Geraldine Cancel-Tassin

List of Publications by Year
in descending order

45 papers	4,141 citations	218592 26 h-index	206029 48 g-index
52 all docs	52 docs citations	52 times ranked	7059 citing authors

#	Article	IF	Citations
1	Multiple loci identified in a genome-wide association study of prostate cancer. Nature Genetics, 2008, 40, 310-315.	9.4	871
2	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	9.4	652
3	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. Nature Genetics, 2010, 42, 978-984.	9.4	493
4	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	9.4	264
5	HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). Human Genetics, 2013, 132, 5-14.	1.8	166
6	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. Journal of the National Cancer Institute, 2015, 107, djv279.	3.0	152
7	Genome-wide association study identifies multiple loci associated with bladder cancer risk. Human Molecular Genetics, 2014, 23, 1387-1398.	1.4	137
8	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724.	5.8	106
9	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. Human Molecular Genetics, 2014, 23, 6616-6633.	1.4	90
10	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. Nature Communications, 2018, 9, 2256.	5.8	88
11	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
12	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. Human Molecular Genetics, 2012, 21, 456-462.	1.4	81
13	Homologous recombination deficiency (HRD) score in germline BRCA2- versus ATM-altered prostate cancer. Modern Pathology, 2021, 34, 1185-1193.	2.9	61
14	The influence of obesity-related factors in the etiology of renal cell carcinomaâ€"A mendelian randomization study. PLoS Medicine, 2019, 16, e1002724.	3.9	59
15	PCAP is the major known prostate cancer predisposing locus in families from south and west Europe. European Journal of Human Genetics, 2001, 9, 135-142.	1.4	58
16	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. Journal of the National Cancer Institute, 2017, 109, .	3.0	57
17	Genome-wide interaction study of smoking and bladder cancer risk. Carcinogenesis, 2014, 35, 1737-1744.	1.3	50
18	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. Human Molecular Genetics, 2011, 20, 2869-2878.	1.4	43

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19	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. Nature Communications, 2018, 9, 4616.	5.8	43
20	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. Human Molecular Genetics, 2016, 25, 1203-1214.	1.4	38
21	Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. European Urology Oncology, 2021, 4, 570-579.	2.6	38
22	Comprehensive molecular classification of localized prostate adenocarcinoma reveals a tumour subtype predictive of non-aggressive disease. Annals of Oncology, 2018, 29, 1814-1821.	0.6	35
23	Screening, diagnosis and monitoring of sarcopenia: When to use which tool?. Clinical Nutrition ESPEN, 2022, 48, 36-44.	0.5	34
24	Mutational Profile of Aggressive, Localised Prostate Cancer from African Caribbean Men Versus European Ancestry Men. European Urology, 2019, 75, 11-15.	0.9	32
25	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. European Urology, 2020, 78, 316-320.	0.9	32
26	Genetic Variability in 8q24 Confers Susceptibility to Urothelial Carcinoma of the Upper Urinary Tract and is Linked With Patterns of Disease Aggressiveness at Diagnosis. Journal of Urology, 2012, 187, 424-428.	0.2	27
27	Sex specific associations in genome wide association analysis of renal cell carcinoma. European Journal of Human Genetics, 2019, 27, 1589-1598.	1.4	27
28	Genetic susceptibility to prostate cancer. BJU International, 2005, 96, 1380-1385.	1.3	24
29	Testosterone replacement therapy (TRT) and prostate cancer: An updated systematic review with a focus on previous or active localized prostate cancer. Urologic Oncology: Seminars and Original Investigations, 2020, 38, 661-670.	0.8	24
30	Africanâ€specific improvement of a polygenic hazard score for age at diagnosis of prostate cancer. International Journal of Cancer, 2021, 148, 99-105.	2.3	24
31	Association Study of Polymorphisms in the Human Estrogen Receptor Alpha Gene and Prostate Cancer Risk. European Urology, 2003, 44, 487-490.	0.9	23
32	Marital status and prostate cancer incidence: a pooled analysis of 12 caseâ€“control studies from the PRACTICAL consortium. European Journal of Epidemiology, 2021, 36, 913-925.	2.5	23
33	A Rare Germline HOXB13 Variant Contributes to Risk of Prostate Cancer in Men of African Ancestry. European Urology, 2022, 81, 458-462.	0.9	22
34	No evidence of linkage to HPC20 on chromosome 20q13 in hereditary prostate cancer. International Journal of Cancer, 2001, 93, 455-456.	2.3	20
35	Assessment of Xpert Bladder Cancer Monitor test performance for the detection of recurrence during non-muscle invasive bladder cancer follow-up. World Journal of Urology, 2021, 39, 3329-3335.	1.2	19
36	Urothelial Cancers with Small Cell Variant Histology Have Confirmed High Tumor Mutational Burden, Frequent TP53 and RB Mutations, and a Unique Gene Expression Profile. European Urology Oncology, 2021, 4, 297-300.	2.6	18

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37	A GWAS in uveal melanoma identifies risk polymorphisms in the CLPTM1L locus. Npj Genomic Medicine, 2017, 2, .	1.7	17
38	The Genetic Complexity of Prostate Cancer. Genes, 2020, 11, 1396.	1.0	9
39	Performance of African-ancestry-specific polygenic hazard score varies according to local ancestry in 8q24. Prostate Cancer and Prostatic Diseases, 2022, 25, 229-237.	2.0	9
40	Bayesian predictive model to assess BRCA2 mutational status according to clinical history: Early onset, metastatic phenotype or family history of breast/ovary cancer. Prostate, 2021, 81, 318-325.	1.2	7
41	Different Pigmentation Risk Loci for High-Risk Monosomy 3 and Low-Risk Disomy 3 Uveal Melanomas. Journal of the National Cancer Institute, 2022, 114, 302-309.	3.0	5
42	Genetic variability in 13q33 and 9q34 is linked to aggressiveness patterns and a higher risk of progression of nonâ€muscleâ€invasive bladder cancer at the time of diagnosis. BJU International, 2021, 127, 375-383.	1.3	2
43	A study of the immunohistochemical profile of bladder cancer in neuro-urological patients by the French Association of Urology. World Journal of Urology, 2022, , 1.	1.2	2
44	DNA damage repair gene germline profiling for metastatic prostate cancer patients of different ancestries. Prostate, 2022, 82, 1196-1201.	1.2	2
45	Diagnosis of prostate cancer in one day: The benefits of cytology in tumour detection. Cytopathology, 2021, 32, 211-216.	0.4	0