

# Tokhir Dadaev

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6351912/publications.pdf>

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  | 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        |
| 2  | 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        |
| 3  | 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       |
| 4  | Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. <i>European Urology Oncology</i> , 2021, 4, 570-579.  | 5.4  | 38        |
| 5  | 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         |
| 6  | 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        |
| 7  | Homeobox B13 G84E Mutation and Prostate Cancer Risk. <i>European Urology</i> , 2019, 75, 834-845.  | 1.9  | 28        |
| 8  | 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        |
| 9  | Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , 2018, 9, 4616.   | 12.8 | 43        |
| 10 | 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        |
| 11 | 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       |
| 12 | 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        |
| 13 | 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        |
| 14 | 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        |
| 15 | 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        |
| 16 | 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        |
| 17 | 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        |
| 18 | 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        |

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 19 | Effect of BRCA 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       |
| 20 | 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        |
| 21 | 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       |
| 22 | Germline BRCA 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       |
| 23 | 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       |
| 24 | Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. <i>Human Molecular Genetics</i> , 2013, 22, 2520-2528.                                    | 2.9  | 100       |
| 25 | Clinical implications of family history of prostate cancer and genetic risk single nucleotide polymorphism (SNP) profiles in an active surveillance cohort. <i>BJU International</i> , 2013, 112, 666-673.  | 2.5  | 34        |
| 26 | Abstract 2546: Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. , 2013, , .  |      | 1         |
| 27 | Abstract 2612: The PROFILE study; Germline genetic profiling: Correlation with targeted prostate cancer screening and treatment. , 2012, , .  |      | 0         |
| 28 | Abstract 4495: Clinical implications of family history of prostate cancer in an active surveillance cohort. , 2012, , .   |      | 0         |
| 29 | 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       |