

Dagmar Wieczorek

List of Publications by Citations

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225
papers

12,049
citations

59
h-index

102
g-index

239
ext. papers

15,031
ext. citations

7.9
avg, IF

5.5
L-index

#	Paper	IF	Citations
225	Range of genetic mutations associated with severe non-syndromic sporadic intellectual disability: an exome sequencing study. <i>Lancet, The</i> , 2012 , 380, 1674-82	40	765
224	Deep sequencing reveals 50 novel genes for recessive cognitive disorders. <i>Nature</i> , 2011 , 478, 57-63	50.4	649
223	Severe COVID-19 Is Marked by a Dysregulated Myeloid Cell Compartment. <i>Cell</i> , 2020 , 182, 1419-1440.e236.2	36.2	558
222	Germline KRAS and BRAF mutations in cardio-facio-cutaneous syndrome. <i>Nature Genetics</i> , 2006 , 38, 294-6	36.3	437
221	Mutations in GRIN2A and GRIN2B encoding regulatory subunits of NMDA receptors cause variable neurodevelopmental phenotypes. <i>Nature Genetics</i> , 2010 , 42, 1021-6	36.3	347
220	Mutations in genes encoding subunits of RNA polymerases I and III cause Treacher Collins syndrome. <i>Nature Genetics</i> , 2011 , 43, 20-2	36.3	239
219	SARS-CoV-2 targets neurons of 3D human brain organoids. <i>EMBO Journal</i> , 2020 , 39, e106230	13	206
218	De novo nonsense mutations in ASXL1 cause Bohring-Opitz syndrome. <i>Nature Genetics</i> , 2011 , 43, 729-31	36.3	198
217	CEP152 is a genome maintenance protein disrupted in Seckel syndrome. <i>Nature Genetics</i> , 2011 , 43, 23-6	36.3	185
216	Haploinsufficiency of ARID1B, a member of the SWI/SNF-a chromatin-remodeling complex, is a frequent cause of intellectual disability. <i>American Journal of Human Genetics</i> , 2012 , 90, 565-72	11	182
215	Mutations in NSUN2 cause autosomal-recessive intellectual disability. <i>American Journal of Human Genetics</i> , 2012 , 90, 847-55	11	179
214	The core FOXP1 syndrome phenotype consists of postnatal microcephaly, severe mental retardation, absent language, dyskinesia, and corpus callosum hypogenesis. <i>Journal of Medical Genetics</i> , 2011 , 48, 396-406	5.8	179
213	Mutations in U4atac snRNA, a component of the minor spliceosome, in the developmental disorder MOPD I. <i>Science</i> , 2011 , 332, 238-40	33.3	179
212	Heterozygous missense mutations in SMARCA2 cause Nicolaides-Baraitser syndrome. <i>Nature Genetics</i> , 2012 , 44, 445-9, S1	36.3	170
211	Oculo-auriculo-vertebral spectrum (OAVS): clinical evaluation and severity scoring of 53 patients and proposal for a new classification. <i>European Journal of Medical Genetics</i> , 2005 , 48, 397-411	2.6	150
210	Expanding the phenotypic spectrum of lupus erythematosus in Aicardi-Goutières syndrome. <i>Arthritis and Rheumatism</i> , 2010 , 62, 1469-77		145
209	ZMYND10 is mutated in primary ciliary dyskinesia and interacts with LRRC6. <i>American Journal of Human Genetics</i> , 2013 , 93, 336-45	11	144

