 55-65.	1.3	114
87	Decreased gene expression of steroid 5 alpha-reductase 2 in human prostate cancer: Implications for finasteride therapy of prostate carcinoma. <i>Prostate</i> , 2003, 57, 134-139.	1.2	111
88	H6D Polymorphism in Macrophage-Inhibitory Cytokine-1 Gene Associated With Prostate Cancer. <i>Journal of the National Cancer Institute</i> , 2004, 96, 1248-1254.	3.0	111
89	Sequence Variants at 22q13 Are Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2009, 69, 10-15.	0.4	109
90	Intraductal/ductal histology and lymphovascular invasion are associated with germline DNA repair gene mutations in prostate cancer. <i>Prostate</i> , 2018, 78, 401-407.	1.2	105

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91	Physical Mapping of Chromosome 8p22 Markers and Their Homozygous Deletion in a Metastatic Prostate Cancer. <i>Genomics</i> , 1996, 35, 46-54.	1.3	104
92	Fine mapping association study and functional analysis implicate a SNP in MSMB at 10q11 as a causal variant for prostate cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1368-1375.	1.4	103
93	Inherited genetic variant predisposes to aggressive but not indolent prostate cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 2136-2140.	3.3	100
94	Alpha-methylacyl-CoA racemase as an androgen-independent growth modifier in prostate cancer. <i>Cancer Research</i> , 2003, 63, 7365-76.	0.4	100
95	Homozygous Deletions and Recurrent Amplifications Implicate New Genes Involved in Prostate Cancer. <i>Neoplasia</i> , 2008, 10, 897-IN37.	2.3	99
96	COX-2 gene promoter haplotypes and prostate cancer risk. <i>Carcinogenesis</i> , 2004, 25, 961-966.	1.3	95
97	Common Sequence Variants of the Macrophage Scavenger Receptor 1 Gene Are Associated with Prostate Cancer Risk. <i>American Journal of Human Genetics</i> , 2003, 72, 208-212.	2.6	94
98	Evaluation of Linkage and Association of HPC2/ELAC2 in Patients with Familial or Sporadic Prostate Cancer. <i>American Journal of Human Genetics</i> , 2001, 68, 901-911.	2.6	93
99	Individual and cumulative effect of prostate cancer risk-associated variants on clinicopathologic variables in 5,895 prostate cancer patients. <i>Prostate</i> , 2009, 69, 1195-1205.	1.2	93
100	Modulation of CXCL14 (BRAK) expression in prostate cancer. <i>Prostate</i> , 2005, 64, 67-74.	1.2	92
101	The Role of Genetic Markers in the Management of Prostate Cancer. <i>European Urology</i> , 2012, 62, 577-587.	0.9	92
102	DIAPH3 governs the cellular transition to the amoeboid tumour phenotype. <i>EMBO Molecular Medicine</i> , 2012, 4, 743-760.	3.3	92
103	Assembly of Inflammation-Related Genes for Pathway-Focused Genetic Analysis. <i>PLoS ONE</i> , 2007, 2, e1035.	1.1	89
104	A Polymorphism in the CDKN1B Gene Is Associated with Increased Risk of Hereditary Prostate Cancer. <i>Cancer Research</i> , 2004, 64, 1997-1999.	0.4	88
105	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
106	Linkage of prostate cancer susceptibility loci to chromosome 1. <i>Human Genetics</i> , 2001, 108, 335-345.	1.8	86
107	Trefoil factor 3 overexpression in prostatic carcinoma: Prognostic importance using tissue microarrays. <i>Prostate</i> , 2004, 61, 215-227.	1.2	85
108	Potential Impact of Adding Genetic Markers to Clinical Parameters in Predicting Prostate Biopsy Outcomes in Men Following an Initial Negative Biopsy: Findings from the REDUCE Trial. <i>European Urology</i> , 2012, 62, 953-961.	0.9	85

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109	Focus on prostate cancer. <i>Cancer Cell</i> , 2002, 2, 113-116.	7.7	83
110	Relation between aberrant $\beta$ -catenin expression and loss of E-cadherin function in prostate cancer. , 1997, 74, 374-377.		82
111	Genetic markers associated with early cancer-specific mortality following prostatectomy. <i>Cancer</i> , 2013, 119, 2405-2412.	2.0	81
112	Molecular and cellular changes associated with the acquisition of metastatic ability by prostatic cancer cells. <i>Prostate</i> , 1994, 25, 249-265.	1.2	79
113	Genome-wide association study identifies a new locus JMJD1C at 10q21 that may influence serum androgen levels in men. <i>Human Molecular Genetics</i> , 2012, 21, 5222-5228.	1.4	79
114	Acne and risk of prostate cancer. <i>International Journal of Cancer</i> , 2007, 121, 2688-2692.	2.3	78
115	Evaluation of Serum and Seminal Plasma Markers in the Diagnosis of Canine Prostatic Disorders. <i>Journal of Veterinary Internal Medicine</i> , 1995, 9, 149-153.	0.6	77
116	Estimation of absolute risk for prostate cancer using genetic markers and family history. <i>Prostate</i> , 2009, 69, 1565-1572.	1.2	76
117	Frequent Loss of Chromosome Arms 8p and 13q in Collecting Duct Carcinoma (CDC) of the Kidney. <i>Genes Chromosomes and Cancer</i> , 1995, 12, 76-80.	1.5	75
118	Genome-wide scan for prostate cancer susceptibility genes in the Johns Hopkins hereditary prostate cancer families. <i>Prostate</i> , 2003, 57, 320-325.	1.2	75
119	Phenotypic characterization of telomerase-immortalized primary non-malignant and malignant tumor-derived human prostate epithelial cell lines. <i>Experimental Cell Research</i> , 2006, 312, 831-843.	1.2	75
120	In Swedish Families with Hereditary Prostate Cancer, Linkage to the HPC1 Locus on Chromosome 1q24-25 Is Restricted to Families with Early-Onset Prostate Cancer. <i>American Journal of Human Genetics</i> , 1999, 65, 134-140.	2.6	73
121	Association of a Germ-Line Copy Number Variation at 2p24.3 and Risk for Aggressive Prostate Cancer. <i>Cancer Research</i> , 2009, 69, 2176-2179.	0.4	73
122	Combined Genome-Wide Scan for Prostate Cancer Susceptibility Genes. <i>Journal of the National Cancer Institute</i> , 2004, 96, 1240-1247.	3.0	72
123	XMRV: A New Virus in Prostate Cancer?. <i>Cancer Research</i> , 2010, 70, 10028-10033.	0.4	72
124	Molecular advances in prostate cancer. <i>Current Opinion in Oncology</i> , 1997, 9, 101-107.	1.1	71
125	Polymorphic GGC repeats in the androgen receptor gene are associated with hereditary and sporadic prostate cancer risk. <i>Human Genetics</i> , 2002, 110, 122-129.	1.8	71
126	Identification of miR-30b-3p and miR-30d-5p as direct regulators of androgen receptor signaling in prostate cancer by complementary functional microRNA library screening. <i>Oncotarget</i> , 2016, 7, 72593-72607.	0.8	71



