Saurav Guha

List of Publications by Year
in descending order

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	Best practices for the interpretation and reporting of clinical whole genome sequencing. Npj Genomic Medicine, 2022, 7, 27.	3.8	48
2	Addendum: Technical standards and guidelines for spinal muscular atrophy testing. Genetics in Medicine, 2021, 23, 2462.	2.4	2
3	Variant Classification Concordance using the ACMG-AMP Variant Interpretation Guidelines across Nine Genomic Implementation Research Studies. American Journal of Human Genetics, 2020, 107, 932-941.	6.2	51
4	Consensus interpretation of the p.Met34Thr and p.Val37Ile variants in GJB2 by the ClinGen Hearing Loss Expert Panel. Genetics in Medicine, 2019, 21, 2442-2452.	2.4	56
5	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 363-373.	1.7	25
6	Brain White Matter Development Is Associated with a Human-Specific Haplotype Increasing the Synthesis of Long Chain Fatty Acids. Journal of Neuroscience, 2014, 34, 6367-6376.	3.6	27
7	A schizophrenia risk gene, ZNF804A, is associated with brain white matter microstructure. Schizophrenia Research, 2014, 155, 15-20.	2.0	22
8	Molecular genetic evidence for overlap between general cognitive ability and risk for schizophrenia: a report from the Cognitive Genomics consorTium (COGENT). Molecular Psychiatry, 2014, 19, 168-174.	7.9	178
9	Genome-wide mapping of IBD segments in an Ashkenazi PD cohort identifies associated haplotypes. Human Molecular Genetics, 2014, 23, 4693-4702.	2.9	49
10	Sequencing an Ashkenazi reference panel supports population-targeted personal genomics and illuminates Jewish and European origins. Nature Communications, 2014, 5, 4835.	12.8	156
11	Excess of homozygosity in the major histocompatibility complex in schizophrenia. Human Molecular Genetics, 2014, 23, 6088-6095.	2.9	18
12	High rate of disease-related copy number variations in childhood onset schizophrenia. Molecular Psychiatry, 2014, 19, 568-572.	7.9	116
13	Genetic variants associated with breast cancer risk for Ashkenazi Jewish women with strong family histories but no identifiable BRCA1/2 mutation. Human Genetics, 2013, 132, 523-536.	3.8	26
14	Implication of a Rare Deletion at Distal 16p11.2 in Schizophrenia. JAMA Psychiatry, 2013, 70, 253.	11.0	69
15	Genome-wide association study implicates NDST3 in schizophrenia and bipolar disorder. Nature Communications, 2013, 4, 2739.	12.8	101
16	Implications for health and disease in the genetic signature of the Ashkenazi Jewish population. Genome Biology, 2012, 13, R2.	9.6	48
17	De novo copy number variants are associated with congenital diaphragmatic hernia. Journal of Medical Genetics, 2012, 49, 650-659.	3.2	68
18	Effects of Inclusion of Relatives in DNA Databases: Empirical Observations from 13K SNPs in Hap-Map Population Data. Handbook of Statistics, 2012, , 353-366.	0.6	0

#	Article	IF	Citations
19	Genetic architecture of prostate cancer in the Ashkenazi Jewish population. British Journal of Cancer, 2011, 105, 864-869.	6.4	10
20	A complete deficiency of Hyaluronoglucosaminidase 1 (HYAL1) presenting as familial juvenile idiopathic arthritis. Journal of Inherited Metabolic Disease, 2011, 34, 1013-1022.	3.6	68
21	Novel Splice Mutation in Microthalmia-Associated Transcription Factor in Waardenburg Syndrome. Genetic Testing and Molecular Biomarkers, 2011, 15, 525-529.	0.7	5
22	Common variants in 8q24 are associated with risk for prostate cancer and tumor aggressiveness in men of European ancestry. Prostate, 2009, 69, 1548-1556.	2.3	41
23	Association of Tagging Single Nucleotide Polymorphisms on 8 Candidate Genes in Dopaminergic Pathway with Schizophrenia in Croatian Population. Croatian Medical Journal, 2009, 50, 361-369.	0.7	27
24	Correlation Analyses Reveal a Substantial Influence of Allelic Gaps on the Investigation of Genetic Diversity of Modern Human Populations with Microsatellites. Annals of Human Genetics, 2008, 72, 644-653.	0.8	0
25	Association of ApoE genetic variants with obstructive sleep apnea in children. Sleep Medicine, 2008, 9, 260-265.	1.6	42
26	COMMON VARIANTS ON MLH1 GENE AND THEIR ASSOCIATIONS WITH PROSTATE CANCER IN EUROPEAN AMERICANS. Journal of Urology, 2008, 179, 227-227.	0.4	0
27	GENETIC VARIATION ON 8q24 AND SUSCEPTIBILITY TO PROSTATE CANCER IN CAUCASIAN MEN. Journal of Urology, 2008, 179, 226-226.	0.4	0
28	Molecular phylogeny of musk deer: A genomic view with mitochondrial 16S rRNA and cytochrome b gene. Molecular Phylogenetics and Evolution, 2007, 42, 585-597.	2.7	43
29	Tagging SNPs in the kallikrein genesÂ3 and 2 on 19q13 and their associations with prostate cancer in men of European origin. Human Genetics, 2007, 122, 251-259.	3.8	35
30	Genetic structure of Indian populations based on fifteen autosomal microsatellite loci. BMC Genetics, 2006, 7, 28.	2.7	28
31	Genomic variation in the mitochondrially encoded cytochrome b (MT-CYB) and 16S rRNA (MT-RNR2) genes: characterization of eight endangered Pecoran species. Animal Genetics, 2006, 37, 262-265.	1.7	14
32	Molecular identification of lizard by RAPD & FINS of mitochondrial 16s rRNA gene. Legal Medicine, 2006, 8, 5-10.	1.3	26
33	Development of novel heminested PCR assays based on mitochondrial 16s rRNA gene for identification of seven pecora species. , 2005, 6, 42.		22
34	Concordance Study on 15 STR Loci in Three Major Populations of Himalayan State Sikkim. Journal of Forensic Sciences, 2002, 47, 1-5.	1.6	19
35	Concordance study on 15 STR loci in three major populations of Himalayan State Sikkim. Journal of Forensic Sciences, 2002, 47, 1163-7.	1.6	5