

Diptasri Mandal

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

16
papers

1,037
citations

9
h-index

18
g-index

18
ext. papers

1,574
ext. citations

5.7
avg, IF

2.54
L-index

#	Paper	IF	Citations
16	REVEL: An Ensemble Method for Predicting the Pathogenicity of Rare Missense Variants. <i>American Journal of Human Genetics</i> , 2016 , 99, 877-885	11	722
15	HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). <i>Human Genetics</i> , 2013 , 132, 5-14	6.3	134
14	A recurrent mutation in PARK2 is associated with familial lung cancer. <i>American Journal of Human Genetics</i> , 2015 , 96, 301-8	11	47
13	Genome-wide association of familial prostate cancer cases identifies evidence for a rare segregating haplotype at 8q24.21. <i>Human Genetics</i> , 2016 , 135, 923-38	6.3	27
12	Genome-wide association study of familial lung cancer. <i>Carcinogenesis</i> , 2018 , 39, 1135-1140	4.6	24
11	Focused Analysis of Exome Sequencing Data for Rare Germline Mutations in Familial and Sporadic Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2016 , 11, 52-61	8.9	18
10	Familial Lung Cancer: A Brief History from the Earliest Work to the Most Recent Studies. <i>Genes</i> , 2017 , 8,	4.2	16
9	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2020 , 78, 316-320	10.2	13
8	Rare Variants in Known Susceptibility Loci and Their Contribution to Risk of Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2018 , 13, 1483-1495	8.9	12
7	Two-stage Study of Familial Prostate Cancer by Whole-exome Sequencing and Custom Capture Identifies 10 Novel Genes Associated with the Risk of Prostate Cancer. <i>European Urology</i> , 2021 , 79, 353-361	10.2	9
6	Whole Exome Sequencing of Highly Aggregated Lung Cancer Families Reveals Linked Loci for Increased Cancer Risk on Chromosomes 12q, 7p, and 4q. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 434-442	4	6
5	gsSKAT: Rapid gene set analysis and multiple testing correction for rare-variant association studies using weighted linear kernels. <i>Genetic Epidemiology</i> , 2017 , 41, 297-308	2.6	5
4	Post hoc Analysis for Detecting Individual Rare Variant Risk Associations Using Probit Regression Bayesian Variable Selection Methods in Case-Control Sequencing Studies. <i>Genetic Epidemiology</i> , 2016 , 40, 461-9	2.6	3
3	Genome-wide interaction analysis identified low-frequency variants with sex disparity in lung cancer risk.. <i>Human Molecular Genetics</i> , 2022 ,	5.6	1
2	Rare deleterious germline variants and risk of lung cancer. <i>Npj Precision Oncology</i> , 2021 , 5, 12	9.8	0
1	Situs Inversus Totalis: From Anatomical Variation to Clinical Studies. <i>FASEB Journal</i> , 2015 , 29, 552.11	0.9	