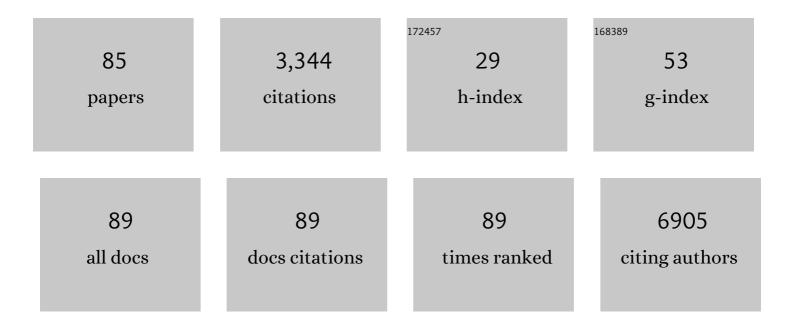
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Accelerating Novel Candidate Gene Discovery in Neurogenetic Disorders via Whole-Exome Sequencing of Prescreened Multiplex Consanguineous Families. Cell Reports, 2015, 10, 148-161.	6.4	375
2	Clinical exome sequencing: results from 2819 samples reflecting 1000 families. European Journal of Human Genetics, 2017, 25, 176-182.	2.8	291
3	The landscape of genetic diseases in Saudi Arabia based on the first 1000 diagnostic panels and exomes. Human Genetics, 2017, 136, 921-939.	3.8	209
4	Lessons Learned from Large-Scale, First-Tier Clinical Exome Sequencing in a Highly Consanguineous Population. American Journal of Human Genetics, 2019, 104, 1182-1201.	6.2	184
5	IFT27, encoding a small GTPase component of IFT particles, is mutated in a consanguineous family with Bardet-Biedl syndrome. Human Molecular Genetics, 2014, 23, 3307-3315.	2.9	134
6	Expanding the genetic heterogeneity of intellectual disability. Human Genetics, 2017, 136, 1419-1429.	3.8	122
7	Neu-Laxova Syndrome, an Inborn Error of Serine Metabolism, Is Caused by Mutations in PHGDH. American Journal of Human Genetics, 2014, 94, 898-904.	6.2	93
8	Expanding the clinical and genetic heterogeneity of hereditary disorders of connective tissue. Human Genetics, 2016, 135, 525-540.	3.8	89
9	Genomic and phenotypic delineation of congenital microcephaly. Genetics in Medicine, 2019, 21, 545-552.	2.4	85
10	Molecular autopsy in maternal–fetal medicine. Genetics in Medicine, 2018, 20, 420-427.	2.4	84
11	Autozygome and high throughput confirmation of disease genes candidacy. Genetics in Medicine, 2019, 21, 736-742.	2.4	81
12	Ciliary Genes <i>TBC1D32</i> / <i>C6orf170</i> and <i>SCLT1</i> are Mutated in Patients with OFD Type IX. Human Mutation, 2014, 35, 36-40.	2.5	78
13	Identification of novel loci for pediatric cholestatic liver disease defined by KIF12, PPM1F, USP53, LSR, and WDR83OS pathogenic variants. Genetics in Medicine, 2019, 21, 1164-1172.	2.4	71
14	ARL3 Mutations Cause Joubert Syndrome by Disrupting Ciliary Protein Composition. American Journal of Human Genetics, 2018, 103, 612-620.	6.2	70
15	A multicenter clinical exome study in unselected cohorts from a consanguineous population of Saudi Arabia demonstrated a high diagnostic yield. Molecular Genetics and Metabolism, 2017, 121, 91-95.	1.1	68
16	The morbid genome of ciliopathies: an update. Genetics in Medicine, 2020, 22, 1051-1060.	2.4	68
17	Successful application of genome sequencing in a diagnostic setting: 1007 index cases from a clinically heterogeneous cohort. European Journal of Human Genetics, 2021, 29, 141-153.	2.8	66
18	Homozygous KCNMA1 mutation as a cause of cerebellar atrophy, developmental delay and seizures. Human Genetics, 2016, 135, 1295-1298.	3.8	65

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19	Analysis of transcript-deleterious variants in Mendelian disorders: implications for RNA-based diagnostics. Genome Biology, 2020, 21, 145.	8.8	59
20	KIAA0556 is a novel ciliary basal body component mutated in Joubert syndrome. Genome Biology, 2015, 16, 293.	8.8	56
21	Accelerating matchmaking of novel dysmorphology syndromes through clinical and genomic characterization of a large cohort. Genetics in Medicine, 2016, 18, 686-695.	2.4	55
22	Severe CNS involvement in <i>WWOX</i> mutations: Description of five new cases. American Journal of Medical Genetics, Part A, 2015, 167, 3209-3213.	1.2	50
