

# Katta Mohan Girisha

## List of Publications by Year in descending order

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245  
papers

5,238  
citations

136885

32  
h-index

128225

60  
g-index

252  
all docs

252  
docs citations

252  
times ranked

10379  
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterozygous STAT1 gain-of-function mutations underlie an unexpectedly broad clinical phenotype. <i>Blood</i> , 2016, 127, 3154-3164.	0.6	465
2	Biallelic mutations in the death domain of PIDD1 impair caspase-2 activation and are associated with intellectual disability. <i>Translational Psychiatry</i> , 2021, 11, 1.	2.4	334
3	Characterization of Greater Middle Eastern genetic variation for enhanced disease gene discovery. <i>Nature Genetics</i> , 2016, 48, 1071-1076.	9.4	314
4	Association of <i>MTOR</i> Mutations With Developmental Brain Disorders, Including Megalencephaly, Focal Cortical Dysplasia, and Pigmentary Mosaicism. <i>JAMA Neurology</i> , 2016, 73, 836.	4.5	234
5	Genetic heterogeneity in Cornelia de Lange syndrome (CdLS) and CdLS-like phenotypes with observed and predicted levels of mosaicism. <i>Journal of Medical Genetics</i> , 2014, 51, 659-668.	1.5	141
6	RSPO2 inhibition of RNF43 and ZNRF3 governs limb development independently of LGR4/5/6. <i>Nature</i> , 2018, 557, 564-569.	13.7	141
7	PIK3CA-associated developmental disorders exhibit distinct classes of mutations with variable expression and tissue distribution. <i>JCI Insight</i> , 2016, 1, .	2.3	134
8	The promise of discovering population-specific disease-associated genes in South Asia. <i>Nature Genetics</i> , 2017, 49, 1403-1407.	9.4	129
9	Cortical-Bone Fragility Insights from sFRP4 Deficiency in Pyle's Disease. <i>New England Journal of Medicine</i> , 2016, 374, 2553-2562.	13.9	119
10	Genomic and phenotypic delineation of congenital microcephaly. <i>Genetics in Medicine</i> , 2019, 21, 545-552.	1.1	85
11	Down syndrome in diverse populations. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 42-53.	0.7	75
12	The Genomics of Arthrogyrosis, a Complex Trait: Candidate Genes and Further Evidence for Oligogenic Inheritance. <i>American Journal of Human Genetics</i> , 2019, 105, 132-150.	2.6	74
13	Down syndrome. <i>Archives of Medical Research</i> , 2004, 35, 31-35.	1.5	72
14	Microduplications encompassing the Sonic hedgehog limb enhancer <i>ZRS</i> are associated with Haas-type polysyndactyly and Laurin-Sandrow syndrome. <i>Clinical Genetics</i> , 2014, 86, 318-325.	1.0	72
15	A dyadic approach to the delineation of diagnostic entities in clinical genomics. <i>American Journal of Human Genetics</i> , 2021, 108, 8-15.	2.6	71
16	Phenotypes and genotypes in individuals with <i>SMC1A</i> variants. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2108-2125.	0.7	69
17	A homozygous nonsense variant in <i>IFT52</i> is associated with a human skeletal ciliopathy. <i>Clinical Genetics</i> , 2016, 90, 536-539.	1.0	68
18	Noonan syndrome in diverse populations. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2323-2334.	0.7	68