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127	BIOLOGICAL AGGRESSIVENESS OF HEREDITARY PROSTATE CANCER: LONG-TERM EVALUATION FOLLOWING RADICAL PROSTATECTOMY. <i>Journal of Urology</i> , 1998, 160, 660-663.	0.2	69
128	GOLPH2 and MYO6: Putative prostate cancer markers localized to the Golgi apparatus. <i>Prostate</i> , 2008, 68, 1387-1395.	1.2	69
129	Leveraging population admixture to characterize the heritability of complex traits. <i>Nature Genetics</i> , 2014, 46, 1356-1362.	9.4	69
130	Genome-wide Scan of 29,141 African Americans Finds No Evidence of Directional Selection since Admixture. <i>American Journal of Human Genetics</i> , 2014, 95, 437-444.	2.6	69
131	Genetic Variants in the <i>LEPR</i> , <i>CRY1</i> , <i>RNASEL</i> , <i>IL4</i> , and <i>ARVCF</i> Genes Are Prognostic Markers of Prostate Cancer-Specific Mortality. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1928-1936.	1.1	68
132	Identification of Aryl Hydrocarbon Receptor as a Putative Wnt/ $\beta$ -Catenin Pathway Target Gene in Prostate Cancer Cells. <i>Cancer Research</i> , 2004, 64, 2523-2533.	0.4	66
133	The Effects of Basic Fibroblast Growth Factor and suramin on Cell Motility and Growth of Rat Prostate Cancer Cells. <i>Journal of Urology</i> , 1991, 145, 199-202.	0.2	65
134	Titin, a huge, elastic sarcomeric protein with a probable role in morphogenesis. <i>BioEssays</i> , 1991, 13, 157-161.	1.2	65
135	Identification of New Differentially Methylated Genes That Have Potential Functional Consequences in Prostate Cancer. <i>PLoS ONE</i> , 2012, 7, e48455.	1.1	65
136	Systematic replication study of reported genetic associations in prostate cancer: Strong support for genetic variation in the androgen pathway. <i>Prostate</i> , 2006, 66, 1729-1743.	1.2	64
137	Comprehensive assessment of DNA copy number alterations in human prostate cancers using Affymetrix 100K SNP mapping array. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 1018-1032.	1.5	64
138	Constitutively active androgen receptor splice variants AR-V3, AR-V7 and AR-V9 are co-expressed in castration-resistant prostate cancer metastases. <i>British Journal of Cancer</i> , 2018, 119, 347-356.	2.9	63
139	Joint effect of HSD3B1 and HSD3B2 genes is associated with hereditary and sporadic prostate cancer susceptibility. <i>Cancer Research</i> , 2002, 62, 1784-9.	0.4	63
140	Genomic Organization of the Human KAI1 Metastasis-Suppressor Gene. <i>Genomics</i> , 1997, 41, 25-32.	1.3	62
141	Deletion of a Small Consensus Region at 6q15, Including the <i>MAP3K7</i> Gene, Is Significantly Associated with High-Grade Prostate Cancers. <i>Clinical Cancer Research</i> , 2007, 13, 5028-5033.	3.2	62
142	Monocyte chemotactic protein-1 (MCP-1/CCL2) is associated with prostatic growth dysregulation and benign prostatic hyperplasia. <i>Prostate</i> , 2010, 70, 473-481.	1.2	62
143	Generalizability of established prostate cancer risk variants in men of African ancestry. <i>International Journal of Cancer</i> , 2015, 136, 1210-1217.	2.3	62
144	Analytic, Preanalytic, and Clinical Validation of p53 IHC for Detection of <i>TP53</i> Missense Mutation in Prostate Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 4693-4703.	3.2	62

