Edward J Saunders

List of Publications by Year
in descending order

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	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	21.4	652
2	Germline BRCA Mutations Are Associated With Higher Risk of Nodal Involvement, Distant Metastasis, and Poor Survival Outcomes in Prostate Cancer. Journal of Clinical Oncology, 2013, 31, 1748-1757.	1.6	641
3	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. Nature Genetics, 2013, 45, 385-391.	21.4	492
4	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	21.4	408
5	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. Nature Genetics, 2009, 41, 1116-1121.	21.4	389
6	Effect of BRCA Mutations on Metastatic Relapse and Cause-specific Survival After Radical Treatment for Localised Prostate Cancer. European Urology, 2015, 68, 186-193.	1.9	279
7	Multiple loci on 8q24 associated with prostate cancer susceptibility. Nature Genetics, 2009, 41, 1058-1060.	21.4	273
8	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. Nature Genetics, 2011, 43, 785-791.	21.4	265
9	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.	21.4	264
10	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. Human Molecular Genetics, 2013, 22, 408-415.	2.9	118
11	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. Human Molecular Genetics, 2013, 22, 2520-2528.	2.9	100
12	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. Nature Communications, 2018, 9, 2256.	12.8	88
13	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
14	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. Human Molecular Genetics, 2015, 24, 5589-5602.	2.9	67
15	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1121-1129.	2.5	56
16	12 new susceptibility loci for prostate cancer identified by genome-wide association study in Japanese population. Nature Communications, 2019, 10, 4422.	12.8	49
17	Germline DNA Repair Gene Mutations in Young-onset Prostate Cancer Cases in the UK: Evidence for a More Extensive Genetic Panel. European Urology, 2019, 76, 329-337.	1.9	48
18	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. Nature Communications, 2018, 9, 4616.	12.8	43

#	Article	IF	Citations
19	Rare germline variants in DNA repair genes and the angiogenesis pathway predispose prostate cancer patients to develop metastatic disease. British Journal of Cancer, 2018, 119, 96-104.	6.4	40
20	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. Journal of the National Cancer Institute, 2021, 113, 616-625.	6.3	40
21	Rare Germline Variants in ATM Predispose to Prostate Cancer: A PRACTICAL Consortium Study. European Urology Oncology, 2021, 4, 570-579.	5.4	38
22	Clinical implications of family history of prostate cancer and genetic risk single nucleotide polymorphism (<scp>SNP</scp>) profiles in an active surveillance cohort. BJU International, 2013, 112, 666-673.	2.5	34
23	Fine-Mapping the HOXB Region Detects Common Variants Tagging a Rare Coding Allele: Evidence for Synthetic Association in Prostate Cancer. PLoS Genetics, 2014, 10, e1004129.	3.5	34
24	The PROFILE Feasibility Study: Targeted Screening of Men With a Family History of Prostate Cancer. Oncologist, 2016, 21, 716-722.	3.7	27
25	Identification of Germline Genetic Variants that Increase Prostate Cancer Risk and Influence Development of Aggressive Disease. Cancers, 2021, 13, 760.	3.7	22
26	Gene and pathway level analyses of germline DNA-repair gene variants and prostate cancer susceptibility using the iCOGS-genotyping array. British Journal of Cancer, 2016, 114, 945-952.	6.4	17
27	Prostate cancer risk in men of differing genetic ancestry and approaches to disease screening and management in these groups. British Journal of Cancer, 2022, 126, 1366-1373.	6.4	12
28	Effect of germ-line BRCA mutations in biochemical relapse and survival after treatment for localized prostate cancer.. Journal of Clinical Oncology, 2013, 31, 29-29.	1.6	2
29	Rare Germline Variants Are Associated with Rapid Biochemical Recurrence After Radical Prostate Cancer Treatment: A Pan Prostate Cancer Group Study. European Urology, 2022, 82, 201-211.	1.9	2
30	Abstract 2546: Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates withTERTexpression.. , 2013, , .		1
31	Abstract 3810: BRCA2 is a moderate penetrance gene contributing to young onset prostate cancer, but not disease over 65 years. , 2011, , .		0
32	Abstract 2612: The PROFILE study; Germline genetic profiling: Correlation with targeted prostate cancer screening and treatment. , 2012, , .		0
33	Abstract 4495: Clinical implications of family history of prostate cancer in an active surveillance cohort. , 2012, , .		0
34	The PROFILE feasibility study: Genetic prostate cancer risk stratification for targeted screening.. Journal of Clinical Oncology, 2014, 32, 22-22.	1.6	0
35	Abstract 5065: Fine-mapping theHOXBregion detects common variants tagging a rare coding allele: Evidence for synthetic association in prostate cancer. , 2014, , .		0
36	Prostate-specific antigen velocity as a predictive biomarker in a prospective prostate cancer screening study (IMPACT study).. Journal of Clinical Oncology, 2015, 33, 16-16.	1.6	0

#	Article	IF	Citations
37	Abstract 4606: Fine mapping of 64 prostate cancer GWAS regions identifies multiple novel association signals. , 2015, , .		0
38	DNA repair gene panel mutations in young onset prostate cancer cases in the.. Journal of Clinical Oncology, 2018, 36, 18-18.	1.6	0
39	Germline sequencing of advanced prostate cancer patients in the BARCODE2 study.. Journal of Clinical Oncology, 2018, 36, e13617-e13617.	1.6	0