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19	Biallelic Loss of Proprioception-Related PIEZO2 Causes Muscular Atrophy with Perinatal Respiratory Distress, Arthrogyposis, and Scoliosis. <i>American Journal of Human Genetics</i> , 2016, 99, 1206-1216.	2.6	65
20	Phenotypic spectrum and prevalence of INPP5E mutations in Joubert Syndrome and related disorders. <i>European Journal of Human Genetics</i> , 2013, 21, 1074-1078.	1.4	64
21	Analysis of the <i>WISP3</i> gene in Indian families with progressive pseudorheumatoid dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 2820-2828.	0.7	63
22	Mutations in EBF3 Disturb Transcriptional Profiles and Cause Intellectual Disability, Ataxia, and Facial Dysmorphism. <i>American Journal of Human Genetics</i> , 2017, 100, 117-127.	2.6	62
23	Genotype-phenotype spectrum of PYCR1-related autosomal recessive cutis laxa. <i>Molecular Genetics and Metabolism</i> , 2013, 110, 352-361.	0.5	57
24	Williams-Beuren syndrome in diverse populations. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1128-1136.	0.7	55
25	The homozygous variant c.797G>A/p.(Cys266Tyr) in <i>PISD</i> is associated with a Spondyloepimetaphyseal dysplasia with large epiphyses and disturbed mitochondrial function. <i>Human Mutation</i> , 2019, 40, 299-309.	1.1	54
26	Homozygous p.(Glu87Lys) variant in ISCA1 is associated with a multiple mitochondrial dysfunctions syndrome. <i>Journal of Human Genetics</i> , 2017, 62, 723-727.	1.1	53
27	NAD(P)HX dehydratase (NAXD) deficiency: a novel neurodegenerative disorder exacerbated by febrile illnesses. <i>Brain</i> , 2019, 142, 50-58.	3.7	51
28	The SMAD-binding domain of SKI: a hotspot for de novo mutations causing Shprintzen-Goldberg syndrome. <i>European Journal of Human Genetics</i> , 2015, 23, 224-228.	1.4	48
29	Cornelia de Lange syndrome in diverse populations. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 150-158.	0.7	40
30	Diagnostic strategies and genotype-phenotype correlation in a large Indian cohort of osteogenesis imperfecta. <i>Bone</i> , 2018, 110, 368-377.	1.4	38
31	Mutations in patients with osteogenesis imperfecta from consanguineous Indian families. <i>European Journal of Medical Genetics</i> , 2015, 58, 21-27.	0.7	37
32	SCUBE3 loss-of-function causes a recognizable recessive developmental disorder due to defective bone morphogenetic protein signaling. <i>American Journal of Human Genetics</i> , 2021, 108, 115-133.	2.6	37
33	Genetic abnormalities in a large cohort of Coffin-Siris syndrome patients. <i>Journal of Human Genetics</i> , 2019, 64, 1173-1186.	1.1	36
34	Glycosaminoglycan levels in dried blood spots of patients with mucopolysaccharidoses and mucopolipidoses. <i>Molecular Genetics and Metabolism</i> , 2017, 120, 247-254.	0.5	35
35	Clinical utility of fetal autopsy and its impact on genetic counseling. <i>Prenatal Diagnosis</i> , 2015, 35, 685-691.	1.1	34
36	Identification of a novel LRRK1 mutation in a family with osteosclerotic metaphyseal dysplasia. <i>Journal of Human Genetics</i> , 2017, 62, 437-441.	1.1	33

