Anuranjan Anand

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/2195797/publications.pdf

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19	411	933447	888059
papers	citations	10 h-index	17 g-index

20	20	20	706
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Functional consequences of novel connexin 26 mutations associated with hereditary hearing loss. European Journal of Human Genetics, 2009, 17, 502-509.	2.8	66
2	An idiopathic epilepsy syndrome linked to 3q13.3â€q21 and missense mutations in the extracellular calcium sensing receptor gene. Annals of Neurology, 2008, 64, 158-167.	5.3	65
3	Non-Syndromic Hearing Impairment in India: High Allelic Heterogeneity among Mutations in TMPRSS3, TMC1, USHIC, CDH23 and TMIE. PLoS ONE, 2014, 9, e84773.	2.5	50
4	The polyglutamine motif is highly conserved at the Clock locus in various organisms and is not polymorphic in humans. Human Genetics, 2001, 109, 136-142.	3.8	49
5	A locus for autosomal dominant reflex epilepsy precipitated by hot water maps at chromosome 10q21.3-q22.3. Human Genetics, 2009, 125, 541-549.	3.8	40
6	Familial autosomal dominant reflex epilepsy triggered by hot water maps to 4q24-q28. Human Genetics, 2009, 126, 677-683.	3.8	37
7	Association analysis of CAG repeats at theKCNN3 locus in Indian patients with bipolar disorder and schizophrenia. American Journal of Medical Genetics Part A, 2000, 96, 744-748.	2.4	26
8	A locus for juvenile myoclonic epilepsy maps to 2q33â€“q36. Human Genetics, 2010, 128, 123-130.	3.8	14
9	A novel genetic locus for juvenile myoclonic epilepsy at chromosome 5q12â€“q14. Human Genetics, 2007, 121, 655-662.	3.8	12
10	Rare SLC1A1 variants in hot water epilepsy. Human Genetics, 2017, 136, 693-703.	3.8	12
11	Mutations in & genes as major causes of hearing impairment in Dhadkai village, Jammu & Kashmir, India. Indian Journal of Medical Research, 2017, 146, 489-497.	1.0	9
12	Microtubule-associated defects caused by EFHC1 mutations in juvenile myoclonic epilepsy. Human Mutation, 2017, 38, 816-826.	2.5	7
13	A novel locus DFNA59 for autosomal dominant nonsyndromic hearing loss maps at chromosome 11p14.2â€“q12.3. Human Genetics, 2009, 124, 669-675.	3.8	6
14	A genetic locus for sensory epilepsy precipitated by contact with hot water maps to chromosome 9p24.3-p23. Journal of Genetics, 2018, 97, 391-398.	0.7	6
15	A linkage and exome study implicates rare variants of KANK4 and CAP2 in bipolar disorder in a multiplex family. Bipolar Disorders, 2020, 22, 70-78.	1.9	6
16	Identification of a novel homozygous mutation in transmembrane channel like 1 () gene, one of the second-tier hearing loss genes after in India. Indian Journal of Medical Research, 2017, 145, 492-497.	1.0	3
17	Functional Analysis of a Novel Connexin30 Mutation in a Large Family with Hearing Loss, Pesplanus, Ichthyosis, Cutaneous Nodules, and Keratoderma. Annals of Human Genetics, 2016, 80, 11-19.	0.8	2
18	A genetic locus for sensory epilepsy precipitated by contact with hot water maps to chromosome 9p24.3-p23. Journal of Genetics, 2018, 97, 391-398.	0.7	1

#	Article	IF	Citations
19	Sex determining signal inDrosophila melanogaster. Journal of Genetics, 2004, 83, 121-123.	0.7	0