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145	Association of Prostate Cancer Risk Variants with Clinicopathologic Characteristics of the Disease. <i>Clinical Cancer Research</i> , 2008, 14, 5819-5824.	3.2	61
146	Prostate cancer risk-associated variants reported from genome-wide association studies: Meta-analysis and their contribution to genetic Variation. <i>Prostate</i> , 2010, 70, 1729-1738.	1.2	61
147	Homologous recombination deficiency (HRD) score in germline BRCA2- versus ATM-altered prostate cancer. <i>Modern Pathology</i> , 2021, 34, 1185-1193.	2.9	61
148	Stronger Association between Obesity and Biochemical Progression after Radical Prostatectomy among Men Treated in the Last 10 Years. <i>Clinical Cancer Research</i> , 2005, 11, 2883-2888.	3.2	60
149	Sexually Transmitted Infections and Prostatic Inflammation/Cell Damage as Measured by Serum Prostate Specific Antigen Concentration. <i>Journal of Urology</i> , 2006, 175, 1937-1942.	0.2	60
150	Adding genetic risk score to family history identifies twice as many high-risk men for prostate cancer: Results from the prostate cancer prevention trial. <i>Prostate</i> , 2016, 76, 1120-1129.	1.2	60
151	Role of androgen receptor splice variant-7 (AR-V7) in prostate cancer resistance to 2nd-generation androgen receptor signaling inhibitors. <i>Oncogene</i> , 2020, 39, 6935-6949.	2.6	60
152	Prostate Cancer Risk Associated Loci in African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2145-2149.	1.1	57
153	Increased gene copy number of ERG on chromosome 21 but not TMPRSS2-ERG fusion predicts outcome in prostatic adenocarcinomas. <i>Modern Pathology</i> , 2011, 24, 1511-1520.	2.9	57
154	A comprehensive evaluation of CHEK2 germline mutations in men with prostate cancer. <i>Prostate</i> , 2018, 78, 607-615.	1.2	57
155	VITAMIN D RECEPTOR POLYMORPHISMS AND LETHAL PROSTATE CANCER. <i>Journal of Urology</i> , 1998, 160, 1405-1409.	0.2	56
156	Polymorphisms in the CYP1A1 gene are associated with prostate cancer risk. <i>International Journal of Cancer</i> , 2003, 106, 375-378.	2.3	56
157	Cyclin D1 Loss Distinguishes Prostatic Small-Cell Carcinoma from Most Prostatic Adenocarcinomas. <i>Clinical Cancer Research</i> , 2015, 21, 5619-5629.	3.2	56
158	LOSS OF HETEROZYGOSITY AT 12P12-13 IN PRIMARY AND METASTATIC PROSTATE ADENOCARCINOMA. <i>Journal of Urology</i> , 2000, 164, 192-196.	0.2	55
159	Evidence for a prostate cancer linkage to chromosome 20 in 159 hereditary prostate cancer families. <i>Human Genetics</i> , 2001, 108, 430-435.	1.8	53
160	Design, Synthesis, and In Vitro Testing of Î±-Methylacyl-CoA Racemase Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2007, 50, 2700-2707.	2.9	52
161	Association of 17 prostate cancer susceptibility loci with prostate cancer risk in Chinese men. <i>Prostate</i> , 2010, 70, 425-432.	1.2	52
162	Cytokine profiling of prostatic fluid from cancerous prostate glands identifies cytokines associated with extent of tumor and inflammation. <i>Prostate</i> , 2008, 68, 872-882.	1.2	51

#	ARTICLE	IF	CITATIONS
163	Endoglin (CD105) as a urinary and serum marker of prostate cancer. <i>International Journal of Cancer</i> , 2009, 124, 664-669.	2.3	51
164	Evaluation of PPP2R2A as a prostate cancer susceptibility gene: a comprehensive germline and somatic study. <i>Cancer Genetics</i> , 2011, 204, 375-381.	0.2	51
165	Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. <i>Nature Communications</i> , 2015, 6, 8469.	5.8	51
166	A Novel Prostate Cancer Susceptibility Locus at 19q13. <i>Cancer Research</i> , 2009, 69, 2720-2723.	0.4	50
167	The G84E mutation of HOXB13 is associated with increased risk for prostate cancer: results from the REDUCE trial. <i>Carcinogenesis</i> , 2013, 34, 1260-1264.	1.3	50
168	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , 2015, 24, 5603-5618.	1.4	50
169	No evidence for a role of BRCA1 or BRCA2 mutations in Ashkenazi Jewish families with hereditary prostate cancer. <i>Cancer Research</i> , 1999, 59, 280-284.		49
170	Variation in IL10 and Other Genes Involved in the Immune Response and in Oxidation and Prostate Cancer Recurrence. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1774-1782.	1.1	49
171	Screening for familial and hereditary prostate cancer. <i>International Journal of Cancer</i> , 2016, 138, 2579-2591.	2.3	49
172	High mobility group protein I(Y): a candidate architectural protein for chromosomal rearrangements in prostate cancer cells. <i>Cancer Research</i> , 2002, 62, 647-51.	0.4	49
173	Interleukin-2 transfected prostate cancer cells generate a local antitumor effect in vivo. <i>Prostate</i> , 1994, 24, 244-251.	1.2	48
174	Linkage and association of CYP17 gene in hereditary and sporadic prostate cancer. <i>International Journal of Cancer</i> , 2001, 95, 354-359.	2.3	48
175	Germ-Line Mutation of NKX3.1 Cosegregates with Hereditary Prostate Cancer and Alters the Homeodomain Structure and Function. <i>Cancer Research</i> , 2006, 66, 69-77.	0.4	48
176	TMPRSS2-ERG gene fusion status in minute (minimal) prostatic adenocarcinoma. <i>Modern Pathology</i> , 2009, 22, 1415-1422.	2.9	48
177	A comprehensive association study for genes in inflammation pathway provides support for their roles in prostate cancer risk in the CAPS study. <i>Prostate</i> , 2006, 66, 1556-1564.	1.2	47
178	Cumulative effect of five genetic variants on prostate cancer risk in multiple study populations. <i>Prostate</i> , 2008, 68, 1257-1262.	1.2	47
179	The HOXB13 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
180	Genetic variability in inflammation pathways and prostate cancer risk. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2007, 25, 250-259.	0.8	46