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37	Mutations in SREBF1, Encoding Sterol Regulatory Element Binding Transcription Factor 1, Cause Autosomal-Dominant IFAP Syndrome. American Journal of Human Genetics, 2020, 107, 34-45.	2.6	33
38	<i>GALNS</i> mutations in Indian patients with mucopolysaccharidosis IVA. American Journal of Medical Genetics, Part A, 2014, 164, 2793-2801.	0.7	31
39	Clinical and mutation profile of multicentric osteolysis nodulosis and arthropathy. American Journal of Medical Genetics, Part A, 2016, 170, 410-417.	0.7	31
40	Organization for rare diseases India (ORDI)â€™â€™addressing the challenges and opportunities for the Indian rare diseases' community. Genetical Research, 2014, 96, e009.	0.3	30
41	Expanding the phenotype associated with 17q12 duplication: Case report and review of the literature. American Journal of Medical Genetics, Part A, 2013, 161, 352-359.	0.7	29
42	BGN Mutations in X-Linked Spondyloepimetaphyseal Dysplasia. American Journal of Human Genetics, 2016, 98, 1243-1248.	2.6	29
43	Autosomal recessive spinocerebellar ataxia 20: Report of a new patient and review of literature. European Journal of Medical Genetics, 2017, 60, 118-123.	0.7	29
44	Novel subtype of mucopolysaccharidosis caused by arylsulfatase K (ARSK) deficiency. Journal of Medical Genetics, 2022, 59, 957-964.	1.5	29
45	A novel sequence variant in SFRP4 causing Pyle disease. Journal of Human Genetics, 2017, 62, 575-576.	1.1	27
46	Gaucher disease: single gene molecular characterization of one-hundred Indian patients reveals novel variants and the most prevalent mutation. BMC Medical Genetics, 2019, 20, 31.	2.1	27
47	Homozygosity for a nonsense variant in AIMP2 is associated with a progressive neurodevelopmental disorder with microcephaly, seizures, and spastic quadriplegia. Journal of Human Genetics, 2018, 63, 19-25.	1.1	26
48	A data set of variants derived from 1455 clinical and research exomes is efficient in variant prioritization for early-onset monogenic disorders in Indians. Human Mutation, 2021, 42, e15-e61.	1.1	25
49	Mutation spectrum of <i>COL1A1</i> and <i>COL1A2</i> genes in Indian patients with osteogenesis imperfecta. American Journal of Medical Genetics, Part A, 2014, 164, 1482-1489.	0.7	24
50	Identification and characterization of 20 novel pathogenic variants in 60 unrelated Indian patients with mucopolysaccharidoses type I and type <sc>II</sc>. Clinical Genetics, 2016, 90, 496-508.	1.0	23
51	Human IFT52 mutations uncover a novel role for the protein in microtubule dynamics and centrosome cohesion. Human Molecular Genetics, 2019, 28, 2720-2737.	1.4	23
52	Costello syndrome with severe cutis laxa and mosaic <i>HRAS</i> G12S mutation. American Journal of Medical Genetics, Part A, 2010, 152A, 2861-2864.	0.7	22
53	Recurrent and novel GLB1 mutations in India. Gene, 2015, 567, 173-181.	1.0	22
54	Clinical and genetic spectrum of AMPD2-related pontocerebellar hypoplasia type 9. European Journal of Human Genetics, 2018, 26, 695-708.	1.4	22

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55	Novel and recurrent mutations in <i>WISP3</i> and an atypical phenotype. American Journal of Medical Genetics, Part A, 2015, 167, 2481-2484.	0.7	21
56	Novel KIAA0753 mutations extend the phenotype of skeletal ciliopathies. Scientific Reports, 2017, 7, 15585.	1.6	21
57	Mutations in MYLPF Cause a Novel Segmental Amyoplasia that Manifests as Distal Arthrogryposis. American Journal of Human Genetics, 2020, 107, 293-310.	2.6	21
58	T1 and M1 polymorphism in glutathione S-transferase gene and coronary artery disease in North Indian population. Indian Journal of Medical Sciences, 2004, 58, 520-6.	0.1	20
59	A neurodegenerative mitochondrial disease phenotype due to biallelic loss-of-function variants in <i>PNPLA8</i> encoding calcium-independent phospholipase A2 <sup>3</sup> . American Journal of Medical Genetics, Part A, 2018, 176, 1232-1237.	0.7	19
60	High diagnostic yield in skeletal ciliopathies using massively parallel genome sequencing, structural variant screening and RNA analyses. Journal of Human Genetics, 2021, 66, 995-1008.	1.1	19
61	Eight years experience from a skeletal dysplasia referral center in a tertiary hospital in Southern India: A model for the diagnosis and treatment of rare diseases in a developing country. American Journal of Medical Genetics, Part A, 2014, 164, 2317-2323.	0.7	18
62	Phenotype and genotype in patients with Larsen syndrome: clinical homogeneity and allelic heterogeneity in seven patients. BMC Medical Genetics, 2016, 17, 27.	2.1	18
63	Second family provides further evidence for causation of Steel syndrome by biallelic mutations in <i>COL27A1</i> . Clinical Genetics, 2017, 92, 323-326.	1.0	18
64	Bi-allelic TMEM94 Truncating Variants Are Associated with Neurodevelopmental Delay, Congenital Heart Defects, and Distinct Facial Dysmorphism. American Journal of Human Genetics, 2018, 103, 948-967.	2.6	18
65	FLNA mutations in surviving males presenting with connective tissue findings: two new case reports and review of the literature. BMC Medical Genetics, 2018, 19, 140.	2.1	18
66	Locus and allelic heterogeneity and phenotypic variability in Waardenburg syndrome. Clinical Genetics, 2019, 95, 398-402.	1.0	18
67	Synpolydactyly and HOXD13 polyalanine repeat: addition of 2 alanine residues is without clinical consequences. BMC Medical Genetics, 2007, 8, 78.	2.1	17
68	Additional three patients with Smith-McCort dysplasia due to novel <i>RAB33B</i> mutations. American Journal of Medical Genetics, Part A, 2017, 173, 588-595.	0.7	17
69	Heterozygous ANKRD17 loss-of-function variants cause a syndrome with intellectual disability, speech delay, and dysmorphism. American Journal of Human Genetics, 2021, 108, 1138-1150.	2.6	17
70	A novel multiple joint dislocation syndrome associated with a homozygous nonsense variant in the EXOC6B gene. European Journal of Human Genetics, 2016, 24, 1206-1210.	1.4	16
71	Hypomorphic Mutations in TONSL Cause SPONASTRIME Dysplasia. American Journal of Human Genetics, 2019, 104, 439-453.	2.6	16
72	Biallelic missense variant, p.Ser35Leu in <i>EXOSC1</i> is associated with pontocerebellar hypoplasia. Clinical Genetics, 2021, 99, 594-600.	1.0	16

