

Sarah E Heron

List of Publications by Year in descending order

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53
papers

5,182
citations

136950

32
h-index

182427

51
g-index

53
all docs

53
docs citations

53
times ranked

4856
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of <i>SLC32A1</i> Missense Variants With Genetic Epilepsy With Febrile Seizures Plus. <i>Neurology</i> , 2021, 96, e2251-e2260.	1.1	13
2	Integrated in silico and experimental assessment of disease relevance of <i>PCDH19</i> missense variants. <i>Human Mutation</i> , 2021, 42, 1030-1041.	2.5	1
3	Two novel intragenic variants in the <i>FMR1</i> gene in patients with suspect clinical diagnosis of Fragile X syndrome and no CGG repeat expansion. <i>European Journal of Medical Genetics</i> , 2020, 63, 104010.	1.3	4
4	Benign infantile seizures and paroxysmal dyskinesia caused by an <i>SCN8A</i> mutation. <i>Annals of Neurology</i> , 2016, 79, 428-436.	5.3	159
5	<i>BRAT1</i> associated neurodegeneration: Intrafamilial phenotypic differences in siblings. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 3033-3038.	1.2	18
6	Reply. <i>Annals of Neurology</i> , 2016, 80, 168-169.	5.3	0
7	<i>KCNT1</i> mutations in seizure disorders: the phenotypic spectrum and functional effects. <i>Journal of Medical Genetics</i> , 2016, 53, 217-225.	3.2	94
8	Multiplex families with epilepsy. <i>Neurology</i> , 2016, 86, 713-722.	1.1	23
9	Mutations in the mammalian target of rapamycin pathway regulators <i>NPRL2</i> and <i>NPRL3</i> cause focal epilepsy. <i>Annals of Neurology</i> , 2016, 79, 120-131.	5.3	190
10	Familial neonatal seizures in 36 families: Clinical and genetic features correlate with outcome. <i>Epilepsia</i> , 2015, 56, 1071-1080.	5.1	94
11	Single Nucleotide Variations in <i>CLCN6</i> Identified in Patients with Benign Partial Epilepsies in Infancy and/or Febrile Seizures. <i>PLoS ONE</i> , 2015, 10, e0118946.	2.5	13
12	Mutations in <i>KCNT1</i> cause a spectrum of focal epilepsies. <i>Epilepsia</i> , 2015, 56, e114-20.	5.1	117
13	A recurrent de novo mutation in <i>KCNC1</i> causes progressive myoclonus epilepsy. <i>Nature Genetics</i> , 2015, 47, 39-46.	21.4	245
14	<i>KCNT1</i> gain of function in 2 epilepsy phenotypes is reversed by quinidine. <i>Annals of Neurology</i> , 2014, 75, 581-590.	5.3	249
15	Mutations in mammalian target of rapamycin regulator <i>DEPDC5</i> cause focal epilepsy with brain malformations. <i>Annals of Neurology</i> , 2014, 75, 782-787.	5.3	193
16	Genetics of epilepsy. <i>Neurology</i> , 2014, 83, 1042-1048.	1.1	61
17	A variant of <i>KCC2</i> from patients with febrile seizures impairs neuronal Cl ⁻ extrusion and dendritic spine formation. <i>EMBO Reports</i> , 2014, 15, 723-729.	4.5	163
18	Mutations in <i>DEPDC5</i> cause familial focal epilepsy with variable foci. <i>Nature Genetics</i> , 2013, 45, 546-551.	21.4	301