208	Haploinsufficiency of a spliceosomal GTPase encoded by EFTUD2 causes mandibulofacial dysostosis with microcephaly. <i>American Journal of Human Genetics</i> , 2012 , 90, 369-77	11	143
207	A comprehensive molecular study on Coffin-Siris and Nicolaides-Baraitser syndromes identifies a broad molecular and clinical spectrum converging on altered chromatin remodeling. <i>Human Molecular Genetics</i> , 2013 , 22, 5121-35	5.6	138
206	Mutations in SRCAP, encoding SNF2-related CREBBP activator protein, cause Floating-Harbor syndrome. <i>American Journal of Human Genetics</i> , 2012 , 90, 308-13	11	130
205	Spectrum of mutations in PTPN11 and genotype-phenotype correlation in 96 patients with Noonan syndrome and five patients with cardio-facio-cutaneous syndrome. <i>European Journal of Human Genetics</i> , 2003 , 11, 201-6	5.3	128
204	Genotyping in 46 patients with tentative diagnosis of Treacher Collins syndrome revealed unexpected phenotypic variation. <i>European Journal of Human Genetics</i> , 2004 , 12, 879-90	5.3	124
203	Diagnostic approach to microcephaly in childhood: a two-center study and review of the literature. <i>Developmental Medicine and Child Neurology</i> , 2014 , 56, 732-41	3.3	123
202	Diagnostic Yield and Novel Candidate Genes by Exome Sequencing in 152 Consanguineous Families With Neurodevelopmental Disorders. <i>JAMA Psychiatry</i> , 2017 , 74, 293-299	14.5	116
201	Mutations in the histone methyltransferase gene KMT2B cause complex early-onset dystonia. <i>Nature Genetics</i> , 2017 , 49, 223-237	36.3	116
200	Longitudinal Multi-omics Analyses Identify Responses of Megakaryocytes, Erythroid Cells, and Plasmablasts as Hallmarks of Severe COVID-19. <i>Immunity</i> , 2020 , 53, 1296-1314.e9	32.3	109
199	A specific mutation in the distant sonic hedgehog (SHH) cis-regulator (ZRS) causes Werner mesomelic syndrome (WMS) while complete ZRS duplications underlie Haas type polysyndactyly and preaxial polydactyly (PPD) with or without triphalangeal thumb. <i>Human Mutation</i> , 2010 , 31, 81-9	4.7	104
198	Acetylcholine receptor pathway mutations explain various fetal akinesia deformation sequence disorders. <i>American Journal of Human Genetics</i> , 2008 , 82, 464-76	11	104
197	De novo CCND2 mutations leading to stabilization of cyclin D2 cause megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome. <i>Nature Genetics</i> , 2014 , 46, 510-515	36.3	100
196	The genetic basis of DOORS syndrome: an exome-sequencing study. <i>Lancet Neurology</i> , 2014 , 13, 44-58	24.1	96
195	Swarm Learning for decentralized and confidential clinical machine learning. <i>Nature</i> , 2021 , 594, 265-270	50.4	89
194	Mutation Update for Kabuki Syndrome Genes KMT2D and KDM6A and Further Delineation of X-Linked Kabuki Syndrome Subtype 2. <i>Human Mutation</i> , 2016 , 37, 847-64	4.7	89
193	RPA and Rad51 constitute a cell intrinsic mechanism to protect the cytosol from self DNA. <i>Nature Communications</i> , 2016 , 7, 11752	17.4	88
192	A paternal deletion of MKRN3, MAGEL2 and NDN does not result in Prader-Willi syndrome. <i>European Journal of Human Genetics</i> , 2009 , 17, 582-90	5.3	88
191	A mutation screen in patients with Kabuki syndrome. <i>Human Genetics</i> , 2011 , 130, 715-24	6.3	87