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73	A novel mutation (g.106737G>T) in zone of polarizing activity regulatory sequence (ZRS) causes variable limb phenotypes in Werner mesomelia. American Journal of Medical Genetics, Part A, 2014, 164, 898-906.	0.7	15
74	Spectrum of <i>SMPD1</i> mutations in Asian-Indian patients with acid sphingomyelinase (ASM)-deficient Niemann-Pick disease. American Journal of Medical Genetics, Part A, 2016, 170, 2719-2730.	0.7	15
75	Report of four novel variants in <i>ASNS</i> causing asparagine synthetase deficiency and review of literature. Congenital Anomalies (discontinued), 2018, 58, 181-182.	0.3	15
76	Genetic diversity of NDUFV1-dependent mitochondrial complex I deficiency. European Journal of Human Genetics, 2018, 26, 1582-1587.	1.4	15
77	Meckel syndrome: Clinical and mutation profile in six fetuses. Clinical Genetics, 2019, 96, 560-565.	1.0	15
78	Turner syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2020, 182, 303-313.	0.7	15
79	Torg-Winchester syndrome: lack of efficacy of pamidronate therapy. Clinical Dysmorphology, 2007, 16, 95-100.	0.1	14
80	A biallelic 36-bp insertion in PIBF1 is associated with Joubert syndrome. Journal of Human Genetics, 2018, 63, 935-939.	1.1	14
81	Identification of a novel homozygous variant confirms <i>ITPA</i> as a developmental and epileptic encephalopathy gene. American Journal of Medical Genetics, Part A, 2019, 179, 857-861.	0.7	14
82	Bain type of X-linked syndromic mental retardation in a male with a pathogenic variant in HNRNPH2. American Journal of Medical Genetics, Part A, 2020, 182, 183-188.	0.7	14
83	Bi-allelic variants in IPO8 cause a connective tissue disorder associated with cardiovascular defects, skeletal abnormalities, and immune dysregulation. American Journal of Human Genetics, 2021, 108, 1126-1137.	2.6	14
84	SOX11 variants cause a neurodevelopmental disorder with infrequent ocular malformations and hypogonadotropic hypogonadism and with distinct DNA methylation profile. Genetics in Medicine, 2022, 24, 1261-1273.	1.1	14
85	Report of second case and clinical and molecular characterization of Eiken syndrome. Clinical Genetics, 2018, 94, 457-460.	1.0	13
86	Development, behaviour and autism in individuals with <i>SMC1A</i> variants. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2019, 60, 305-313.	3.1	13
87	Genome sequencing in families with congenital limb malformations. Human Genetics, 2021, 140, 1229-1239.	1.8	13
88	Fetal akinesia deformation sequence: Expanding the phenotypic spectrum. American Journal of Medical Genetics, Part A, 2014, 164, 2643-2648.	0.7	12
89	A Novel Frameshift Mutation in TWIST2 Gene Causing Setleis Syndrome. Indian Journal of Pediatrics, 2014, 81, 302-304.	0.3	12
90	Clinical and genetic spectrum of 104 Indian families with central nervous system white matter abnormalities. Clinical Genetics, 2021, 100, 542-550.	1.0	12