#	ARTICLE	IF	CITATIONS
181	High-Throughput Screen Identifies Novel Inhibitors of Cancer Biomarker $\beta$ -Methylacyl Coenzyme A Racemase (AMACR/P504S). <i>Molecular Cancer Therapeutics</i> , 2011, 10, 825-838.	1.9	46
182	Mannose Receptor $\alpha$ 4 positive Macrophage Infiltration Correlates with Prostate Cancer Onset and Metastatic Castration-resistant Disease. <i>European Urology Oncology</i> , 2019, 2, 429-436.	2.6	46
183	Looking Beyond Morphology: Cancer Gene Expression Profiling Using DNA Microarrays. <i>Cancer Investigation</i> , 2003, 21, 937-949.	0.6	45
184	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
185	Prostate Cancer Predisposition Loci and Risk of Metastatic Disease and Prostate Cancer Recurrence. <i>Clinical Cancer Research</i> , 2011, 17, 1075-1081.	3.2	44
186	Assignment of the Human $\beta$ -Catenin Gene (CTNNA1) to Chromosome 5q21-q22. <i>Genomics</i> , 1994, 19, 188-190.	1.3	43
187	Genome-wide screen for prostate cancer susceptibility genes in men with clinically significant disease. <i>Prostate</i> , 2005, 64, 356-361.	1.2	43
188	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. <i>Human Molecular Genetics</i> , 2011, 20, 2869-2878.	1.4	43
189	Sequence variants of alpha-methylacyl-CoA racemase are associated with prostate cancer risk. <i>Cancer Research</i> , 2002, 62, 6485-8.	0.4	43
190	Deletion mapping at 12p12-13 in metastatic prostate cancer. , 1999, 25, 270-276.		42
191	Immunomodulatory IL-18 binding protein is produced by prostate cancer cells and its levels in urine and serum correlate with tumor status. <i>International Journal of Cancer</i> , 2011, 129, 424-432.	2.3	42
192	A Genome-Wide Assessment of Variability in Human Serum Metabolism. <i>Human Mutation</i> , 2013, 34, 515-524.	1.1	42
193	Infections and inflammation in prostate cancer. <i>American Journal of Clinical and Experimental Urology</i> , 2013, 1, 3-11.	0.4	42
194	Multiple genomic alterations on 21q22 predict various <i>TMPRSS2/ERG</i> fusion transcripts in human prostate cancers. <i>Genes Chromosomes and Cancer</i> , 2007, 46, 972-980.	1.5	41
195	Association between sequence variants at 17q12 and 17q24.3 and prostate cancer risk in European and African Americans. <i>Prostate</i> , 2008, 68, 691-697.	1.2	41
196	Nucleotide resolution analysis of <i>TMPRSS2</i> and <i>ERG</i> rearrangements in prostate cancer. <i>Journal of Pathology</i> , 2013, 230, 174-183.	2.1	41
197	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
198	Inflammation, Microbiota, and Prostate Cancer. <i>European Urology Focus</i> , 2016, 2, 374-382.	1.6	40

#	ARTICLE	IF	CITATIONS
199	HOXB13 interaction with MEIS1 modifies proliferation and gene expression in prostate cancer. <i>Prostate</i> , 2019, 79, 414-424.	1.2	39
200	Dynamic structure of the SPANX gene cluster mapped to the prostate cancer susceptibility locus HPCX at Xq27. <i>Genome Research</i> , 2005, 15, 1477-1486.	2.4	38
201	An evaluation of PCR primer sets used for detection of <i>Propionibacterium acnes</i> in prostate tissue samples. <i>Prostate</i> , 2008, 68, 1492-1495.	1.2	38
202	Inherited genetic markers discovered to date are able to identify a significant number of men at considerably elevated risk for prostate cancer. <i>Prostate</i> , 2011, 71, 421-430.	1.2	38
203	A genome-wide search for loci interacting with known prostate cancer risk-associated genetic variants. <i>Carcinogenesis</i> , 2012, 33, 598-603.	1.3	38
204	The expression of AURKA is androgen regulated in castration-resistant prostate cancer. <i>Scientific Reports</i> , 2017, 7, 17978.	1.6	38
205	Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. <i>European Urology Oncology</i> , 2021, 4, 570-579.	2.6	38
206	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
207	Meta-analysis of association of rare mutations and common sequence variants in the MSR1 gene and prostate cancer risk. <i>Prostate</i> , 2006, 66, 728-737.	1.2	36
208	Chromosome 8q24 risk variants in hereditary and non-hereditary prostate cancer patients. <i>Prostate</i> , 2008, 68, 489-497.	1.2	36
209	Association of reported prostate cancer risk alleles with PSA levels among men without a diagnosis of prostate cancer. <i>Prostate</i> , 2009, 69, 419-427.	1.2	36
210	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
211	Methylation and mutational analysis of p27kip1 in prostate carcinoma. <i>Prostate</i> , 2001, 48, 248-253.	1.2	35
212	Genetic Variants and Family History Predict Prostate Cancer Similar to Prostate-Specific Antigen. <i>Clinical Cancer Research</i> , 2009, 15, 1105-1111.	3.2	35
213	Sequence variation within the 5' regulatory regions of the vitamin D binding protein and receptor genes and prostate cancer risk. <i>Prostate</i> , 2005, 64, 272-282.	1.2	34
214	Appraising the relevance of DNA copy number loss and gain in prostate cancer using whole genome DNA sequence data. <i>PLoS Genetics</i> , 2017, 13, e1007001.	1.5	34
215	Genome-wide copy-number variation analysis identifies common genetic variants at 20p13 associated with aggressiveness of prostate cancer. <i>Carcinogenesis</i> , 2011, 32, 1057-1062.	1.3	33
216	Integration of Somatic Deletion Analysis of Prostate Cancers and Germline Linkage Analysis of Prostate Cancer Families Reveals Two Small Consensus Regions for Prostate Cancer Genes at 8p. <i>Cancer Research</i> , 2007, 67, 4098-4103.	0.4	32

