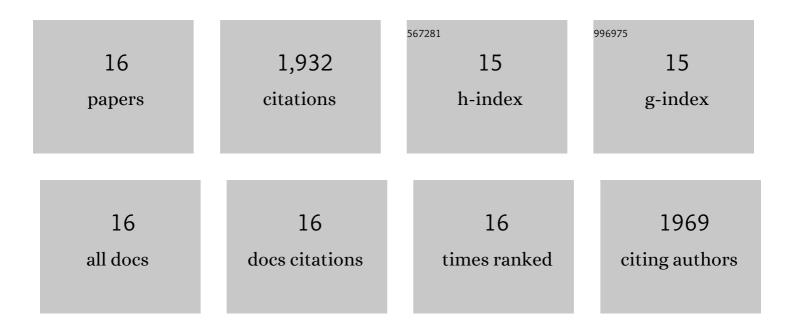
Uppala Radhakrishna

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12140585/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Duplications of <i>BHLHA9</i> are associated with ectrodactyly and tibia hemimelia inherited in non-Mendelian fashion. Journal of Medical Genetics, 2012, 49, 119-125.	3.2	81
2	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. American Journal of Human Genetics, 2007, 80, 105-111.	6.2	30
3	Autosomal Dominant Nonsyndromic Cleft Lip and Palate: Significant Evidence of Linkage at 18q21.1. American Journal of Human Genetics, 2007, 81, 180-188.	6.2	17
4	Molecular and Clinical Characteristics in 46 Families Affected with Peutz–Jeghers Syndrome. Digestive Diseases and Sciences, 2007, 52, 1924-1933.	2.3	56
5	Genomewide Scan for Nonsyndromic Cleft Lip and Palate in Multigenerational Indian Families Reveals Significant Evidence of Linkage at 13q33.1-34. American Journal of Human Genetics, 2006, 79, 580-585.	6.2	29
6	Split-hand/split-foot malformation 3 (SHFM3) at 10q24, development of rapid diagnostic methods and gene expression from the region. American Journal of Medical Genetics, Part A, 2006, 140A, 1384-1395.	1.2	33
7	A genomic rearrangement resulting in a tandem duplication is associated with split hand-split foot malformation 3 (SHFM3) at 10q24. Human Molecular Genetics, 2003, 12, 1959-1971.	2.9	88
8	Insertion of β-satellite repeats identifies a transmembrane protease causing both congenital and childhood onset autosomal recessive deafness. Nature Genetics, 2001, 27, 59-63.	21.4	208
9	Novel mutations of TMPRSS3 in four DFNB8/B10 families segregating congenital autosomal recessive deafness. Journal of Medical Genetics, 2001, 38, 396-400.	3.2	55
10	Mutations in GJB6 cause hidrotic ectodermal dysplasia. Nature Genetics, 2000, 26, 142-144.	21.4	270
11	Clouston hidrotic ectodermal dysplasia (HED): genetic homogeneity, presence of a founder effect in the French Canadian population and fine genetic mapping. European Journal of Human Genetics, 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	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. American Journal of Human Genetics, 1999, 65, 645-655.	6.2	175
14	Schizophrenia susceptibility loci on chromosomes 13q32 and 8p21. Nature Genetics, 1998, 20, 70-73.	21.4	506
15	Peutz-Jeghers Syndrome: Confirmation of Linkage to Chromosome 19p13.3 and Identification of a Potential Second Locus, on 19q13.4. American Journal of Human Genetics, 1997, 61, 1327-1334.	6.2	145
16	Mutation in GLI3 in postaxial polydactyly type A. Nature Genetics, 1997, 17, 269-271.	21.4	163