

Tokhir Dadaev

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

3,743
citations

331670

21
h-index

526287

27
g-index

32
all docs

32
docs citations

32
times ranked

6650
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	21.4	652
2	Germline <i>BRCA</i> Mutations Are Associated With Higher Risk of Nodal Involvement, Distant Metastasis, and Poor Survival Outcomes in Prostate Cancer. <i>Journal of Clinical Oncology</i> , 2013, 31, 1748-1757.	1.6	641
3	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	21.4	492
4	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014, 46, 1103-1109.	21.4	408
5	Effect of <i>BRCA</i> Mutations on Metastatic Relapse and Cause-specific Survival After Radical Treatment for Localised Prostate Cancer. <i>European Urology</i> , 2015, 68, 186-193.	1.9	279
6	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	21.4	265
7	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.	21.4	264
8	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with <i>TERT</i> expression. <i>Human Molecular Genetics</i> , 2013, 22, 2520-2528.	2.9	100
9	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	12.8	88
10	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. <i>Human Molecular Genetics</i> , 2015, 24, 5589-5602.	2.9	67
11	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.	2.5	56
12	Germline DNA Repair Gene Mutations in Young-onset Prostate Cancer Cases in the UK: Evidence for a More Extensive Genetic Panel. <i>European Urology</i> , 2019, 76, 329-337.	1.9	48
13	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , 2018, 9, 4616.	12.8	43
14	Rare germline variants in DNA repair genes and the angiogenesis pathway predispose prostate cancer patients to develop metastatic disease. <i>British Journal of Cancer</i> , 2018, 119, 96-104.	6.4	40
15	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. <i>Journal of the National Cancer Institute</i> , 2021, 113, 616-625.	6.3	40
16	Rare Germline Variants in <i>ATM</i> Predispose to Prostate Cancer: A PRACTICAL Consortium Study. <i>European Urology Oncology</i> , 2021, 4, 570-579.	5.4	38
17	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.	3.8	37
18	Clinical implications of family history of prostate cancer and genetic risk single nucleotide polymorphism (<i>SNP</i>) profiles in an active surveillance cohort. <i>BJU International</i> , 2013, 112, 666-673.	2.5	34

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19	Fine-Mapping the HOXB Region Detects Common Variants Tagging a Rare Coding Allele: Evidence for Synthetic Association in Prostate Cancer. <i>PLoS Genetics</i> , 2014, 10, e1004129.	3.5	34
20	Homeobox B13 G84E Mutation and Prostate Cancer Risk. <i>European Urology</i> , 2019, 75, 834-845.	1.9	28
21	The PROFILE Feasibility Study: Targeted Screening of Men With a Family History of Prostate Cancer. <i>Oncologist</i> , 2016, 21, 716-722.	3.7	27
22	Gene and pathway level analyses of germline DNA-repair gene variants and prostate cancer susceptibility using the iCOGS-genotyping array. <i>British Journal of Cancer</i> , 2016, 114, 945-952.	6.4	17
23	LocusExplorer: a user-friendly tool for integrated visualization of human genetic association data and biological annotations. <i>Bioinformatics</i> , 2016, 32, 949-951.	4.1	13
24	Prostate-specific antigen velocity in a prospective prostate cancer screening study of men with genetic predisposition. <i>British Journal of Cancer</i> , 2018, 118, 266-276.	6.4	12
25	Prostate cancer risk in men of differing genetic ancestry and approaches to disease screening and management in these groups. <i>British Journal of Cancer</i> , 2022, 126, 1366-1373.	6.4	12
26	Abstract 2546: Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. , 2013, , .		1
27	Relationship of self-reported body size and shape with risk for prostate cancer: A UK case-control study. <i>PLoS ONE</i> , 2020, 15, e0238928.	2.5	0
28	Abstract 2612: The PROFILE study; Germline genetic profiling: Correlation with targeted prostate cancer screening and treatment. , 2012, , .		0
29	Abstract 4495: Clinical implications of family history of prostate cancer in an active surveillance cohort. , 2012, , .		0