

Edward J Saunders

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

4,574
citations

257450

24
h-index

477307

29
g-index

40
all docs

40
docs citations

40
times ranked

7235
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	Rare Germline Variants Are Associated with Rapid Biochemical Recurrence After Radical Prostate Cancer Treatment: A Pan Prostate Cancer Group Study. <i>European Urology</i> , 2022, 82, 201-211.	1.9	2
3	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
4	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
5	Identification of Germline Genetic Variants that Increase Prostate Cancer Risk and Influence Development of Aggressive Disease. <i>Cancers</i> , 2021, 13, 760.	3.7	22
6	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
7	12 new susceptibility loci for prostate cancer identified by genome-wide association study in Japanese population. <i>Nature Communications</i> , 2019, 10, 4422.	12.8	49
8	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
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	DNA repair gene panel mutations in young onset prostate cancer cases in the.. <i>Journal of Clinical Oncology</i> , 2018, 36, 18-18.	1.6	0
14	Germline sequencing of advanced prostate cancer patients in the BARCODE2 study.. <i>Journal of Clinical Oncology</i> , 2018, 36, e13617-e13617.	1.6	0
15	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
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

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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	Prostate-specific antigen velocity as a predictive biomarker in a prospective prostate cancer screening study (IMPACT study).. <i>Journal of Clinical Oncology</i> , 2015, 33, 16-16.	1.6	0
21	Abstract 4606: Fine mapping of 64 prostate cancer GWAS regions identifies multiple novel association signals. , 2015, , .		0
22	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
23	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
24	The PROFILE feasibility study: Genetic prostate cancer risk stratification for targeted screening.. <i>Journal of Clinical Oncology</i> , 2014, 32, 22-22.	1.6	0
25	Abstract 5065: Fine-mapping theHOXBregion detects common variants tagging a rare coding allele: Evidence for synthetic association in prostate cancer. , 2014, , .		0
26	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
27	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
28	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.	2.9	118
29	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
30	Clinical implications of family history of prostate cancer and genetic risk single nucleotide polymorphism (<scp>SNP</scp>) profiles in an active surveillance cohort. <i>BJU International</i> , 2013, 112, 666-673.	2.5	34
31	Abstract 2546: Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates withTERTexpression.. , 2013, , .		1
32	Effect of germ-line BRCA mutations in biochemical relapse and survival after treatment for localized prostate cancer.. <i>Journal of Clinical Oncology</i> , 2013, 31, 29-29.	1.6	2
33	Abstract 2612: The PROFILE study; Germline genetic profiling: Correlation with targeted prostate cancer screening and treatment. , 2012, , .		0
34	Abstract 4495: Clinical implications of family history of prostate cancer in an active surveillance cohort. , 2012, , .		0
35	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
36	Abstract 3810: BRCA2 is a moderate penetrance gene contributing to young onset prostate cancer, but not disease over 65 years. , 2011, , .		0

#	ARTICLE	IF	CITATIONS
37	The rs10993994 Risk Allele for Prostate Cancer Results in Clinically Relevant Changes in Microseminoprotein-Beta Expression in Tissue and Urine. PLoS ONE, 2010, 5, e13363.	2.5	73
38	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. Nature Genetics, 2009, 41, 1116-1121.	21.4	389
39	Multiple loci on 8q24 associated with prostate cancer susceptibility. Nature Genetics, 2009, 41, 1058-1060.	21.4	273