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91	Exome Sequencing Identifies a Dominant <i>TNNT3</i> Mutation in a Large Family with Distal Arthrogyriposis. <i>Molecular Syndromology</i> , 2014, 5, 218-228.	0.3	11
92	White matter changes in GM1 gangliosidosis. <i>Indian Pediatrics</i> , 2015, 52, 155-156.	0.2	11
93	Spectrum of urorectal septum malformation sequence. <i>Congenital Anomalies (discontinued)</i> , 2016, 56, 119-126.	0.3	11
94	Variants in the transcriptional corepressor <i>BCORL1</i> are associated with an X-linked disorder of intellectual disability, dysmorphic features, and behavioral abnormalities. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 870-874.	0.7	11
95	Mucopolipidosis type II with a homozygous missense mutation in the <i>GNPTAB</i> gene. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 1225-1228.	0.7	10
96	Hyperphosphatasia with Mental Retardation Syndrome Due to a Novel Mutation in PGAP3. <i>Journal of Pediatric Genetics</i> , 2017, 06, 191-193.	0.3	10
97	Seven additional families with spondylocarpotarsal synostosis syndrome with novel biallelic deleterious variants in <i>FLNB</i> . <i>Clinical Genetics</i> , 2018, 94, 159-164.	1.0	10
98	Confirmation of a Rare Genetic Leukoencephalopathy due to a Novel Bi-allelic Variant in RPIA. <i>European Journal of Medical Genetics</i> , 2019, 62, 103708.	0.7	10
99	Phenotyping and genotyping of skeletal dysplasias: Evolution of a center and a decade of experience in India. <i>Bone</i> , 2019, 120, 204-211.	1.4	10
100	Preimplantation diagnosis of genetic diseases. <i>Journal of Postgraduate Medicine</i> , 2010, 56, 317-320.	0.2	10
101	Novel biallelic variants expand the SLC5A6-related phenotypic spectrum. <i>European Journal of Human Genetics</i> , 2022, 30, 439-449.	1.4	10
102	A novel <i>EDARADD</i> splice site mutation resulting in activation of two alternate cryptic splice sites causes autosomal recessive Hypohidrotic Ectodermal Dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1639-1641.	0.7	9
103	Novel variant p.(Ala102Thr) in <i>SDHB</i> causes mitochondrial complex II deficiency: Case report and review of the literature. <i>Annals of Human Genetics</i> , 2020, 84, 345-351.	0.3	9
104	Rhizomelic chondrodysplasia punctata type 1: report of mutations in 3 children from India. <i>Journal of Applied Genetics</i> , 2010, 51, 107-110.	1.0	8
105	Placental Teratoma Presenting as a Lobulated Mass behind the Neck of Fetus: A Case Report. <i>Case Reports in Obstetrics and Gynecology</i> , 2012, 2012, 1-2.	0.2	8
106	Profile of patients with Von Gierke disease from India. <i>Indian Pediatrics</i> , 2012, 49, 228-230.	0.2	8
107	The novel <i>EDAR</i> p.L397H missense mutation causes autosomal dominant hypohidrotic ectodermal dysplasia. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2017, 31, e17-e20.	1.3	8
108	Homozygous c.359del variant in <i>MGME1</i> is associated with early onset cerebellar ataxia. <i>European Journal of Medical Genetics</i> , 2017, 60, 533-535.	0.7	8



