

Saurav Guha

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

1,445
citations

331670

21
h-index

454955

30
g-index

35
all docs

35
docs citations

35
times ranked

3867
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular genetic evidence for overlap between general cognitive ability and risk for schizophrenia: a report from the Cognitive Genomics consortium (COGENT). <i>Molecular Psychiatry</i> , 2014, 19, 168-174.	7.9	178
2	Sequencing an Ashkenazi reference panel supports population-targeted personal genomics and illuminates Jewish and European origins. <i>Nature Communications</i> , 2014, 5, 4835.	12.8	156
3	High rate of disease-related copy number variations in childhood onset schizophrenia. <i>Molecular Psychiatry</i> , 2014, 19, 568-572.	7.9	116
4	Genome-wide association study implicates NDST3 in schizophrenia and bipolar disorder. <i>Nature Communications</i> , 2013, 4, 2739.	12.8	101
5	Implication of a Rare Deletion at Distal 16p11.2 in Schizophrenia. <i>JAMA Psychiatry</i> , 2013, 70, 253.	11.0	69
6	A complete deficiency of Hyaluronoglucosaminidase 1 (<i>HYAL1</i>) presenting as familial juvenile idiopathic arthritis. <i>Journal of Inherited Metabolic Disease</i> , 2011, 34, 1013-1022.	3.6	68
7	De novo copy number variants are associated with congenital diaphragmatic hernia. <i>Journal of Medical Genetics</i> , 2012, 49, 650-659.	3.2	68
8	Consensus interpretation of the p.Met34Thr and p.Val37Ile variants in GJB2 by the ClinGen Hearing Loss Expert Panel. <i>Genetics in Medicine</i> , 2019, 21, 2442-2452.	2.4	56
9	Variant Classification Concordance using the ACMG-AMP Variant Interpretation Guidelines across Nine Genomic Implementation Research Studies. <i>American Journal of Human Genetics</i> , 2020, 107, 932-941.	6.2	51
10	Genome-wide mapping of IBD segments in an Ashkenazi PD cohort identifies associated haplotypes. <i>Human Molecular Genetics</i> , 2014, 23, 4693-4702.	2.9	49
11	Implications for health and disease in the genetic signature of the Ashkenazi Jewish population. <i>Genome Biology</i> , 2012, 13, R2.	9.6	48
12	Best practices for the interpretation and reporting of clinical whole genome sequencing. <i>Npj Genomic Medicine</i> , 2022, 7, 27.	3.8	48
13	Molecular phylogeny of musk deer: A genomic view with mitochondrial 16S rRNA and cytochrome b gene. <i>Molecular Phylogenetics and Evolution</i> , 2007, 42, 585-597.	2.7	43
14	Association of ApoE genetic variants with obstructive sleep apnea in children. <i>Sleep Medicine</i> , 2008, 9, 260-265.	1.6	42
15	Common variants in 8q24 are associated with risk for prostate cancer and tumor aggressiveness in men of European ancestry. <i>Prostate</i> , 2009, 69, 1548-1556.	2.3	41
16	Tagging SNPs in the kallikrein genes $\text{K}1$ and 2 on 19q13 and their associations with prostate cancer in men of European origin. <i>Human Genetics</i> , 2007, 122, 251-259.	3.8	35
17	Genetic structure of Indian populations based on fifteen autosomal microsatellite loci. <i>BMC Genetics</i> , 2006, 7, 28.	2.7	28
18	Association of Tagging Single Nucleotide Polymorphisms on 8 Candidate Genes in Dopaminergic Pathway with Schizophrenia in Croatian Population. <i>Croatian Medical Journal</i> , 2009, 50, 361-369.	0.7	27

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19	Brain White Matter Development Is Associated with a Human-Specific Haplotype Increasing the Synthesis of Long Chain Fatty Acids. <i>Journal of Neuroscience</i> , 2014, 34, 6367-6376.	3.6	27
20	Molecular identification of lizard by RAPD & FINS of mitochondrial 16s rRNA gene. <i>Legal Medicine</i> , 2006, 8, 5-10.	1.3	26
21	Genetic variants associated with breast cancer risk for Ashkenazi Jewish women with strong family histories but no identifiable BRCA1/2 mutation. <i>Human Genetics</i> , 2013, 132, 523-536.	3.8	26
22	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 363-373.	1.7	25
23	Development of novel heminested PCR assays based on mitochondrial 16s rRNA gene for identification of seven pecora species. , 2005, 6, 42.		22
24	A schizophrenia risk gene, ZNF804A, is associated with brain white matter microstructure. <i>Schizophrenia Research</i> , 2014, 155, 15-20.	2.0	22
25	Concordance Study on 15 STR Loci in Three Major Populations of Himalayan State Sikkim. <i>Journal of Forensic Sciences</i> , 2002, 47, 1-5.	1.6	19
26	Excess of homozygosity in the major histocompatibility complex in schizophrenia. <i>Human Molecular Genetics</i> , 2014, 23, 6088-6095.	2.9	18
27	Genomic variation in the mitochondrially encoded cytochrome b (MT-CYB) and 16S rRNA (MT-RNR2) genes: characterization of eight endangered Pecoran species. <i>Animal Genetics</i> , 2006, 37, 262-265.	1.7	14
28	Genetic architecture of prostate cancer in the Ashkenazi Jewish population. <i>British Journal of Cancer</i> , 2011, 105, 864-869.	6.4	10
29	Novel Splice Mutation in Microphthalmia-Associated Transcription Factor in Waardenburg Syndrome. <i>Genetic Testing and Molecular Biomarkers</i> , 2011, 15, 525-529.	0.7	5
30	Concordance study on 15 STR loci in three major populations of Himalayan State Sikkim. <i>Journal of Forensic Sciences</i> , 2002, 47, 1163-7.	1.6	5
31	Addendum: Technical standards and guidelines for spinal muscular atrophy testing. <i>Genetics in Medicine</i> , 2021, 23, 2462.	2.4	2
32	Correlation Analyses Reveal a Substantial Influence of Allelic Gaps on the Investigation of Genetic Diversity of Modern Human Populations with Microsatellites. <i>Annals of Human Genetics</i> , 2008, 72, 644-653.	0.8	0
33	COMMON VARIANTS ON MLH1 GENE AND THEIR ASSOCIATIONS WITH PROSTATE CANCER IN EUROPEAN AMERICANS. <i>Journal of Urology</i> , 2008, 179, 227-227.	0.4	0
34	GENETIC VARIATION ON 8q24 AND SUSCEPTIBILITY TO PROSTATE CANCER IN CAUCASIAN MEN. <i>Journal of Urology</i> , 2008, 179, 226-226.	0.4	0
35	Effects of Inclusion of Relatives in DNA Databases: Empirical Observations from 13K SNPs in Hap-Map Population Data. <i>Handbook of Statistics</i> , 2012, , 353-366.	0.6	0