

Geraldine Cancel-Tassin

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

41
papers

2,723
citations

20
h-index

52
g-index

52
ext. papers

3,560
ext. citations

9.5
avg, IF

3.29
L-index

#	Paper	IF	Citations
41	A study of the immunohistochemical profile of bladder cancer in neuro-urological patients by the French Association of Urology.. <i>World Journal of Urology</i> , 2022 , 1	4	0
40	Screening, diagnosis and monitoring of sarcopenia: When to use which tool?. <i>Clinical Nutrition ESPEN</i> , 2022 , 48, 36-44	1.3	5
39	Assessment of Xpert Bladder Cancer Monitor test performance for the detection of recurrence during non-muscle invasive bladder cancer follow-up. <i>World Journal of Urology</i> , 2021 , 39, 3329-3335	4	8
38	Marital status and prostate cancer incidence: a pooled analysis of 12 case-control studies from the PRACTICAL consortium. <i>European Journal of Epidemiology</i> , 2021 , 36, 913-925	12.1	2
37	Urothelial Cancers with Small Cell Variant Histology Have Confirmed High Tumor Mutational Burden, Frequent TP53 and RB Mutations, and a Unique Gene Expression Profile. <i>European Urology Oncology</i> , 2021 , 4, 297-300	6.7	4
36	African-specific improvement of a polygenic hazard score for age at diagnosis of prostate cancer. <i>International Journal of Cancer</i> , 2021 , 148, 99-105	7.5	7
35	Genetic variability in 13q33 and 9q34 is linked to aggressiveness patterns and a higher risk of progression of non-muscle-invasive bladder cancer at the time of diagnosis. <i>BJU International</i> , 2021 , 127, 375-383	5.6	1
34	Diagnosis of prostate cancer in one day: The benefits of cytology in tumour detection. <i>Cytopathology</i> , 2021 , 32, 211-216	1.3	
33	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	36.3	62
32	Homologous recombination deficiency (HRD) score in germline BRCA2- versus ATM-altered prostate cancer. <i>Modern Pathology</i> , 2021 , 34, 1185-1193	9.8	15
31	Bayesian predictive model to assess BRCA2 mutational status according to clinical history: Early onset, metastatic phenotype or family history of breast/ovary cancer. <i>Prostate</i> , 2021 , 81, 318-325	4.2	3
30	Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. <i>European Urology Oncology</i> , 2021 , 4, 570-579	6.7	12
29	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	10.2	13
28	Testosterone replacement therapy (TRT) and prostate cancer: An updated systematic review with a focus on previous or active localized prostate cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2020 , 38, 661-670	2.8	10
27	The Genetic Complexity of Prostate Cancer. <i>Genes</i> , 2020 , 11,	4.2	2
26	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
25	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019 , 27, 1589-1598	5.3	15

24	The influence of obesity-related factors in the etiology of renal cell carcinoma-A mendelian randomization study. <i>PLoS Medicine</i> , 2019 , 16, e1002724	11.6	38
23	Mutational Profile of Aggressive, Localised Prostate Cancer from African Caribbean Men Versus European Ancestry Men. <i>European Urology</i> , 2019 , 75, 11-15	10.2	13
22	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018 , 50, 928-936	36.3	340
21	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018 , 9, 2256	17.4	57
20	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , 2018 , 9, 4616	17.4	30
19	Comprehensive molecular classification of localized prostate adenocarcinoma reveals a tumour subtype predictive of non-aggressive disease. <i>Annals of Oncology</i> , 2018 , 29, 1814-1821	10.3	18
18	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
17	A GWAS in uveal melanoma identifies risk polymorphisms in the locus. <i>Npj Genomic Medicine</i> , 2017 , 2,	6.2	9
16	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	38
15	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. <i>Human Molecular Genetics</i> , 2016 , 25, 1203-14	5.6	20
14	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015 , 107, djv279	9.7	107
13	Genome-wide interaction study of smoking and bladder cancer risk. <i>Carcinogenesis</i> , 2014 , 35, 1737-44	4.6	33
12	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
11	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014 , 23, 6616-33	5.6	77
10	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	6.3	134
9	Genetic variability in 8q24 confers susceptibility to urothelial carcinoma of the upper urinary tract and is linked with patterns of disease aggressiveness at diagnosis. <i>Journal of Urology</i> , 2012 , 187, 424-8	2.5	20
8	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012 , 21, 456-62	5.6	74
7	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, 2869-78	5.6	39

6	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 978-84	36.3	408
5	Multiple loci identified in a genome-wide association study of prostate cancer. <i>Nature Genetics</i> , 2008 , 40, 310-5	36.3	787
4	Genetic susceptibility to prostate cancer. <i>BJU International</i> , 2005 , 96, 1380-5	5.6	16
3	Association study of polymorphisms in the human estrogen receptor alpha gene and prostate cancer risk. <i>European Urology</i> , 2003 , 44, 487-90	10.2	21
2	No evidence of linkage to HPC20 on chromosome 20q13 in hereditary prostate cancer. <i>International Journal of Cancer</i> , 2001 , 93, 455-6	7.5	19
1	PCAP is the major known prostate cancer predisposing locus in families from south and west Europe. <i>European Journal of Human Genetics</i> , 2001 , 9, 135-42	5.3	50