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109	Variable presentation of Fraser syndrome in two fetuses and a novel mutation in <i>FRAS1</i> . <i>Congenital Anomalies (discontinued)</i> , 2017, 57, 83-85.	0.3	8
110	Report of the Third Family with Multiple Mitochondrial Dysfunctions Syndrome 5 Caused by the Founder Variant p.(Glu87Lys) in ISCA1. <i>Journal of Pediatric Genetics</i> , 2018, 07, 130-133.	0.3	8
111	Homozygous variant, p.(Arg643Trp) in VAC14 causes striatonigral degeneration: report of a novel variant and review of VAC14-related disorders. <i>Journal of Human Genetics</i> , 2019, 64, 1237-1242.	1.1	8
112	Phenotypic diversity and genetic complexity of PAX3 related Waardenburg syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2951-2958.	0.7	8
113	Recurrent bi-allelic splicing variant c.454+3A>G in TRAPPC4 is associated with progressive encephalopathy and muscle involvement. <i>Brain</i> , 2020, 143, e29.	3.7	8
114	<i>LACC1</i> gene mutation in three sisters with polyarthritis without systemic features. <i>Annals of the Rheumatic Diseases</i> , 2020, 79, 425-426.	0.5	8
115	PPA2-associated sudden cardiac death: extending the clinical and allelic spectrum in 20 new families. <i>Genetics in Medicine</i> , 2021, 23, 2415-2425.	1.1	8
116	Diagnosis and Management of Global Development Delay: Consensus Guidelines of Growth, Development and Behavioral Pediatrics Chapter, Neurology Chapter and Neurodevelopment Pediatrics Chapter of the Indian Academy of Pediatrics. <i>Indian Pediatrics</i> , 2022, 59, 401-415.	0.2	8
117	Immunological response to two hepatitis B vaccines administered in two different schedules. <i>Indian Journal of Pediatrics</i> , 2006, 73, 489-491.	0.3	7
118	Novel mutation and atlantoaxial dislocation in two siblings from India with Dyggve-Melchior-Clausen syndrome. <i>European Journal of Medical Genetics</i> , 2008, 51, 251-256.	0.7	7
119	Long-term efficacy of oral deferiprone in management of iron overload in beta thalassemia major. <i>Hematology</i> , 2008, 13, 77-82.	0.7	7
120	Massive cranial osteolysis, skin changes, growth retardation and developmental delay: Gorham syndrome with systemic manifestations?. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 759-763.	0.7	7
121	Second report of slipped capital femoral epiphysis in Rubinstein-Taybi syndrome. <i>Clinical Dysmorphology</i> , 2011, 20, 55-57.	0.1	7
122	Occurrence of Synpolydactyly and Omphalocele in a Fetus with a HOXD13 Mutation. <i>Journal of Pediatric Genetics</i> , 2017, 06, 194-197.	0.3	7
123	Bi-allelic c.181_183delTCT in BTB domain of KLHL7 is associated with overlapping phenotypes of Crisponi/CISS1-like and Bohring-Opitz like syndrome. <i>European Journal of Medical Genetics</i> , 2019, 62, 103528.	0.7	7
124	The third family with Eiken syndrome. <i>Clinical Genetics</i> , 2019, 96, 378-379.	1.0	7
125	A novel bi-allelic loss-of-function variant in <i>MYOD1</i> : Further evidence for gene-disease association and phenotypic variability in <i>MYOD1</i> -related myopathy. <i>Clinical Genetics</i> , 2019, 96, 276-277.	1.0	7
126	Locus and allelic heterogeneity in five families with hereditary spastic paraplegia. <i>Journal of Human Genetics</i> , 2019, 64, 17-21.	1.1	7