#	ARTICLE	IF	CITATIONS
217	Genetic and epigenetic inactivation of <i>LPL</i> gene in human prostate cancer. <i>International Journal of Cancer</i> , 2009, 124, 734-738.	2.3	32
218	A systematic comparison of exercise training protocols on animal models of cardiovascular capacity. <i>Life Sciences</i> , 2019, 217, 128-140.	2.0	32
219	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
220	Molecular genetics and chromosomal alterations in prostate cancer. <i>Cancer</i> , 1995, 75, 2004-2012.	2.0	31
221	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
222	A major locus for hereditary prostate cancer in Finland: localization by linkage disequilibrium of a haplotype in the HPCX region. <i>Human Genetics</i> , 2005, 117, 307-316.	1.8	30
223	Fine-mapping the putative chromosome 17q21 prostate cancer susceptibility gene to a 10cM region based on linkage analysis. <i>Human Genetics</i> , 2007, 121, 49-55.	1.8	30
224	DNA methylation, molecular genetic, and linkage studies in prostate cancer. <i>Prostate</i> , 1996, 29, 36-44.	1.2	29
225	Comprehensive genetic evaluation of common E-cadherin sequence variants and prostate cancer risk: strong confirmation of functional promoter SNP. <i>Human Genetics</i> , 2005, 118, 339-347.	1.8	29
226	Germline Variants in Asporin Vary by Race, Modulate the Tumor Microenvironment, and Are Differentially Associated with Metastatic Prostate Cancer. <i>Clinical Cancer Research</i> , 2016, 22, 448-458.	3.2	29
227	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , 2011, 20, 3322-3329.	1.4	28
228	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
229	Evaluation of Association of HNF1B Variants with Diverse Cancers: Collaborative Analysis of Data from 19 Genome-Wide Association Studies. <i>PLoS ONE</i> , 2010, 5, e10858.	1.1	28
230	Telomere length as a risk factor for hereditary prostate cancer. <i>Prostate</i> , 2014, 74, 359-364.	1.2	27
231	Familial aggregation of bothersome benign prostatic hyperplasia symptoms. <i>Urology</i> , 2003, 61, 781-785.	0.5	26
232	Refining the Prostate Cancer Genetic Association within the <i>JAZF1</i> Gene on Chromosome 7p15.2. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1349-1355.	1.1	26
233	DNA-Repair Gene Mutations in Metastatic Prostate Cancer. <i>New England Journal of Medicine</i> , 2016, 375, 1802-1805.	13.9	26
234	Genetic and epigenetic inactivation of <i>TNFRSF10C</i> in human prostate cancer. <i>Prostate</i> , 2009, 69, 327-335.	1.2	25

#	ARTICLE	IF	CITATIONS
235	Functional annotation of risk loci identified through genome-wide association studies for prostate cancer. <i>Prostate</i> , 2011, 71, 955-963.	1.2	25
236	Genome-Wide Association Scan for Variants Associated with Early-Onset Prostate Cancer. <i>PLoS ONE</i> , 2014, 9, e93436.	1.1	25
237	Variation in genes involved in the immune response and prostate cancer risk in the placebo arm of the Prostate Cancer Prevention Trial. <i>Prostate</i> , 2015, 75, 1403-1418.	1.2	25
238	Physical and Transcript Map of the Hereditary Prostate Cancer Region at Xq27. <i>Genomics</i> , 2002, 79, 41-50.	1.3	24
239	Evaluation of SRD5A2 sequence variants in susceptibility to hereditary and sporadic prostate cancer. <i>Prostate</i> , 2003, 56, 37-44.	1.2	24
240	Improved Biomarkers for Prostate Cancer: A Definite Need. <i>Journal of the National Cancer Institute</i> , 2004, 96, 813-815.	3.0	24
241	RNASEL Arg462Gln polymorphism and prostate cancer in PLCO. <i>Prostate</i> , 2007, 67, 849-854.	1.2	24
242	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
243	Integrated clinical, whole-genome, and transcriptome analysis of multisampled lethal metastatic prostate cancer. <i>Journal of Physical Education and Sports Management</i> , 2016, 2, a000752.	0.5	24
244	Effect of pentosan, a novel cancer chemotherapeutic agent, on prostate cancer cell growth and motility. <i>Prostate</i> , 1992, 20, 233-241.	1.2	23
245	Identification of a prostate cancer susceptibility locus on chromosome 7q11.21 in Jewish families. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 1939-1944.	3.3	23
246	$\beta$ -Catenin overrides Src-dependent activation of $\beta$ -catenin oncogenic signaling. <i>Molecular Cancer Therapeutics</i> , 2008, 7, 1386-1397.	1.9	23
247	Validation of a prostate cancer polygenic risk score. <i>Prostate</i> , 2020, 80, 1314-1321.	1.2	23
248	Mutational analysis of PINX1 in hereditary prostate cancer. <i>Prostate</i> , 2004, 60, 298-302.	1.2	22
249	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
250	Rare Variation in <i>TET2</i> Is Associated with Clinically Relevant Prostate Carcinoma in African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1456-1463.	1.1	22
251	Resistance to androgen receptor signaling inhibition does not necessitate development of neuroendocrine prostate cancer. <i>JCI Insight</i> , 2021, 6, .	2.3	22
252	Germline sequence variants of the LZTS1 gene are associated with prostate cancer risk. <i>Cancer Genetics and Cytogenetics</i> , 2002, 137, 1-7.	1.0	21

