

Uppala Radhakrishna

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

1,932
citations

567281

15
h-index

996975

15
g-index

16
all docs

16
docs citations

16
times ranked

1969
citing authors

#	ARTICLE	IF	CITATIONS
1	Schizophrenia susceptibility loci on chromosomes 13q32 and 8p21. <i>Nature Genetics</i> , 1998, 20, 70-73.	21.4	506
2	Mutations in GJB6 cause hidrotic ectodermal dysplasia. <i>Nature Genetics</i> , 2000, 26, 142-144.	21.4	270
3	Insertion of \hat{I}^2 -satellite repeats identifies a transmembrane protease causing both congenital and childhood onset autosomal recessive deafness. <i>Nature Genetics</i> , 2001, 27, 59-63.	21.4	208
4	The Phenotypic Spectrum of GLI3 Morphopathies Includes Autosomal Dominant Preaxial Polydactyly Type-IV and Postaxial Polydactyly Type-A/B; No Phenotype Prediction from the Position of GLI3 Mutations. <i>American Journal of Human Genetics</i> , 1999, 65, 645-655.	6.2	175
5	Mutation in GLI3 in postaxial polydactyly type A. <i>Nature Genetics</i> , 1997, 17, 269-271.	21.4	163
6	Peutz-Jeghers Syndrome: Confirmation of Linkage to Chromosome 19p13.3 and Identification of a Potential Second Locus, on 19q13.4. <i>American Journal of Human Genetics</i> , 1997, 61, 1327-1334.	6.2	145
7	A genomic rearrangement resulting in a tandem duplication is associated with split hand-split foot malformation 3 (SHFM3) at 10q24. <i>Human Molecular Genetics</i> , 2003, 12, 1959-1971.	2.9	88
8	Duplications of <i>BHLHA9</i> are associated with ectrodactyly and tibia hemimelia inherited in non-Mendelian fashion. <i>Journal of Medical Genetics</i> , 2012, 49, 119-125.	3.2	81
9	Molecular and Clinical Characteristics in 46 Families Affected with Peutz-Jeghers Syndrome. <i>Digestive Diseases and Sciences</i> , 2007, 52, 1924-1933.	2.3	56
10	Novel mutations of TMRSS3 in four DFNB8/B10 families segregating congenital autosomal recessive deafness. <i>Journal of Medical Genetics</i> , 2001, 38, 396-400.	3.2	55
11	Clouston hidrotic ectodermal dysplasia (HED): genetic homogeneity, presence of a founder effect in the French Canadian population and fine genetic mapping. <i>European Journal of Human Genetics</i> , 2000, 8, 372-380.	2.8	43
12	Lack of linkage or association between schizophrenia and the polymorphic trinucleotide repeat within the KCNN3 gene on chromosome 1q21. , 1999, 88, 348-351.		33
13	Split-hand/split-foot malformation 3 (SHFM3) at 10q24, development of rapid diagnostic methods and gene expression from the region. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 1384-1395.	1.2	33
14	Genomewide Linkage Scan for Split-Hand/Foot Malformation with Long-Bone Deficiency in a Large Arab Family Identifies Two Novel Susceptibility Loci on Chromosomes 1q42.2-q43 and 6q14.1. <i>American Journal of Human Genetics</i> , 2007, 80, 105-111.	6.2	30
15	Genomewide Scan for Nonsyndromic Cleft Lip and Palate in Multigenerational Indian Families Reveals Significant Evidence of Linkage at 13q33.1-34. <i>American Journal of Human Genetics</i> , 2006, 79, 580-585.	6.2	29
16	Autosomal Dominant Nonsyndromic Cleft Lip and Palate: Significant Evidence of Linkage at 18q21.1. <i>American Journal of Human Genetics</i> , 2007, 81, 180-188.	6.2	17