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127	Clinical Characteristics, Molecular Profile, and Outcomes in Indian Patients with Glutaric Aciduria Type 1. <i>Journal of Pediatric Genetics</i> , 2021, 10, 213-221.	0.3	7
128	Genetic disorders with central nervous system white matter abnormalities: An update. <i>Clinical Genetics</i> , 2021, 99, 119-132.	1.0	7
129	Deficiency of TMEM53 causes a previously unknown sclerosing bone disorder by dysregulation of BMP-SMAD signaling. <i>Nature Communications</i> , 2021, 12, 2046.	5.8	7
130	Multilocus disease-causing genomic variations for Mendelian disorders: role of systematic phenotyping and implications on genetic counselling. <i>European Journal of Human Genetics</i> , 2021, 29, 1774-1780.	1.4	7
131	Clinically relevant variants in a large cohort of Indian patients with Marfan syndrome and related disorders identified by next-generation sequencing. <i>Scientific Reports</i> , 2021, 11, 764.	1.6	7
132	Familial 7q11.23 duplication with variable phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2727-2730.	0.7	6
133	Clinical Variability in Familial X-Linked Ohdo Syndrome—Maat-Kievit-Brunner Type with MED12 Mutation. <i>Journal of Pediatric Genetics</i> , 2017, 06, 198-204.	0.3	6
134	Pycnodysostosis: Novel Variants in CTSK and Occurrence of Giant Cell Tumor. <i>Journal of Pediatric Genetics</i> , 2018, 07, 009-013.	0.3	6
135	Biallelic variants p.Arg1133Cys and p.Arg1379Cys in <i>COL2A1</i> : Further delineation of phenotypic spectrum of recessive Type 2 collagenopathies. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 338-347.	0.7	6
136	Late onset Pompe Disease in India — Beyond the Caucasian phenotype. <i>Neuromuscular Disorders</i> , 2021, 31, 431-441.	0.3	6
137	Functional characterization of novel variants in <i>SMPD1</i> in Indian patients with acid sphingomyelinase deficiency. <i>Human Mutation</i> , 2021, 42, 1336-1350.	1.1	6
138	Milder form of pachydermoperiostosis: a report of four cases. <i>Clinical Dysmorphology</i> , 2009, 18, 85-89.	0.1	5
139	A syndrome of facial dysmorphism, cubital pterygium, short distal phalanges, swan neck deformity of fingers, and scoliosis. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1035-1040.	0.7	5
140	Is coloboma a feature of fetal valproate syndrome?. <i>Clinical Dysmorphology</i> , 2014, 23, 24-25.	0.1	5
141	Hunter syndrome with late age of presentation: clinical description of a case and review of the literature. <i>BMJ Case Reports</i> , 2015, 2015, bcr2015209305-bcr2015209305.	0.2	5
142	Homozygous deletion of exons 2 and 3 of NPC2 associated with Niemann—Pick disease type C. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2486-2489.	0.7	5
143	Intrafamilial variability in syndromic microphthalmia type 5 caused by a novel variation in <i>OTX2</i> . <i>Ophthalmic Genetics</i> , 2017, 38, 533-536.	0.5	5
144	Severe Form of Brachydactyly Type A1 in a Child with a c.298G>>A Mutation in IHH Gene. <i>Journal of Pediatric Genetics</i> , 2017, 06, 177-180.	0.3	5

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145	p.Arg69Trp in <i>RNASEH2C</i> is a founder variant in three Indian families with Aicardi's "Goutières" syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 156-160.	0.7	5
146	Novel pathogenic variants in <i>GBE1</i> causing fetal akinesia deformation sequence and severe neuromuscular form of glycogen storage disease type IV. <i>Clinical Dysmorphology</i> , 2019, 28, 17-21.	0.1	5
147	Identification of novel variants in a large cohort of children with Tay-Sachs disease: An initiative of a multicentric task force on lysosomal storage disorders by Government of India. <i>Journal of Human Genetics</i> , 2019, 64, 985-994.	1.1	5
148	<i>GATAD2B</i> -related intellectual disability due to parental mosaicism and review of literature. <i>Clinical Dysmorphology</i> , 2019, 28, 190-194.	0.1	5
149	Recurrent 1q21.1 deletion syndrome: report on variable expression, nonpenetrance and review of literature. <i>Clinical Dysmorphology</i> , 2020, 29, 127-131.	0.1	5
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