#	ARTICLE	IF	CITATIONS
253	Expression mapping at 12p12-13 in advanced prostate carcinoma. <i>International Journal of Cancer</i> , 2004, 109, 668-672.	2.3	21
254	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
255	Prevalence of the <i>HOXB13</i> G84E prostate cancer risk allele in men treated with radical prostatectomy. <i>BJU International</i> , 2014, 113, 830-835.	1.3	21
256	Association between variants in genes involved in the immune response and prostate cancer risk in men randomized to the finasteride arm in the Prostate Cancer Prevention Trial. <i>Prostate</i> , 2017, 77, 908-919.	1.2	21
257	Sequence variants in the human 25-hydroxyvitamin D3 1- $\alpha$ -hydroxylase (CYP27B1) gene are not associated with prostate cancer risk. <i>Prostate</i> , 2002, 53, 175-178.	1.2	20
258	Germline <i>ATBF1</i> mutations and prostate cancer risk. <i>Prostate</i> , 2006, 66, 1082-1085.	1.2	20
259	Polymorphic variants in <i>methylenetetrahydrofolate dehydrogenase</i> and prostate cancer. <i>Prostate</i> , 2007, 67, 1487-1497.	1.2	20
260	Single-Nucleotide Polymorphism-Based Genetic Risk Score and Patient Age at Prostate Cancer Diagnosis. <i>JAMA Network Open</i> , 2019, 2, e1918145.	2.8	20
261	Androgen receptor splice variant, AR-V7, and resistance to enzalutamide and abiraterone in men with metastatic castration-resistant prostate cancer (mCRPC). <i>Journal of Clinical Oncology</i> , 2014, 32, 5001-5001.	0.8	20
262	Mutational analysis of <i>ETV6</i> in prostate carcinoma. <i>Prostate</i> , 2002, 52, 305-310.	1.2	19
263	Mutational analysis of <i>SPANX</i> genes in families with X-Linked prostate cancer. <i>Prostate</i> , 2007, 67, 820-828.	1.2	19
264	Identification and characterization of novel SNPs in <i>CHEK2</i> in Ashkenazi Jewish men with prostate cancer. <i>Cancer Letters</i> , 2008, 270, 173-180.	3.2	19
265	Comprehensive mutational analysis and mRNA isoform quantification of <i>TP63</i> in normal and neoplastic human prostate cells. <i>Prostate</i> , 2009, 69, 559-569.	1.2	19
266	A genetic variant near <i>GATA3</i> implicated in inherited susceptibility and etiology of benign prostatic hyperplasia (BPH) and lower urinary tract symptoms (LUTS). <i>Prostate</i> , 2017, 77, 1213-1220.	1.2	19
267	Multiple antibodies to titin immunoreact with <i>AHNAK</i> and localize to the mitotic spindle machinery. <i>Cytoskeleton</i> , 2001, 50, 101-113.	4.4	18
268	Association of <i>CASP8 D302H</i> polymorphism with reduced risk of aggressive prostate carcinoma. <i>Prostate</i> , 2010, 70, 646-653.	1.2	18
269	A Peripheral Circulating TH1 Cytokine Profile Is Inversely Associated with Prostate Cancer Risk in CLUE II. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2561-2567.	1.1	18
270	Identification of a novel germline <i>SPOP</i> mutation in a family with hereditary prostate cancer. <i>Prostate</i> , 2014, 74, 983-990.	1.2	18



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271	Concept and benchmarks for assessing narrow-sense validity of genetic risk score values. <i>Prostate</i> , 2019, 79, 1099-1105.	1.2	18
272	Genomic and Clinicopathologic Characterization of ATM-deficient Prostate Cancer. <i>Clinical Cancer Research</i> , 2020, 26, 4869-4881.	3.2	18
273	Differential effects of growth factor antagonists on neoplastic and normal prostatic cells. <i>Prostate</i> , 1990, 17, 327-336.	1.2	17
274	Molecular and cellular markers for metastatic prostate cancer. <i>Cancer and Metastasis Reviews</i> , 1993, 12, 3-10.	2.7	17
275	Interaction effect of PTEN and CDKN1B chromosomal regions on prostate cancer linkage. <i>Human Genetics</i> , 2003, -1, 1-1.	1.8	17
276	Genome-wide expression analysis of recently processed formalin-fixed paraffin embedded human prostate tissues. <i>Prostate</i> , 2009, 69, 214-218.	1.2	17
277	Genome-wide two-locus epistasis scans in prostate cancer using two European populations. <i>Human Genetics</i> , 2012, 131, 1225-1234.	1.8	17
278	Health inequity drives disease biology to create disparities in prostate cancer outcomes. <i>Journal of Clinical Investigation</i> , 2022, 132, .	3.9	17
279	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
280	G1/S cell cycle proteins as markers of aggressive prostate carcinoma. <i>Urology</i> , 2000, 55, 316-322.	0.5	15
281	Comparison of Two Methods for Estimating Absolute Risk of Prostate Cancer Based on Single Nucleotide Polymorphisms and Family History. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1083-1088.	1.1	15
282	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
283	Association of prostate cancer risk with snps in regions containing androgen receptor binding sites captured by ChIP-on-chip analyses. <i>Prostate</i> , 2012, 72, 376-385.	1.2	15
284	Germline mutations in DNA repair genes are associated with bladder cancer risk and unfavourable prognosis. <i>BJU International</i> , 2018, 122, 808-813.	1.3	15
285	Germline BLM mutations and metastatic prostate cancer. <i>Prostate</i> , 2020, 80, 235-237.	1.2	15
286	Xq27-28 deletions in prostate carcinoma. <i>Genes Chromosomes and Cancer</i> , 2003, 37, 381-388.	1.5	14
287	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
288	AR splice variant 7 (AR-V7) and response to taxanes in men with metastatic castration-resistant prostate cancer (mCRPC).. <i>Journal of Clinical Oncology</i> , 2015, 33, 138-138.	0.8	14