23	Loss of SMPD4 Causes a Developmental Disorder Characterized by Microcephaly and Congenital Arthrogryposis. American Journal of Human Genetics, 2019, 105, 689-705.	6.2	48
24	Expanding the phenome and variome of skeletal dysplasia. Genetics in Medicine, 2018, 20, 1609-1616.	2.4	46
25	Mutations in DDX59 Implicate RNA Helicase in the Pathogenesis of Orofaciodigital Syndrome. American Journal of Human Genetics, 2013, 93, 555-560.	6.2	45
26	Identification of a novel MKS locus defined by <i>TMEM107</i> mutation. Human Molecular Genetics, 2015, 24, 5211-5218.	2.9	42
27	Genetic variants in components of the NALCN–UNC80–UNC79 ion channel complex cause a broad clinical phenotype (NALCN channelopathies). Human Genetics, 2018, 137, 753-768.	3.8	38
28	Vici syndrome associated with unilateral lung hypoplasia and myopathy. American Journal of Medical Genetics, Part A, 2010, 152A, 1849-1853.	1.2	33
29	Severe early-onset epileptic encephalopathy due to mutations in the KCNA2 gene: Expansion of the genotypic and phenotypic spectrum. European Journal of Paediatric Neurology, 2016, 20, 657-660.	1.6	33
30	MDH1 deficiency is a metabolic disorder of the malate–aspartate shuttle associated with early onset severe encephalopathy. Human Genetics, 2019, 138, 1247-1257.	3.8	31
31	Congenital disorders of glycosylation: The Saudi experience. American Journal of Medical Genetics, Part A, 2017, 173, 2614-2621.	1.2	31
32	Combining exome/genome sequencing with data repository analysis reveals novel gene–disease associations for a wide range of genetic disorders. Genetics in Medicine, 2021, 23, 1551-1568.	2.4	30
33	Effect of consanguinity on birth defects in Saudi women: Results from a nested caseâ€control study. Birth Defects Research Part A: Clinical and Molecular Teratology, 2015, 103, 100-104.	1.6	28
34	FARS2 deficiency; new cases, review of clinical, biochemical, and molecular spectra, and variants interpretation based on structural, functional, and evolutionary significance. Molecular Genetics and Metabolism, 2018, 125, 281-291.	1.1	28
35	Biallelic Mutations in Tetratricopeptide Repeat Domain 26 (Intraflagellar Transport 56) Cause Severe Biliary Ciliopathy in Humans. Hepatology, 2020, 71, 2067-2079.	7.3	28
36	Biallelic variants in the transcription factor PAX7 are a new genetic cause of myopathy. Genetics in Medicine, 2019, 21, 2521-2531.	2.4	25

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37	Recessive AFG3L2 Mutation Causes Progressive Microcephaly, Early Onset Seizures, Spasticity, and Basal Ganglia Involvement. Pediatric Neurology, 2017, 71, 24-28.	2.1	19
38	Categorized Genetic Analysis in Childhood-Onset Cardiomyopathy. Circulation Genomic and Precision Medicine, 2020, 13, 504-514.	3.6	18
39	A Wide Clinical Phenotype Spectrum in Patients With <i>ATP1A2</i> Mutations. Journal of Child Neurology, 2014, 29, 265-268.	1.4	17
40	Further Delineation of the ALG9-CDG Phenotype. JIMD Reports, 2015, 27, 107-112.	1.5	17
41	Congenital anomalies and associated risk factors in a Saudi population: a cohort study from pregnancy to age 2 years. BMJ Open, 2019, 9, e026351.	1.9	17
42	An exome-first approach to aid in the diagnosis of primary ciliary dyskinesia. Human Genetics, 2020, 139, 1273-1283.	3.8	16
43	The Leukodystrophy Spectrum in Saudi Arabia: Epidemiological, Clinical, Radiological, and Genetic Data. Frontiers in Pediatrics, 2021, 9, 633385.	1.9	15
