

Hardik Shah

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1421486/publications.pdf>

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20
papers

1,552
citations

623734

14
h-index

839539

18
g-index

20
all docs

20
docs citations

20
times ranked

4534
citing authors

#	ARTICLE	IF	CITATIONS
1	Downregulation of exhausted cytotoxic T cells in gene expression networks of multisystem inflammatory syndrome in children. <i>Nature Communications</i> , 2021, 12, 4854.	12.8	42
2	Molecular Profiling of Coronavirus Disease 2019 (COVID-19) Autopsies Uncovers Novel Disease Mechanisms. <i>American Journal of Pathology</i> , 2021, 191, 2064-2071.	3.8	14
3	Elucidation of de novo small insertion/deletion biology with parentâ€œorigin phasing. <i>Human Mutation</i> , 2020, 41, 800-806.	2.5	3
4	Genetic associations with clozapine-induced myocarditis in patients with schizophrenia. <i>Translational Psychiatry</i> , 2020, 10, 37.	4.8	24
5	Double <i>PIK3CA</i> mutations in cis increase oncogenicity and sensitivity to PI3K inhibitors. <i>Science</i> , 2019, 366, 714-723.	12.6	185
6	O1.3. DIFFERENTIAL HISTONE MODIFICATIONS IN 250 SCHIZOPHRENIA CASES AND 330 CONTROLS. <i>Schizophrenia Bulletin</i> , 2019, 45, S159-S160.	4.3	0
7	High-coverage, long-read sequencing of Han Chinese trio reference samples. <i>Scientific Data</i> , 2019, 6, 91.	5.3	13
8	Neuronal impact of patient-specific aberrant <i>NRXN1</i> splicing. <i>Nature Genetics</i> , 2019, 51, 1679-1690.	21.4	91
9	A physical and genetic map of <i>Cannabis sativa</i> identifies extensive rearrangements at the <i>THC/CBD</i> acid synthase loci. <i>Genome Research</i> , 2019, 29, 146-156.	5.5	160
10	Cell-specific histone modification maps in the human frontal lobe link schizophrenia risk to the neuronal epigenome. <i>Nature Neuroscience</i> , 2018, 21, 1126-1136.	14.8	112
11	A functional genomics predictive network model identifies regulators of inflammatory bowel disease. <i>Nature Genetics</i> , 2017, 49, 1437-1449.	21.4	199
12	Analysis of 589,306 genomes identifies individuals resilient to severe Mendelian childhood diseases. <i>Nature Biotechnology</i> , 2016, 34, 531-538.	17.5	273
13	Development and clinical application of an integrative genomic approach to personalized cancer therapy. <i>Genome Medicine</i> , 2016, 8, 62.	8.2	71
14	How do students react to analyzing their own genomes in a whole-genome sequencing course?: outcomes of a longitudinal cohort study. <i>Genetics in Medicine</i> , 2015, 17, 866-874.	2.4	29
15	Preparing the next generation of genomicists: a laboratory-style course in medical genomics. <i>BMC Medical Genomics</i> , 2015, 8, 47.	1.5	16
16	AN INTEGRATIVE PIPELINE FOR MULTI-MODAL DISCOVERY OF DISEASE RELATIONSHIPS. , 2014, , .		15
17	Draft Genome Sequence of <i>Bacillus alcalophilus</i> AV1934, a Classic Alkaliphile Isolated from Human Feces in 1934. <i>Genome Announcements</i> , 2014, 2, .	0.8	5
18	microRNA-181a has a critical role in ovarian cancer progression through the regulation of the epithelialâ€œmesenchymal transition. <i>Nature Communications</i> , 2014, 5, 2977.	12.8	226

#	ARTICLE	IF	CITATIONS
19	Personalized Ovarian Cancer Disease Surveillance and Detection of Candidate Therapeutic Drug Target in Circulating Tumor DNA. <i>Neoplasia</i> , 2014, 16, 97-W29.	5.3	45
20	Informed decision-making among students analyzing their personal genomes on a whole genome sequencing course: a longitudinal cohort study. <i>Genome Medicine</i> , 2013, 5, 113.	8.2	29