190	SOS1 is the second most common Noonan gene but plays no major role in cardio-facio-cutaneous syndrome. <i>Journal of Medical Genetics</i> , 2007 , 44, 651-6	5.8	85
189	Neu-Laxova syndrome is a heterogeneous metabolic disorder caused by defects in enzymes of the L-serine biosynthesis pathway. <i>American Journal of Human Genetics</i> , 2014 , 95, 285-93	11	82
188	A novel microdeletion syndrome involving 5q14.3-q15: clinical and molecular cytogenetic characterization of three patients. <i>European Journal of Human Genetics</i> , 2009 , 17, 1592-9	5.3	81
187	Mutation and phenotypic spectrum in patients with cardio-facio-cutaneous and Costello syndrome. <i>Clinical Genetics</i> , 2008 , 73, 62-70	4	81
186	Molecular and clinical characterization of cardio-facio-cutaneous (CFC) syndrome: overlapping clinical manifestations with Costello syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2007 , 143A, 799-807	2.5	79
185	Genetics of intellectual disability in consanguineous families. <i>Molecular Psychiatry</i> , 2019 , 24, 1027-1039	15.1	79
184	Effects of RANK-ligand antibody (denosumab) treatment on bone turnover markers in a girl with juvenile Paget's disease. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013 , 98, 3121-6	5.6	78
183	Molecular and clinical analysis of RAF1 in Noonan syndrome and related disorders: dephosphorylation of serine 259 as the essential mechanism for mutant activation. <i>Human Mutation</i> , 2010 , 31, 284-94	4.7	77
182	Genotype-phenotype correlation of Coffin-Siris syndrome caused by mutations in SMARCB1, SMARCA4, SMARCE1, and ARID1A. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014 , 166C, 262-75	3.1	74
181	Deficiency of ECHS1 causes mitochondrial encephalopathy with cardiac involvement. <i>Annals of Clinical and Translational Neurology</i> , 2015 , 2, 492-509	5.3	69
180	A noncoding, regulatory mutation implicates HCFC1 in nonsyndromic intellectual disability. <i>American Journal of Human Genetics</i> , 2012 , 91, 694-702	11	69
179	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. <i>Genetics in Medicine</i> , 2018 , 20, 630-638	8.1	68
178	A family with autosomal dominant oculo-auriculo-vertebral spectrum. <i>Clinical Dysmorphology</i> , 2007 , 16, 1-7	0.9	68
177	Further delineation of Kabuki syndrome in 48 well-defined new individuals 2005 , 132A, 265-72		68
176	UBQLN4 Represses Homologous Recombination and Is Overexpressed in Aggressive Tumors. <i>Cell</i> , 2019 , 176, 505-519.e22	56.2	68
175	Integrative analysis revealed the molecular mechanism underlying RBM10-mediated splicing regulation. <i>EMBO Molecular Medicine</i> , 2013 , 5, 1431-42	12	67
174	Genetic determination of human facial morphology: links between cleft-lips and normal variation. <i>European Journal of Human Genetics</i> , 2011 , 19, 1192-7	5.3	67
173	Loss of the BMP antagonist, SMOC-1, causes Ophthalmo-acromelic (Waardenburg Anophthalmia) syndrome in humans and mice. <i>PLoS Genetics</i> , 2011 , 7, e1002114	6	67

172	Extreme growth failure is a common presentation of ligase IV deficiency. <i>Human Mutation</i> , 2014 , 35, 76-85	4.7	63
171	Mutations in ZIC2 in human holoprosencephaly: description of a novel ZIC2 specific phenotype and comprehensive analysis of 157 individuals. <i>Journal of Medical Genetics</i> , 2010 , 47, 513-24	5.8	63
170	Syndrome identification based on 2D analysis software. <i>European Journal of Human Genetics</i> , 2006 , 14, 1082-9	5.3	63
169	De novo mutations in beta-catenin (CTNNB1) appear to be a frequent cause of intellectual disability: expanding the mutational and clinical spectrum. <i>Human Genetics</i> , 2015 , 134, 97-109	6.3	62
168	The ARID1B phenotype: what we have learned so far. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014 , 166C, 276-89	3.1	62
167	A review of craniofacial disorders caused by spliceosomal defects. <i>Clinical Genetics</i> , 2015 , 88, 405-15	4	60
166	Is there a higher incidence of maternal uniparental disomy 14 [upd(14)mat]? Detection of 10 new patients by methylation-specific PCR. <i>American Journal of Medical Genetics, Part A</i> , 2006 , 140, 2039-49	2.5	57
165	Computer-based recognition of dysmorphic faces. <i>European Journal of Human Genetics</i> , 2003 , 11, 555-60	5.3	56
164	De Novo Mutations in CHAMP1 Cause Intellectual Disability with Severe Speech Impairment. <i>American Journal of Human Genetics</i> , 2015 , 97, 493-500	11	55
163	Loss-of-function variants of SETD5 cause intellectual disability and the core phenotype of microdeletion 3p25.3 syndrome. <i>European Journal of Human Genetics</i> , 2015 , 23, 753-60	5.3	55
162	Histone acetylation dependent allelic expression imbalance of BAPX1 in patients with the oculo-auriculo-vertebral spectrum. <i>Human Molecular Genetics</i> , 2006 , 15, 581-7	5.6	55
161	Clinical and mutation data in 12 patients with the clinical diagnosis of Nager syndrome. <i>Human Genetics</i> , 2013 , 132, 885-98	6.3	54
160	Two patients with EP300 mutations and facial dysmorphism different from the classic Rubinstein-Taybi syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2010 , 152A, 181-4	2.5	52
159	Phenotype and genotype in Nicolaidis-Baraitser syndrome. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014 , 166C, 302-14	3.1	51
158	Microcephaly with or without chorioretinopathy, lymphoedema, or mental retardation (MCLMR): review of phenotype associated with KIF11 mutations. <i>European Journal of Human Genetics</i> , 2014 , 22, 881-7	5.3	51
157	Human facial dysostoses. <i>Clinical Genetics</i> , 2013 , 83, 499-510	4	51
156	Miller (Genee-Wiedemann) syndrome represents a clinically and biochemically distinct subgroup of postaxial acrofacial dysostosis associated with partial deficiency of DHODH. <i>Human Molecular Genetics</i> , 2012 , 21, 3969-83	5.6	51
155	Rare copy number variants are a common cause of short stature. <i>PLoS Genetics</i> , 2013 , 9, e1003365	6	50

