

Richa Gupta

List of Publications by Year in Descending Order

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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.

14
papers

1,817
citations

11
h-index

16
g-index

16
ext. papers

2,496
ext. citations

17.5
avg, IF

2.63
L-index

#	Paper	IF	Citations
14	Curated variation benchmarks for challenging medically relevant autosomal genes.. <i>Nature Biotechnology</i> , 2022 ,	44.5	12
13	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals.. <i>Nature Genetics</i> , 2022 ,	36.3	7
12	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. <i>Nature Genetics</i> , 2021 , 53, 1311-1321	36.3	27
11	Epigenome-wide association study of serum cotinine in current smokers reveals novel genetically driven loci. <i>Clinical Epigenetics</i> , 2019 , 11, 1	7.7	40
10	Genome-wide association study in Finnish twins highlights the connection between nicotine addiction and neurotrophin signaling pathway. <i>Addiction Biology</i> , 2019 , 24, 549-561	4.6	7
9	Neuregulin signaling pathway in smoking behavior. <i>Translational Psychiatry</i> , 2017 , 7, e1212	8.6	5
8	Genetic variants linked to education predict longevity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 13366-13371	11.5	90
7	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. <i>Nature Genetics</i> , 2016 , 48, 624-33	36.3	602
6	Genome-wide association study identifies 74 loci associated with educational attainment. <i>Nature</i> , 2016 , 533, 539-42	50.4	850
5	A Genome-Wide Association Study of a Biomarker of Nicotine Metabolism. <i>PLoS Genetics</i> , 2015 , 11, e1005498	8.0	80
4	Characterization of a glycoside hydrolase family 1 β galactosidase from hot spring metagenome with transglycosylation activity. <i>Applied Biochemistry and Biotechnology</i> , 2012 , 168, 1681-93	3.2	28
3	Characterization of MspNI (G/GWCC) and MspNII (R/GATCY), novel thermostable Type II restriction endonucleases from <i>Meiothermus</i> sp., isoschizomers of <i>Avall</i> and <i>BstYI</i> . <i>Molecular Biology Reports</i> , 2012 , 39, 5607-14	2.8	10
2	Restriction endonucleases: natural and directed evolution. <i>Applied Microbiology and Biotechnology</i> , 2012 , 94, 583-99	5.7	20
1	High AU content: a signature of upregulated miRNA in cardiac diseases. <i>Bioinformatics</i> , 2010 , 5, 132-5	1.1	6