

Anita Rauch

List of Publications by Year in descending order

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256
papers

17,020
citations

15504

65
h-index

19749

117
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270
all docs

270
docs citations

270
times ranked

23224
citing authors

#	ARTICLE	IF	CITATIONS
1	Range of genetic mutations associated with severe non-syndromic sporadic intellectual disability: an exome sequencing study. <i>Lancet, The</i> , 2012, 380, 1674-1682.	13.7	940
2	Germline KRAS mutations cause Noonan syndrome. <i>Nature Genetics</i> , 2006, 38, 331-336.	21.4	670
3	CNTNAP2 and NRXN1 Are Mutated in Autosomal-Recessive Pitt-Hopkins-like Mental Retardation and Determine the Level of a Common Synaptic Protein in <i>Drosophila</i> . <i>American Journal of Human Genetics</i> , 2009, 85, 655-666.	6.2	573
4	Transcription Factor E2-2 Is an Essential and Specific Regulator of Plasmacytoid Dendritic Cell Development. <i>Cell</i> , 2008, 135, 37-48.	28.9	567
5	Mutations in GRIN2A and GRIN2B encoding regulatory subunits of NMDA receptors cause variable neurodevelopmental phenotypes. <i>Nature Genetics</i> , 2010, 42, 1021-1026.	21.4	431
6	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. <i>Nature</i> , 2011, 478, 97-102.	27.8	394
7	Mutations in the Pericentrin (<i>PCNT</i>) Gene Cause Primordial Dwarfism. <i>Science</i> , 2008, 319, 816-819.	12.6	370
8	Diagnostic yield of various genetic approaches in patients with unexplained developmental delay or mental retardation. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 2063-2074.	1.2	343
9	Mutations in STRA6 Cause a Broad Spectrum of Malformations Including Anophthalmia, Congenital Heart Defects, Diaphragmatic Hernia, Alveolar Capillary Dysplasia, Lung Hypoplasia, and Mental Retardation. <i>American Journal of Human Genetics</i> , 2007, 80, 550-560.	6.2	316
10	Contribution of Global Rare Copy-Number Variants to the Risk of Sporadic Congenital Heart Disease. <i>American Journal of Human Genetics</i> , 2012, 91, 489-501.	6.2	272
11	Haploinsufficiency of TCF4 Causes Syndromal Mental Retardation with Intermittent Hyperventilation (Pitt-Hopkins Syndrome). <i>American Journal of Human Genetics</i> , 2007, 80, 994-1001.	6.2	261
12	De novo nonsense mutations in ASXL1 cause Bohring-Opitz syndrome. <i>Nature Genetics</i> , 2011, 43, 729-731.	21.4	236
13	Genotype-phenotype correlations in Noonan syndrome. <i>Journal of Pediatrics</i> , 2004, 144, 368-374.	1.8	227
14	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, 565-572.	6.2	225
15	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.	3.2	220
16	CEP152 is a genome maintenance protein disrupted in Seckel syndrome. <i>Nature Genetics</i> , 2011, 43, 23-26.	21.4	201
17	Human TBX1 Missense Mutations Cause Gain of Function Resulting in the Same Phenotype as 22q11.2 Deletions. <i>American Journal of Human Genetics</i> , 2007, 80, 510-517.	6.2	195
18	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. <i>JAMA Psychiatry</i> , 2016, 73, 20.	11.0	195