154	Diagnostic algorithms in Charcot-Marie-Tooth neuropathies: experiences from a German genetic laboratory on the basis of 1206 index patients. <i>Clinical Genetics</i> , 2016 , 89, 34-43	4	48
153	RAP1-mediated MEK/ERK pathway defects in Kabuki syndrome. <i>Journal of Clinical Investigation</i> , 2015 , 125, 3585-99	15.9	48
152	Exome sequencing unravels unexpected differential diagnoses in individuals with the tentative diagnosis of Coffin-Siris and Nicolaidis-Baraitser syndromes. <i>Human Genetics</i> , 2015 , 134, 553-68	6.3	47
151	The face of Noonan syndrome: Does phenotype predict genotype. <i>American Journal of Medical Genetics, Part A</i> , 2010 , 152A, 1960-6	2.5	47
150	Novel mutations in BCOR in three patients with oculo-facio-cardio-dental syndrome, but none in Lenz microphthalmia syndrome. <i>European Journal of Human Genetics</i> , 2005 , 13, 563-9	5.3	47
149	Acrofacial Dysostosis, Cincinnati Type, a Mandibulofacial Dysostosis Syndrome with Limb Anomalies, Is Caused by POLR1A Dysfunction. <i>American Journal of Human Genetics</i> , 2015 , 96, 765-74	11	46
148	Further delineation of the KAT6B molecular and phenotypic spectrum. <i>European Journal of Human Genetics</i> , 2015 , 23, 1165-70	5.3	45
147	Compound heterozygosity of low-frequency promoter deletions and rare loss-of-function mutations in TXNL4A causes Burn-McKeown syndrome. <i>American Journal of Human Genetics</i> , 2014 , 95, 698-707	11	45
146	De Novo Mutations in SON Disrupt RNA Splicing of Genes Essential for Brain Development and Metabolism, Causing an Intellectual-Disability Syndrome. <i>American Journal of Human Genetics</i> , 2016 , 99, 711-719	11	44
145	Biallelic loss of function of the promyelocytic leukaemia zinc finger (PLZF) gene causes severe skeletal defects and genital hypoplasia. <i>Journal of Medical Genetics</i> , 2008 , 45, 731-7	5.8	43
144	Reproduction abnormalities and twin pregnancies in parents of sporadic patients with oculo-auriculo-vertebral spectrum/Goldenhar syndrome. <i>Human Genetics</i> , 2007 , 121, 369-76	6.3	41
143	Identification of 34 novel and 56 known FOXL2 mutations in patients with Blepharophimosis syndrome. <i>Human Mutation</i> , 2008 , 29, E205-19	4.7	41
142	Altered development of NKT cells, Γ cells, CD8 T cells and NK cells in a PLZF deficient patient. <i>PLoS ONE</i> , 2011 , 6, e24441	3.7	41
141	Mutations in chromatin regulators functionally link Cornelia de Lange syndrome and clinically overlapping phenotypes. <i>Human Genetics</i> , 2017 , 136, 307-320	6.3	40
140	Autosomal recessive mental retardation: homozygosity mapping identifies 27 single linkage intervals, at least 14 novel loci and several mutation hotspots. <i>Human Genetics</i> , 2011 , 129, 141-8	6.3	40
139	Heterozygous HNRNPU variants cause early onset epilepsy and severe intellectual disability. <i>Human Genetics</i> , 2017 , 136, 821-834	6.3	39
138	Oto-facial syndrome and esophageal atresia, intellectual disability and zygomatic anomalies - expanding the phenotypes associated with EFTUD2 mutations. <i>Orphanet Journal of Rare Diseases</i> , 2013 , 8, 110	4.2	39
137	Bohring-Opitz (Oberklaid-Danks) syndrome: clinical study, review of the literature, and discussion of possible pathogenesis. <i>European Journal of Human Genetics</i> , 2011 , 19, 513-9	5.3	38

