Evan H Baugh

List of Publications by Year
in descending order

16	808	933447 10	940533 16
papers	citations	h-index	g-index

16	16	16	2106
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Why are there hotspot mutations in the TP53 gene in human cancers?. Cell Death and Differentiation, 2018, 25, 154-160.	11.2	393
2	Somatic SLC35A2 variants in the brain are associated with intractable neocortical epilepsy. Annals of Neurology, 2018, 83, 1133-1146.	5.3	95
3	Real-Time PyMOL Visualization for Rosetta and PyRosetta. PLoS ONE, 2011, 6, e21931.	2.5	55
4	Robust classification of protein variation using structural modelling and large-scale data integration. Nucleic Acids Research, 2016, 44, 2501-2513.	14.5	52
5	NBEA: Developmental disease gene with early generalized epilepsy phenotypes. Annals of Neurology, 2018, 84, 788-795.	5.3	44
6	Serine biosynthesis defect due to haploinsufficiency of PHGDH causes retinal disease. Nature Metabolism, 2021, 3, 366-377.	11.9	32
7	Role of <scp>WNT</scp>10A in failure of tooth development in humans and zebrafish. Molecular Genetics & Genomic Medicine, 2017, 5, 730-741.	1.2	27
8	Pre-detection history of extensively drug-resistant tuberculosis in KwaZulu-Natal, South Africa. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 23284-23291.	7.1	23
9	Genetic testing in individuals with cerebral palsy. Developmental Medicine and Child Neurology, 2021, 63, 1448-1455.	2.1	19
10	CSNK2B: A broad spectrum of neurodevelopmental disability and epilepsy severity. Epilepsia, 2021, 62, e103-e109.	5.1	13
11	Whole exome sequencing reveals potentially pathogenic variants in a small subset of premenopausal women with idiopathic osteoporosis. Bone, 2022, 154, 116253.	2.9	12
12	The benefit of diagnostic whole genome sequencing in schizophrenia and other psychotic disorders. Molecular Psychiatry, 2022, 27, 1435-1447.	7.9	12
13	A pathogenic variant in the <scp>SETBP1</scp> hotspot results in a formeâ€fruste <scp>Schinzelâ€“Giedion</scp> syndrome. American Journal of Medical Genetics, Part A, 2020, 182, 1947-1951.	1.2	11
14	New insights into tardive dyskinesia genetics: Implementation of whole-exome sequencing approach. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2019, 94, 109659.	4.8	9
15	Rare germline heterozygous missense variants in BRCA1-associated protein 1, BAP1, cause a syndromic neurodevelopmental disorder. American Journal of Human Genetics, 2022, 109, 361-372.	6.2	6
16	Truncating variants in the SHANK1 gene are associated with a spectrum of neurodevelopmental disorders. Genetics in Medicine, 2021, 23, 1912-1921.	2.4	5