

James Knight

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

2,304
citations

567281

15
h-index

610901

24
g-index

28
all docs

28
docs citations

28
times ranked

5587
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrative molecular and clinical profiling of acral melanoma links focal amplification of 22q11.21 to metastasis. <i>Nature Communications</i> , 2022, 13, 898.	12.8	19
2	Determining the serotype composition of mixed samples of pneumococcus using whole-genome sequencing. <i>Microbial Genomics</i> , 2021, 7, .	2.0	10
3	Genetics of agenesis/hypoplasia of the uterus and vagina: narrowing down the number of candidate genes for Mayer-Rokitansky-Kuster-Hauser Syndrome. <i>Human Genetics</i> , 2021, 140, 667-680.	3.8	16
4	<i>DIAPH1</i> Variants in Non-East Asian Patients With Sporadic Moyamoya Disease. <i>JAMA Neurology</i> , 2021, 78, 993.	9.0	33
5	Dibifunctional protein deficiency caused by splicing variants in a neonate with severe peroxisomal dysfunction and persistent hypoglycemia. <i>American Journal of Medical Genetics, Part A</i> , 2021, .	1.2	2
6	Long-read single molecule real-time (SMRT) sequencing of GBA1 locus in Gaucher disease national cohort from Argentina reveals high frequency of complex allele underlying severe skeletal phenotypes: Collaborative study from the Argentine Group for Diagnosis and Treatment of Gaucher Disease. <i>Molecular Genetics and Metabolism Reports</i> , 2021, 29, 100820.	1.1	3
7	Exome sequencing implicates genetic disruption of prenatal neuro-gliogenesis in sporadic congenital hydrocephalus. <i>Nature Medicine</i> , 2020, 26, 1754-1765.	30.7	84
8	Mutations disrupting neuritogenesis genes confer risk for cerebral palsy. <i>Nature Genetics</i> , 2020, 52, 1046-1056.	21.4	96
9	Exome Sequencing Implicates Impaired GABA Signaling and Neuronal Ion Transport in Trigeminal Neuralgia. <i>IScience</i> , 2020, 23, 101552.	4.1	32
10	The omentum of obese girls harbors small adipocytes and browning transcripts. <i>JCI Insight</i> , 2020, 5, .	5.0	8
11	Genomic sites hypersensitive to ultraviolet radiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 24196-24205.	7.1	66
12	The Use of Whole Exome Sequencing in a Cohort of Transgender Individuals to Identify Rare Genetic Variants. <i>Scientific Reports</i> , 2019, 9, 20099.	3.3	18
13	Insights into the evolution and drug susceptibility of <i>Babesia duncani</i> from the sequence of its mitochondrial and apicoplast genomes. <i>International Journal for Parasitology</i> , 2019, 49, 105-113.	3.1	13
14	Mutations in Chromatin Modifier and Ephrin Signaling Genes in Vein of Galen Malformation. <i>Neuron</i> , 2019, 101, 429-443.e4.	8.1	56
15	Hypokalemia Associated With a Claudin 10 Mutation: A Case Report. <i>American Journal of Kidney Diseases</i> , 2019, 73, 425-428.	1.9	24
16	Translocation of a gut pathobiont drives autoimmunity in mice and humans. <i>Science</i> , 2018, 359, 1156-1161.	12.6	608
17	Distinct adaptive mechanisms drive recovery from aneuploidy caused by loss of the Ulp2 SUMO protease. <i>Nature Communications</i> , 2018, 9, 5417.	12.8	21
18	De Novo Mutation in Genes Regulating Neural Stem Cell Fate in Human Congenital Hydrocephalus. <i>Neuron</i> , 2018, 99, 302-314.e4.	8.1	112

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19	Inflammasome Priming Mediated via Toll-Like Receptors 2 and 4, Induces Th1-Like Regulatory T Cells in De Novo Autoimmune Hepatitis. <i>Frontiers in Immunology</i> , 2018, 9, 1612.	4.8	16
20	Clonal evolution analysis of paired anaplastic and well-differentiated thyroid carcinomas reveals shared common ancestor. <i>Genes Chromosomes and Cancer</i> , 2018, 57, 645-652.	2.8	31
21	An improved genome assembly uncovers prolific tandem repeats in Atlantic cod. <i>BMC Genomics</i> , 2017, 18, 95.	2.8	153
22	Contribution of rare inherited and de novo variants in 2,871 congenital heart disease probands. <i>Nature Genetics</i> , 2017, 49, 1593-1601.	21.4	624
23	Integrated genomic characterization of IDH1-mutant glioma malignant progression. <i>Nature Genetics</i> , 2016, 48, 59-66.	21.4	253