

Dennis J Hazelett

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

2,677
citations

279798

23
h-index

414414

32
g-index

38
all docs

38
docs citations

38
times ranked

7114
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	Granulocyte-Monocyte Progenitors and Monocyte-Dendritic Cell Progenitors Independently Produce Functionally Distinct Monocytes. <i>Immunity</i> , 2017, 47, 890-902.e4.	14.3	297
3	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	2.5	278
4	<i>motifbreakR</i> : an R/Bioconductor package for predicting variant effects at transcription factor binding sites. <i>Bioinformatics</i> , 2015, 31, 3847-3849.	4.1	208
5	Comprehensive Functional Annotation of 77 Prostate Cancer Risk Loci. <i>PLoS Genetics</i> , 2014, 10, e1004102.	3.5	167
6	ONECUT2 is a targetable master regulator of lethal prostate cancer that suppresses the androgen axis. <i>Nature Medicine</i> , 2018, 24, 1887-1898.	30.7	113
7	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv431.	6.3	111
8	act up Controls Actin Polymerization to Alter Cell Shape and Restrict Hedgehog Signaling in the <i>Drosophila</i> Eye Disc. <i>Cell</i> , 2000, 101, 271-281.	28.9	105
9	ELMER v.2: an R/Bioconductor package to reconstruct gene regulatory networks from DNA methylation and transcriptome profiles. <i>Bioinformatics</i> , 2019, 35, 1974-1977.	4.1	87
10	A Mosaic Genetic Screen Reveals Distinct Roles for trithorax and Polycomb Group Genes in <i>Drosophila</i> Eye Development. <i>Genetics</i> , 2004, 166, 187-200.	2.9	71
11	Comparison of Parallel High-Throughput RNA Sequencing Between Knockout of TDP-43 and Its Overexpression Reveals Primarily Nonreciprocal and Nonoverlapping Gene Expression Changes in the Central Nervous System of <i>Drosophila</i> . <i>G3: Genes, Genomes, Genetics</i> , 2012, 2, 789-802.	1.8	71
12	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
13	CRISPR-mediated deletion of prostate cancer risk-associated CTCF loop anchors identifies repressive chromatin loops. <i>Genome Biology</i> , 2018, 19, 160.	8.8	60
14	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	57
15	Enrichment of risk SNPs in regulatory regions implicate diverse tissues in Parkinson's disease etiology. <i>Scientific Reports</i> , 2016, 6, 30509.	3.3	53
16	Non-coding somatic mutations converge on the PAX8 pathway in ovarian cancer. <i>Nature Communications</i> , 2020, 11, 2020.	12.8	52
17	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , 2015, 24, 5603-5618.	2.9	50
18	Nucleosome positioning and histone modifications define relationships between regulatory elements and nearby gene expression in breast epithelial cells. <i>BMC Genomics</i> , 2014, 15, 331.	2.8	40

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19	Cell-type-specific enrichment of risk-associated regulatory elements at ovarian cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2015, 24, 3595-3607.	2.9	40
20	GENAVi: a shiny web application for gene expression normalization, analysis and visualization. <i>BMC Genomics</i> , 2019, 20, 745.	2.8	40
21	A Study of High-Grade Serous Ovarian Cancer Origins Implicates the SOX18 Transcription Factor in Tumor Development. <i>Cell Reports</i> , 2019, 29, 3726-3735.e4.	6.4	39
22	The PAX8 cistrome in epithelial ovarian cancer. <i>Oncotarget</i> , 2017, 8, 108316-108332.	1.8	38
23	Motor neuron expression of the voltage-gated calcium channel cacophony restores locomotion defects in a <i>Drosophila</i> , TDP-43 loss of function model of ALS. <i>Brain Research</i> , 2014, 1584, 39-51.	2.2	34
24	Identification of a Novel Mucin Gene <i>HCG22</i> Associated With Steroid-Induced Ocular Hypertension. , 2015, 56, 2737.		28
25	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. <i>Gynecologic Oncology</i> , 2019, 153, 343-355.	1.4	28
26	Common variants at the <i>CHEK2</i> gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015, 36, 1341-1353.	2.8	24
27	Functional Analysis and Fine Mapping of the 9p22.2 Ovarian Cancer Susceptibility Locus. <i>Cancer Research</i> , 2019, 79, 467-481.	0.9	22
28	A rare variant, which destroys a FoxA1 site at 8q24, is associated with prostate cancer risk. <i>Cell Cycle</i> , 2013, 12, 379-380.	2.6	20
29	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1609-1618.	2.5	18
30	Reducing GWAS Complexity. <i>Cell Cycle</i> , 2016, 15, 22-24.	2.6	16
31	Ovarian Cancer Risk Variants Are Enriched in Histotype-Specific Enhancers and Disrupt Transcription Factor Binding Sites. <i>American Journal of Human Genetics</i> , 2020, 107, 622-635.	6.2	14
32	StateHub-StatePaintR: rapid and reproducible chromatin state evaluation for custom genome annotation. <i>F1000Research</i> , 0, 7, 214.	1.6	4
33	Affinity Density: a novel genomic approach to the identification of transcription factor regulatory targets. <i>Bioinformatics</i> , 2009, 25, 1617-1624.	4.1	3
34	Segment-specific muscle degeneration is triggered directly by a steroid hormone during insect metamorphosis. <i>Journal of Neurobiology</i> , 2005, 62, 164-177.	3.6	1
35	A molecular taxonomy of tumors independent of tissue-of-origin. <i>IScience</i> , 2021, 24, 103084.	4.1	0