136	Mutations in the BAF-Complex Subunit DPF2 Are Associated with Coffin-Siris Syndrome. <i>American Journal of Human Genetics</i> , 2018 , 102, 468-479	11	37
135	Genetic screening confirms heterozygous mutations in ACAN as a major cause of idiopathic short stature. <i>Scientific Reports</i> , 2017 , 7, 12225	4.9	37
134	De novo MECP2 duplication in two females with random X-inactivation and moderate mental retardation. <i>European Journal of Human Genetics</i> , 2011 , 19, 507-12	5.3	37
133	BCL11B mutations in patients affected by a neurodevelopmental disorder with reduced type 2 innate lymphoid cells. <i>Brain</i> , 2018 , 141, 2299-2311	11.2	36
132	Autosomal recessive POLR1D mutation with decrease of TCOF1 mRNA is responsible for Treacher Collins syndrome. <i>Genetics in Medicine</i> , 2014 , 16, 720-4	8.1	36
131	Congenital diaphragmatic hernia interval on chromosome 8p23.1 characterized by genetics and protein interaction networks. <i>American Journal of Medical Genetics, Part A</i> , 2012 , 158A, 3148-58	2.5	36
130	A new face of Borjeson-Forssman-Lehmann syndrome? De novo mutations in PHF6 in seven females with a distinct phenotype. <i>Journal of Medical Genetics</i> , 2013 , 50, 838-47	5.8	36
129	The ARID1B spectrum in 143 patients: from nonsyndromic intellectual disability to Coffin-Siris syndrome. <i>Genetics in Medicine</i> , 2019 , 21, 1295-1307	8.1	36
128	WilmsRTumor 1 Gene Expression Using a Standardized European LeukemiaNet-Certified Assay Compared to Other Methods for Detection of Minimal Residual Disease in Myelodysplastic Syndrome and Acute Myelogenous Leukemia after Allogeneic Blood Stem Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2018 , 24, 2337-2343	4.7	36
127	Novel mutations including deletions of the entire OFD1 gene in 30 families with type 1 orofacioidigital syndrome: a study of the extensive clinical variability. <i>Human Mutation</i> , 2013 , 34, 237-47	4.7	35
126	Cardio-facio-cutaneous (CFC) syndrome--a distinct entity? Report of three patients demonstrating the diagnostic difficulties in delineation of CFC syndrome. <i>Clinical Genetics</i> , 1997 , 52, 37-46	4	35
125	Mutations in MAP3K7 that Alter the Activity of the TAK1 Signaling Complex Cause Frontometaphyseal Dysplasia. <i>American Journal of Human Genetics</i> , 2016 , 99, 392-406	11	34
124	Genetic Analysis of PAX6-Negative Individuals with Aniridia or Gillespie Syndrome. <i>PLoS ONE</i> , 2016 , 11, e0153757	3.7	34
123	Heterozygosity for ARID2 loss-of-function mutations in individuals with a Coffin-Siris syndrome-like phenotype. <i>Human Genetics</i> , 2017 , 136, 297-305	6.3	33
122	Bi-allelic Variants in METTL5 Cause Autosomal-Recessive Intellectual Disability and Microcephaly. <i>American Journal of Human Genetics</i> , 2019 , 105, 869-878	11	33
121	DOORS syndrome: phenotype, genotype and comparison with Coffin-Siris syndrome. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014 , 166C, 327-32	3.1	33
120	Expanding the phenotype of IQSEC2 mutations: truncating mutations in severe intellectual disability. <i>European Journal of Human Genetics</i> , 2014 , 22, 289-92	5.3	33
119	WDR26 Haploinsufficiency Causes a Recognizable Syndrome of Intellectual Disability, Seizures, Abnormal Gait, and Distinctive Facial Features. <i>American Journal of Human Genetics</i> , 2017 , 101, 139-148	11	31