44	Truncating ARL6IP1 variant as the genetic cause of fatal complicated hereditary spastic paraplegia. BMC Medical Genetics, 2019, 20, 119.	2.1	14
45	Clinical profile and mutation spectrum of long QT syndrome in Saudi Arabia: The impact of consanguinity. Heart Rhythm, 2017, 14, 1191-1199.	0.7	13
46	Exploiting the Autozygome to Support Previously Published Mendelian Gene-Disease Associations: An Update. Frontiers in Genetics, 2020, 11, 580484.	2.3	13
47	Lethal variants in humans: lessons learned from a large molecular autopsy cohort. Genome Medicine, 2021, 13, 161.	8.2	13
48	Incidence of newborn screening disorders among 56632 infants in Central Saudi Arabia. Journal of King Abdulaziz University, Islamic Economics, 2020, 41, 703-708.	1.1	13
49	6-Pyruvoyltetrahydropterin Synthase Deficiency: Review and Report of 28 Arab Subjects. Pediatric Neurology, 2019, 96, 40-47.	2.1	12
50	The many faces of peroxisomal disorders: Lessons from a large Arab cohort. Clinical Genetics, 2019, 95, 310-319.	2.0	12
51	Biallelic <scp> <i>ZNFX1 </i> </scp> variants are associated with a spectrum of immunoâ€hematological abnormalities. Clinical Genetics, 2022, 101, 247-254.	2.0	12
52	ADAMTS19 â€associated heart valve defects: Novel genetic variants consolidating a recognizable cardiac phenotype. Clinical Genetics, 2020, 98, 56-63.	2.0	11
53	Biallelic variants in KARS1 are associated with neurodevelopmental disorders and hearing loss recapitulated by the knockout zebrafish. Genetics in Medicine, 2021, 23, 1933-1943.	2.4	11
54	Patterns, prevalence, risk factors, and survival of newborns with congenital heart defects in a Saudi population: a three-year, cohort case-control study. Journal of Congenital Cardiology, 2019, 3, .	0.5	10

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55	Munchausen syndrome by proxy mimicking as Gaucher disease. European Journal of Pediatrics, 2010, 169, 1029-1032.	2.7	9
56	Smith–Lemli–Opitz syndrome among Arabs. Clinical Genetics, 2012, 82, 165-172.	2.0	9
57	Further delineation of Temtamy syndrome of corpus callosum and ocular abnormalities. American Journal of Medical Genetics, Part A, 2018, 176, 715-721.	1.2	7
58	Genetic, clinical and biochemical characterization of a large cohort of patients with hyaline fibromatosis syndrome. Orphanet Journal of Rare Diseases, 2019, 14, 209.	2.7	7
59	Novel Homozygous Mutation of the AIMP1 Gene: A Milder Neuroimaging Phenotype With Preservation of the Deep White Matter. Pediatric Neurology, 2019, 91, 57-61.	2.1	7
60	Survey of disorders of sex development in a large cohort of patients with diverse Mendelian phenotypes. American Journal of Medical Genetics, Part A, 2021, 185, 2789-2800.	1.2	7
61	Distal acroosteolysis, poikiloderma and joint stiffness: a novel laminopathy?. European Journal of Human Genetics, 2016, 24, 1220-1222.	2.8	6
62	Recurrent homozygous damaging mutation in <i>TMX2</i> , encoding a protein disulfide isomerase, in four families with microlissencephaly. Journal of Medical Genetics, 2020, 57, 274-282.	3.2	6
63	Neuroimaging manifestations and genetic heterogeneity of Walker-Warburg syndrome in Saudi patients. Brain and Development, 2021, 43, 380-388.	1.1	6
64	Molecular autopsy by proxy in preconception counseling. Clinical Genetics, 2021, 100, 678-691.	2.0	6
