Judith Conroy

List of Publications by Year
in descending order

16 papers	4,108 citations	759233 12 h-index	996975 15 g-index
16 all docs	16 docs citations	16 times ranked	7780 citing authors

#	Article	IF	Citations
1	OC48â€...Re-interrogation of whole exome sequencing data in developmental epileptic encephalopathies. , 2019, , .		0
2	Cost of exome sequencing in epileptic encephalopathy: is it â€˜worth itâ€™?. Archives of Disease in Childhood, 2018, 103, 304-304.	1.9	3
3	Symmetrical thalamic calcification: A trio whole exome sequencing negative series. Brain and Development, 2017, 39, 426-430.	1.1	1
4	Novel <scp>SMC</scp>1A variant and epilepsy of infancy with migrating focal seizures: Expansion of the phenotype. Epilepsia, 2017, 58, 1301-1302.	5.1	12
5	Clinical spectrum and genotypeâ€“phenotype associations of KCNA2-related encephalopathies. Brain, 2017, 140, 2337-2354.	7.6	117
6	Unexplained early onset epileptic encephalopathy: Exome screening and phenotype expansion. Epilepsia, 2016, 57, e12-7.	5.1	164
7	Novel European SLC1A4 variant: infantile spasms and population ancestry analysis. Journal of Human Genetics, 2016, 61, 761-764.	2.3	18
8	Atypical benign partial epilepsy of childhood with acquired neurocognitive, lexical semantic, and autistic spectrum disorder. Epilepsy & Behavior Case Reports, 2016, 6, 42-48.	1.5	10
9	Chromosomal microarray in unexplained severe early onset epilepsy â€“ A single centre cohort. European Journal of Paediatric Neurology, 2015, 19, 390-394.	1.6	14
10	The variable phenotypes of <scp>KCNQ</scp>â€related epilepsy. Epilepsia, 2014, 55, e99-105.	5.1	109
11	TDP2 protects transcription from abortive topoisomerase activity and is required for normal neural function. Nature Genetics, 2014, 46, 516-521.	21.4	122
12	Convergence of Genes and Cellular Pathways Dysregulated in Autism Spectrum Disorders. American Journal of Human Genetics, 2014, 94, 677-694.	6.2	819
13	Towards the identification of a genetic basis for <scp>L</scp>andauâ€<scp>K</scp>leffner <scp>s</scp>yndrome. Epilepsia, 2014, 55, 858-865.	5.1	44
14	Individual common variants exert weak effects on the risk for autism spectrum disorders. Human Molecular Genetics, 2012, 21, 4781-4792.	2.9	334
15	Functional impact of global rare copy number variation in autism spectrum disorders. Nature, 2010, 466, 368-372.	27.8	1,803
16	A genome-wide scan for common alleles affecting risk for autism. Human Molecular Genetics, 2010, 19, 4072-4082.	2.9	538