#	ARTICLE	IF	CITATIONS
289	Distinct Genomic Alterations in Prostate Tumors Derived from African American Men. <i>Molecular Cancer Research</i> , 2020, 18, 1815-1824.	1.5	14
290	A Nonclassic CCAAT Enhancer Element Binding Protein Binding Site Contributes to $\hat{\pm}$ -Methylacyl-CoA Racemase Expression in Prostate Cancer. <i>Molecular Cancer Research</i> , 2005, 3, 110-118.	1.5	13
291	Lactoferrin CpG Island Hypermethylation and Decoupling of mRNA and Protein Expression in the Early Stages of Prostate Carcinogenesis. <i>American Journal of Pathology</i> , 2019, 189, 2311-2322.	1.9	13
292	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
293	Do Environmental Factors Modify the Genetic Risk of Prostate Cancer?. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 213-220.	1.1	12
294	Genetic variants in cell cycle control pathway confer susceptibility to aggressive prostate carcinoma. <i>Prostate</i> , 2016, 76, 479-490.	1.2	12
295	Germline mutations in <i>PPFIBP2</i> are associated with lethal prostate cancer. <i>Prostate</i> , 2018, 78, 1222-1228.	1.2	12
296	Molecular Characterization and Clinical Outcomes of Primary Gleason Pattern 5 Prostate Cancer After Radical Prostatectomy. <i>JCO Precision Oncology</i> , 2019, 3, 1-13.	1.5	12
297	Prostate Cancer Predisposition. <i>Urologic Clinics of North America</i> , 2021, 48, 283-296.	0.8	12
298	Inherited risk assessment and its clinical utility for predicting prostate cancer from diagnostic prostate biopsies. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, 25, 422-430.	2.0	12
299	Sequence variants in the 3'UTR of deoxyribonuclease TREX2: identification in a genetic screen and effects on catalysis by the recombinant proteins. <i>Advances in Enzyme Regulation</i> , 2004, 44, 37-49.	2.9	11
300	Genome-wide Association Study Identifies Loci at ATF7IP and KLK2 Associated with Percentage of Circulating Free PSA. <i>Neoplasia</i> , 2013, 15, 95-103.	2.3	11
301	Infectious mononucleosis, other infections and prostate-specific antigen concentration as a marker of prostate involvement during infection. <i>International Journal of Cancer</i> , 2016, 138, 2221-2230.	2.3	11
302	Genetic factors influencing prostate cancer risk in Norwegian men. <i>Prostate</i> , 2018, 78, 186-192.	1.2	11
303	<i>Trichomonas vaginalis</i> infection and prostate-specific antigen concentration: Insights into prostate involvement and prostate disease risk. <i>Prostate</i> , 2019, 79, 1622-1628.	1.2	11
304	Current progress and questions in germline genetics of prostate cancer. <i>Asian Journal of Urology</i> , 2019, 6, 3-9.	0.5	11
305	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
306	Association of prostate cancer polygenic risk score with number and laterality of tumor cores in active surveillance patients. <i>Prostate</i> , 2021, 81, 703-709.	1.2	11

#	ARTICLE	IF	CITATIONS
307	Transmission/disequilibrium tests of androgen receptor and glutathione S-transferase pi variants in prostate cancer families. <i>International Journal of Cancer</i> , 2002, 98, 938-942.	2.3	10
308	A Genome-Wide Survey over the ChIP-On-Chip Identified Androgen Receptor-Binding Genomic Regions Identifies a Novel Prostate Cancer Susceptibility Locus at 12q13.13. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2396-2403.	1.1	10
309	Germ-line sequence variants of PTEN do not have an important role in hereditary and non-hereditary prostate cancer susceptibility. <i>Journal of Human Genetics</i> , 2011, 56, 496-502.	1.1	10
310	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
311	Truncating Variants in p53/AIP1 Disrupting DNA Damage-Induced Apoptosis Are Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2006, 66, 10302-10307.	0.4	9
312	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
313	Association between pathogenic germline mutations in BRCA2 and ATM and tumor-infiltrating lymphocytes in primary prostate cancer. <i>Cancer Immunology, Immunotherapy</i> , 2022, 71, 943-951.	2.0	9
314	Somatic molecular subtyping of prostate tumors from HOXB13 G84E carriers. <i>Oncotarget</i> , 2017, 8, 22772-22782.	0.8	9
315	Insight into infection-mediated prostate damage: Contrasting patterns of C-reactive protein and prostate-specific antigen levels during infection. <i>Prostate</i> , 2017, 77, 1325-1334.	1.2	8
316	Inherited risk assessment of prostate cancer: it takes three to do it right. <i>Prostate Cancer and Prostatic Diseases</i> , 2020, 23, 59-61.	2.0	8
317	Use of Aspirin and Statins in Relation to Inflammation in Benign Prostate Tissue in the Placebo Arm of the Prostate Cancer Prevention Trial. <i>Cancer Prevention Research</i> , 2020, 13, 853-862.	0.7	8
318	Genome-wide Association Study Identifies Genetic Determinants of Urine PCA3 Levels in Men. <i>Neoplasia</i> , 2013, 15, 448-IN26.	2.3	7
319	E1A Transformed Normal Human Prostate Epithelial Cells Contain a 16q Deletion. <i>Cancer Genetics and Cytogenetics</i> , 1998, 103, 155-163.	1.0	6
320	A multigenic approach to evaluating prostate cancer risk in a systematic replication study. <i>Cancer Genetics and Cytogenetics</i> , 2008, 183, 94-98.	1.0	6
321	Peripheral Zone Inflammation Is Not Strongly Associated With Lower Urinary Tract Symptom Incidence and Progression in the Placebo Arm of the Prostate Cancer Prevention Trial. <i>Prostate</i> , 2016, 76, 1399-1408.	1.2	6
322	Inherited susceptibility for aggressive prostate cancer. <i>Asian Journal of Andrology</i> , 2012, 14, 415-418.	0.8	6
323	Does diabetes mellitus modify the association between 17q12 risk variant and prostate cancer aggressiveness?. <i>BJU International</i> , 2009, 104, 1200-1203.	1.3	5
324	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