65	Bi-allelic loss-of-function variants in PPFIBP1 cause a neurodevelopmental disorder with microcephaly, epilepsy, and periventricular calcifications. American Journal of Human Genetics, 2022, 109, 1421-1435.	6.2	6
66	Crisponi/CISS1 syndrome: A case series. American Journal of Medical Genetics, Part A, 2016, 170, 1236-1241.	1.2	5
67	A homozygous frameshift variant in an alternatively spliced exon of <i>DLG5</i> causes hydrocephalus and renal dysplasia. Clinical Genetics, 2019, 95, 631-633.	2.0	5
68	Fructose-1,6-bisphosphatase deficiency with confirmed molecular diagnosis. Journal of King Abdulaziz University, Islamic Economics, 2020, 41, 199-202.	1.1	5
69	A de novo splicing variant supports the candidacy of TLL1 in ASD pathogenesis. European Journal of Human Genetics, 2020, 28, 525-528.	2.8	4
70	Progressive symmetrical erythrokeratoderma manifesting as harlequin-like ichthyosis with severe thrombocytopenia secondary to a homozygous 3-ketodihydrosphingosine reductase mutation. JAAD Case Reports, 2021, 14, 55-58.	0.8	4
71	Further delineation of <scp><i>SMG9</i></scp> â€related heart and brain malformation syndrome. American Journal of Medical Genetics, Part A, 2021, 185, 1624-1630.	1.2	3
72	Mutations in phospholipase C eta-1 (<i>PLCH1</i>) are associated with holoprosencephaly. Journal of Medical Genetics, 2022, 59, 358-365.	3.2	3

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73	Further clinical and genetic evidence of ASC-1 complex dysfunction in congenital neuromuscular disease. European Journal of Medical Genetics, 2022, 65, 104537.	1.3	3
74	A Biallelic Variant in <i>FRA10AC1</i> Is Associated With Neurodevelopmental Disorder and Growth Retardation. Neurology: Genetics, 2022, 8, e200010.	1.9	2
75	Embryopathy Associated With a Vitamin Therapy. Pediatric Neurology, 2018, 89, 73-74.	2.1	1
76	Hypospadias in ring X syndrome. European Journal of Medical Genetics, 2021, 64, 104225.	1.3	1
77	The phenotypic spectrum of dihydrolipoamide dehydrogenase deficiency in Saudi Arabia. Molecular Genetics and Metabolism Reports, 2021, 29, 100817.	1.1	1
78	An atypical presentation of severe congenital contractures and lack of cerebellar involvement in a patient with a novel LAMA1 mutation. Journal of Biochemical and Clinical Genetics, 0, , 43-46.	0.1	1
79	Molecular and clinical characteristics of very long-chain acyl-CoA dehydrogenase deficiency. Journal of King Abdulaziz University, Islamic Economics, 2020, 41, 590-596.	1.1	1
80	De Novo Ring Chromosome 15: Molecular Cytogenetic and Clinical Characterization of First Case from Saudi Arabia. Journal of Pediatric Genetics, 0, , .	0.7	1
81	Peripheral venous route for administration of ammonul infusion for treatment of acute hyperammonemia. Journal of King Abdulaziz University, Islamic Economics, 2020, 41, 98-101.	1.1	0
82	The genotypic and phenotypic spectrum of pycnodysostosis in Saudi Arabia: Novel variants and clinical findings. American Journal of Medical Genetics, Part A, 2021, 185, 2455-2463.	1.2	0
83	Neuroregression, coarse features, and oligosaccharides in urines. Neurosciences, 2017, 22, 326-328.	0.1	0
84	Epilepsy, neuropsychiatric phenotypes, neuroimaging findings, and genotype-neurophenotype correlation in 22q11.2 deletion syndrome. Journal of King Abdulaziz University, Islamic Economics, 2020, 25, 287-291.	1.1	0
85	Mitochondrial "dysmorphology―in variant classification. Human Genetics, 2022, 141, 55-64.	3.8	0