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19	Role of <i>PRRT2</i> in common paroxysmal neurological disorders: a gene with remarkable pleiotropy. <i>Journal of Medical Genetics</i> , 2013, 50, 133-139.	3.2	88
20	Autosomal dominant vasovagal syncope. <i>Neurology</i> , 2013, 80, 1485-1493.	1.1	20
21	Mutations in <i>PRRT2</i> are not a common cause of infantile epileptic encephalopathies. <i>Epilepsia</i> , 2013, 54, e86-9.	5.1	12
22	Benign Neonatal Sleep Myoclonus. <i>Journal of Child Neurology</i> , 2012, 27, 1260-1263.	1.4	10
23	Missense mutations in the sodium-gated potassium channel gene <i>KCNT1</i> cause severe autosomal dominant nocturnal frontal lobe epilepsy. <i>Nature Genetics</i> , 2012, 44, 1188-1190.	21.4	333
24	Familial focal epilepsy with variable foci mapped to chromosome 22q12: Expansion of the phenotypic spectrum. <i>Epilepsia</i> , 2012, 53, e151-5.	5.1	24
25	<i>PRRT2</i> phenotypic spectrum includes sporadic and fever-related infantile seizures. <i>Neurology</i> , 2012, 79, 2104-2108.	1.1	75
26	<i>PRRT2</i> Mutations Cause Benign Familial Infantile Epilepsy and Infantile Convulsions with Choreoathetosis Syndrome. <i>American Journal of Human Genetics</i> , 2012, 90, 152-160.	6.2	234
27	<i>KCNQ2</i> encephalopathy: Emerging phenotype of a neonatal epileptic encephalopathy. <i>Annals of Neurology</i> , 2012, 71, 15-25.	5.3	427
28	The Role of Seizure-Related <i>SEZ6</i> as a Susceptibility Gene in Febrile Seizures. <i>Neurology Research International</i> , 2011, 2011, 1-4.	1.3	20
29	Proposed genetic classification of the "benign" familial neonatal and infantile epilepsies. <i>Epilepsia</i> , 2011, 52, 649-650.	5.1	9
30	"Blinders, phenotype, and fashionable genetic analysis" Setting the record straight for epilepsy!. <i>Epilepsia</i> , 2011, 52, 1757-1758.	5.1	2
31	Neonatal seizures and long QT Syndrome: A cardiocerebral channelopathy?. <i>Epilepsia</i> , 2010, 51, 293-296.	5.1	61
32	Familial neonatal seizures with intellectual disability caused by a microduplication of chromosome 2q24.3. <i>Epilepsia</i> , 2010, 51, 1865-1869.	5.1	30
33	De novo <i>SCN1A</i> mutations in Dravet syndrome and related epileptic encephalopathies are largely of paternal origin. <i>Journal of Medical Genetics</i> , 2010, 47, 137-141.	3.2	44
34	Novel Mutation in <i>KCNQ2</i> Causing Benign Familial Neonatal Seizures. <i>Pediatric Neurology</i> , 2009, 41, 367-370.	2.1	10
35	Severe autosomal dominant nocturnal frontal lobe epilepsy associated with psychiatric disorders and intellectual disability. <i>Epilepsia</i> , 2008, 49, 2125-2129.	5.1	49
36	Deletions or duplications in <i>KCNQ2</i> can cause benign familial neonatal seizures. <i>Journal of Medical Genetics</i> , 2007, 44, 791-796.	3.2	70

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37	Generalized epilepsy with febrile seizures plus-associated sodium channel β 1 subunit mutations severely reduce β subunit-mediated modulation of sodium channel function. <i>Neuroscience</i> , 2007, 148, 164-174.	2.3	61
38	A childhood epilepsy mutation reveals a role for developmentally regulated splicing of a sodium channel. <i>Molecular and Cellular Neurosciences</i> , 2007, 35, 292-301.	2.2	68
39	Extended spectrum of idiopathic generalized epilepsies associated with <i>CACNA1H</i> functional variants. <i>Annals of Neurology</i> , 2007, 62, 560-568.	5.3	186
40	Channelopathies in idiopathic epilepsy. <i>Neurotherapeutics</i> , 2007, 4, 295-304.	4.4	101
41	Association studies and functional validation or functional validation alone?. <i>Epilepsy Research</i> , 2007, 74, 237-238.	1.6	3
42	SCN2A Mutations and Benign Familial Neonatal-Infantile Seizures: The Phenotypic Spectrum. <i>Epilepsia</i> , 2007, 48, 1138-1142.	5.1	102
43	A polygenic heterogeneity model for common epilepsies with complex genetics. <i>Genes, Brain and Behavior</i> , 2007, 6, 593-597.	2.2	52
44	Is Variation in the GABA(B) Receptor 1 Gene Associated with Temporal Lobe Epilepsy?. <i>Epilepsia</i> , 2005, 46, 778-780.	5.1	15
45	Novel mutations in the KCNQ2 gene link epilepsy to a dysfunction of the KCNQ2-calmodulin interaction. <i>Journal of Medical Genetics</i> , 2004, 41, 35e-35.	3.2	62
46	Generalized Epilepsy with Febrile Seizures Plus (GEFS+): Clinical Spectrum in Seven Italian Families Unrelated to SCN1A, SCN1B, and GABRG2 Gene Mutations. <i>Epilepsia</i> , 2004, 45, 149-158.	5.1	67
47	Febrile Convulsions and Genetic Susceptibility: Role of the Neuronal Nicotinic Acetylcholine Receptor α 4 Subunit. <i>Epilepsia</i> , 2004, 45, 561-561.	5.1	10
48	Genetic variation of <i>CACNA1H</i> in idiopathic generalized epilepsy. <i>Annals of Neurology</i> , 2004, 55, 595-596.	5.3	102
49	Benign familial neonatal-infantile seizures: Characterization of a new sodium channelopathy. <i>Annals of Neurology</i> , 2004, 55, 550-557.	5.3	250
50	Sodium-channel defects in benign familial neonatal-infantile seizures. <i>Lancet, The</i> , 2002, 360, 851-852.	13.7	332
51	CHRNA2 Is the Second Acetylcholine Receptor Subunit Associated with Autosomal Dominant Nocturnal Frontal Lobe Epilepsy*. <i>American Journal of Human Genetics</i> , 2001, 68, 225-231.	6.2	300
52	Absence of Mutations Raises Doubts about the Role of the 70-kD Peroxisomal Membrane Protein in Zellweger Syndrome. <i>American Journal of Human Genetics</i> , 1997, 60, 1535-1539.	6.2	15
53	Absence of Mutations Raises Doubts about the Role of the 70-kD Peroxisomal Membrane Protein in Zellweger Syndrome. <i>American Journal of Human Genetics</i> , 1997, 60, 1535-1538.	6.2	0