#	ARTICLE	IF	CITATIONS
325	Key genes involved in the immune response are generally not associated with intraprostatic inflammation in men without a prostate cancer diagnosis: Results from the prostate cancer prevention trial. <i>Prostate</i> , 2016, 76, 565-574.	1.2	5
326	Updated insights into genetic contribution to prostate cancer predisposition: focus on HOXB13. <i>Canadian Journal of Urology</i> , 2019, 26, 12-13.	0.0	5
327	Polymorphisms Influencing Prostate-Specific Antigen Concentration May Bias Genome-Wide Association Studies on Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 88-93.	1.1	4
328	Sustained influence of infections on prostate-specific antigen concentration: An analysis of changes over 10 years of follow-up. <i>Prostate</i> , 2018, 78, 1024-1034.	1.2	4
329	Feasibility and performance of a novel probe panel to detect somatic DNA copy number alterations in clinical specimens for predicting prostate cancer progression. <i>Prostate</i> , 2020, 80, 1253-1262.	1.2	4
330	Specific Detection of Prostate Cancer Cells in Urine by RNA In Situ Hybridization. <i>Journal of Urology</i> , 2021, 206, 37-43.	0.2	4
331	Genetic Susceptibility for Low Testosterone in Men and Its Implications in Biology and Screening: Data from the UK Biobank. <i>European Urology Open Science</i> , 2021, 29, 36-46.	0.2	4
332	Incorporation of Polygenic Risk Score into Guidelines for Inherited Risk Assessment for Prostate Cancer. <i>European Urology</i> , 2021, 80, 139-141.	0.9	4
333	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
334	Germline copy number polymorphisms involving larger than 100 kb are uncommon in normal subjects. <i>Prostate</i> , 2007, 67, 227-233.	1.2	3
335	A novel method for detection of exfoliated prostate cancer cells in urine by RNA in situ hybridization. <i>Prostate Cancer and Prostatic Diseases</i> , 2021, 24, 220-232.	2.0	3
336	ATM loss in primary prostate cancer: Analysis of >1000 cases using a validated clinical-grade immunohistochemistry (IHC) assay. <i>Journal of Clinical Oncology</i> , 2019, 37, 5069-5069.	0.8	3
337	KLK3 germline mutation I179T complements DNA repair genes for predicting prostate cancer progression. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, , .	2.0	3
338	Identifying Phased Mutations and Complex Rearrangements in Human Prostate Cancer Cell Lines through Linked-Read Whole-Genome Sequencing. <i>Molecular Cancer Research</i> , 2022, 20, 1013-1020.	1.5	3
339	The somatic mutation landscape of germline <i>CHEK2</i> altered prostate cancer. <i>Journal of Clinical Oncology</i> , 2021, 39, 5084-5084.	0.8	2
340	Combined Longitudinal Clinical and Autopsy Phenomic Assessment in Lethal Metastatic Prostate Cancer: Recommendations for Advancing Precision Medicine. <i>European Urology Open Science</i> , 2021, 30, 47-62.	0.2	2
341	Evidence for a general cancer susceptibility locus at 3p24 in families with hereditary prostate cancer. <i>Cancer Letters</i> , 2005, 219, 177-182.	3.2	1
342	What Do Myeloma, Breast Cancer, and Prostate Cancer Have in Common?. <i>European Urology</i> , 2017, 71, 166-167.	0.9	1

#	ARTICLE	IF	CITATIONS
343	Differences in inherited risk among relatives of hereditary prostate cancer patients using genetic risk score. <i>Prostate</i> , 2018, 78, 1063-1068.	1.2	1
344	A snapshot of the expression signature of androgen receptor splicing variants and their distinctive transcriptional activities. , 2011, 71, 1656.		1
345	From the editorial office. <i>Prostate</i> , 2001, 48, 127-127.	1.2	0
346	GENETIC BASIS FOR PROSTATE CANCER. , 2011, , 39-52.		0
347	Association of the <i>HOXB13 G84E</i> mutation with increased risk for prostate cancer and other malignancies.. <i>Journal of Clinical Oncology</i> , 2014, 32, 1558-1558.	0.8	0
348	Effect of germline DNA repair gene mutations on outcomes in men with metastatic castration-resistant prostate cancer receiving first-line abiraterone and enzalutamide.. <i>Journal of Clinical Oncology</i> , 2018, 36, 221-221.	0.8	0
349	Donald S Coffey, a man who meant so much to so many. <i>American Journal of Clinical and Experimental Urology</i> , 2018, 6, 41-42.	0.4	0
350	Germline <i>BRCA2</i> , <i>ATM</i> and <i>CHEK2</i> alterations shape somatic mutation landscapes in prostate cancer.. <i>Journal of Clinical Oncology</i> , 2022, 40, 148-148.	0.8	0
351	The role of genetic testing in prostate cancer screening, diagnosis, and treatment. <i>Current Opinion in Oncology</i> , 2022, Publish Ahead of Print, .	1.1	0