118	WDR73 Mutations Cause Infantile Neurodegeneration and Variable Glomerular Kidney Disease. <i>Human Mutation</i> , 2015 , 36, 1021-8	4.7	31
117	Early IFN- β signatures and persistent dysfunction are distinguishing features of NK cells in severe COVID-19. <i>Immunity</i> , 2021 , 54, 2650-2669.e14	32.3	31
116	Behavioral phenotype in five individuals with de novo mutations within the GRIN2B gene. <i>Behavioral and Brain Functions</i> , 2013 , 9, 20	4.1	30
115	Delineation of PIGV mutation spectrum and associated phenotypes in hyperphosphatasia with mental retardation syndrome. <i>European Journal of Human Genetics</i> , 2014 , 22, 762-7	5.3	30
114	Homozygous and compound-heterozygous mutations in TGDS cause Catel-Manzke syndrome. <i>American Journal of Human Genetics</i> , 2014 , 95, 763-70	11	30
113	Automated syndrome detection in a set of clinical facial photographs. <i>American Journal of Medical Genetics, Part A</i> , 2011 , 155A, 2161-9	2.5	30
112	Microcephalic osteodysplastic primordial dwarfism type I with biallelic mutations in the RNU4ATAC gene. <i>Clinical Genetics</i> , 2012 , 82, 140-6	4	29
111	Identification of new TRIP12 variants and detailed clinical evaluation of individuals with non-syndromic intellectual disability with or without autism. <i>Human Genetics</i> , 2017 , 136, 179-192	6.3	29
110	Goltz-Gorlin (focal dermal hypoplasia) and the microphthalmia with linear skin defects (MLS) syndrome: no evidence of genetic overlap. <i>European Journal of Human Genetics</i> , 2009 , 17, 1207-15	5.3	29
109	Expanding the clinical spectrum of the HDAC8-phenotype: implications for molecular diagnostics, counseling and risk prediction. <i>Clinical Genetics</i> , 2016 , 89, 564-73	4	29
108	Loss-of-function variants in HIVEP2 are a cause of intellectual disability. <i>European Journal of Human Genetics</i> , 2016 , 24, 556-61	5.3	28
107	Bainbridge-Ropers syndrome caused by loss-of-function variants in ASXL3: a recognizable condition. <i>European Journal of Human Genetics</i> , 2017 , 25, 183-191	5.3	28
106	Cohen syndrome diagnosis using whole genome arrays. <i>Journal of Medical Genetics</i> , 2011 , 48, 136-40	5.8	28
105	Infectious and immunologic phenotype of MECP2 duplication syndrome. <i>Journal of Clinical Immunology</i> , 2015 , 35, 168-81	5.7	27
104	POLR1B and neural crest cell anomalies in Treacher Collins syndrome type 4. <i>Genetics in Medicine</i> , 2020 , 22, 547-556	8.1	26
103	Microcephaly, microtia, preauricular tags, choanal atresia and developmental delay in three unrelated patients: a mandibulofacial dysostosis distinct from Treacher Collins syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2009 , 149A, 837-43	2.5	25
102	Genotype-phenotype correlation in eight new patients with a deletion encompassing 2q31.1. <i>American Journal of Medical Genetics, Part A</i> , 2010 , 152A, 1213-24	2.5	25
101	De Novo and Inherited Loss-of-Function Variants in TLK2: Clinical and Genotype-Phenotype Evaluation of a Distinct Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2018 , 102, 1195-1203	11	24

100	Impact of geometry and viewing angle on classification accuracy of 2D based analysis of dysmorphic faces. <i>European Journal of Medical Genetics</i> , 2008 , 51, 44-53	2.6	24
99	Left-ventricular non-compaction (LVNC): a clinical feature more often observed in terminal deletion 1p36 than previously expected. <i>European Journal of Medical Genetics</i> , 2008 , 51, 685-8	2.6	23
98	New insights into the imprinted MEG8-DMR in 14q32 and clinical and molecular description of novel patients with Temple syndrome. <i>European Journal of Human Genetics</i> , 2017 , 25, 935-945	5.3	21
97	Females with de novo aberrations in PHF6: clinical overlap of Borjeson-Forssman-Lehmann with Coffin-Siris syndrome. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2014 , 166C, 290-301	3.1	21
96	Spinocerebellar ataxia 28: a novel AFG3L2 mutation in a German family with young onset, slow progression and saccadic slowing. <i>Cerebellum and Ataxias</i> , 2015 , 2, 19	1.7	21
95	Penetrance and Expressivity in Inherited Cancer Predisposing Syndromes. <i>Trends in Cancer</i> , 2018 , 4, 718-728	7.2	21
94	Elucidating the genetic architecture of Adams-Oliver syndrome in a large European cohort. <i>Human Mutation</i> , 2018 , 39, 1246-1261	4.7	21
93	Genetic predisposition in children with cancer - affected families: acceptance of Trio-WES. <i>European Journal of Pediatrics</i> , 2018 , 177, 53-60	4.1	20
92	De novo nonsense and frameshift variants of TCF20 in individuals with intellectual disability and postnatal overgrowth. <i>European Journal of Human Genetics</i> , 2016 , 24, 1739-1745	5.3	20
91	Wide clinical variability in conditions with coarse facial features and hypertrichosis caused by mutations in ABCC9. <i>American Journal of Medical Genetics, Part A</i> , 2013 , 161A, 295-300	2.5	20
90	Five patients with novel overlapping interstitial deletions in 8q22.2q22.3. <i>American Journal of Medical Genetics, Part A</i> , 2011 , 155A, 1857-64	2.5	20
89	De novo FBXO11 mutations are associated with intellectual disability and behavioural anomalies. <i>Human Genetics</i> , 2018 , 137, 401-411	6.3	20
88	Variants in CPLX1 in two families with autosomal-recessive severe infantile myoclonic epilepsy and ID. <i>European Journal of Human Genetics</i> , 2017 , 25, 889-893	5.3	19
87	Novel microdeletions on chromosome 14q32.2 suggest a potential role for non-coding RNAs in Kagami-Ogata syndrome. <i>European Journal of Human Genetics</i> , 2016 , 24, 1724-1729	5.3	19
86	Genetic variants in components of the NALCN-UNC80-UNC79 ion channel complex cause a broad clinical phenotype (NALCN channelopathies). <i>Human Genetics</i> , 2018 , 137, 753-768	6.3	19
85	Treacher Collins syndrome: clinical implications for the paediatrician--a new mutation in a severely affected newborn and comparison with three further patients with the same mutation, and review of the literature. <i>European Journal of Pediatrics</i> , 2012 , 171, 1611-8	4.1	18
84	Autosomal dominant spinal muscular atrophy with lower extremity predominance: A recognizable phenotype of BICD2 mutations. <i>Muscle and Nerve</i> , 2016 , 54, 496-500	3.4	18
83	De Novo Mutations Affecting the Catalytic C _ε Subunit of PP2A, PPP2CA, Cause Syndromic Intellectual Disability Resembling Other PP2A-Related Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2019 , 104, 139-156	11	18

82	Autosomal dominant intellectual disability. <i>Medizinische Genetik</i> , 2018 , 30, 318-322	0.5	18
81	Splitting versus lumping Temple-Baraitser and Zimmermann-Laband Syndromes. <i>Human Genetics</i> , 2015 , 134, 1089-97	6.3	17
80	Family-based germline sequencing in children with cancer. <i>Oncogene</i> , 2019 , 38, 1367-1380	9.2	17
79	Autosomal dominant frontometaphyseal dysplasia: Delineation of the clinical phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2017 , 173, 1739-1746	2.5	16
78	Next-generation sequencing of 32 genes associated with hereditary aortopathies and related disorders of connective tissue in a cohort of 199 patients. <i>Genetics in Medicine</i> , 2019 , 21, 1832-1841	8.1	16
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