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19	Further clinical and molecular delineation of the 9q subtelomeric deletion syndrome supports a major contribution of EHMT1 haploinsufficiency to the core phenotype. <i>Journal of Medical Genetics</i> , 2009, 46, 598-606.	3.2	194
20	Clinical and molecular delineation of the 17q21.31 microdeletion syndrome. <i>Journal of Medical Genetics</i> , 2008, 45, 710-720.	3.2	191
21	A comprehensive molecular study on Coffinâ€“Siris and Nicolaidesâ€“Baraitser syndromes identifies a broad molecular and clinical spectrum converging on altered chromatin remodeling. <i>Human Molecular Genetics</i> , 2013, 22, 5121-5135.	2.9	190
22	Identification of a novel loss-of-function calcium channel gene mutation in short QT syndrome (SQTS6). <i>European Heart Journal</i> , 2011, 32, 1077-1088.	2.2	178
23	Induction, binding specificity and function of human ICOS. <i>European Journal of Immunology</i> , 2000, 30, 3707-3717.	2.9	166
24	Mutations in MEF2C from the 5q14.3q15 microdeletion syndrome region are a frequent cause of severe mental retardation and diminish MECP2 and CDKL5 expression. <i>Human Mutation</i> , 2010, 31, 722-733.	2.5	163
25	Mutations at the SALL4 locus on chromosome 20 result in a range of clinically overlapping phenotypes, including Okhiro syndrome, Holt-Oram syndrome, acro-renal-ocular syndrome, and patients previously reported to represent thalidomide embryopathy. <i>Journal of Medical Genetics</i> , 2003, 40, 473-478.	3.2	159
26	NEK1 Mutations Cause Short-Rib Polydactyly Syndrome Type Majewski. <i>American Journal of Human Genetics</i> , 2011, 88, 106-114.	6.2	151
27	De Novo Mutations in the Genome Organizer CTCF Cause Intellectual Disability. <i>American Journal of Human Genetics</i> , 2013, 93, 124-131.	6.2	151
28	Genotyping in 46 patients with tentative diagnosis of Treacher Collins syndrome revealed unexpected phenotypic variation. <i>European Journal of Human Genetics</i> , 2004, 12, 879-890.	2.8	149
29	Elastin: mutational spectrum in supravalvular aortic stenosis. <i>European Journal of Human Genetics</i> , 2000, 8, 955-963.	2.8	147
30	PDE3A mutations cause autosomal dominant hypertension with brachydactyly. <i>Nature Genetics</i> , 2015, 47, 647-653.	21.4	146
31	Guidelines for molecular karyotyping in constitutional genetic diagnosis. <i>European Journal of Human Genetics</i> , 2007, 15, 1105-1114.	2.8	144
32	Genetic and neurodevelopmental spectrum of <i>SYNGAP1</i> -associated intellectual disability and epilepsy. <i>Journal of Medical Genetics</i> , 2016, 53, 511-522.	3.2	135
33	Altered TGFÎ² signaling and cardiovascular manifestations in patients with autosomal recessive cutis laxa type I caused by fibulin-4 deficiency. <i>European Journal of Human Genetics</i> , 2010, 18, 895-901.	2.8	132
34	SMCHD1 mutations associated with a rare muscular dystrophy can also cause isolated arhinia and Bosma arhinia microphthalmia syndrome. <i>Nature Genetics</i> , 2017, 49, 238-248.	21.4	131
35	First known microdeletion within the Wolf-Hirschhorn syndrome critical region refines genotype-phenotype correlation. <i>American Journal of Medical Genetics Part A</i> , 2001, 99, 338-342.	2.4	128
36	Comprehensive genotype-phenotype analysis in 230 patients with tetralogy of Fallot. <i>Journal of Medical Genetics</i> , 2010, 47, 321-331.	3.2	126

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37	YY1 Haploinsufficiency Causes an Intellectual Disability Syndrome Featuring Transcriptional and Chromatin Dysfunction. <i>American Journal of Human Genetics</i> , 2017, 100, 907-925.	6.2	125
38	“Mowat-Wilson” syndrome with and without Hirschsprung disease is a distinct, recognizable multiple congenital anomalies-mental retardation syndrome caused by mutations in the zinc finger homeo box 1B gene. <i>American Journal of Medical Genetics Part A</i> , 2002, 108, 177-181.	2.4	122
39	Clinical and Mutational Spectrum of Mowat-Wilson Syndrome. <i>European Journal of Medical Genetics</i> , 2005, 48, 97-111.	1.3	121
40	Systematic assessment of atypical deletions reveals genotype-phenotype correlation in 22q11.2. <i>Journal of Medical Genetics</i> , 2005, 42, 871-876.	3.2	118
41	Severely Incapacitating Mutations in Patients with Extreme Short Stature Identify RNA-Processing Endoribonuclease RMRP as an Essential Cell Growth Regulator. <i>American Journal of Human Genetics</i> , 2005, 77, 795-806.	6.2	117
42	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, 651-656.	3.2	114
43	Expanding the clinical spectrum associated with defects in CNTNAP2 and NRXN1. <i>BMC Medical Genetics</i> , 2011, 12, 106.	2.1	109
44	Molecular karyotyping using an SNP array for genomewide genotyping. <i>Journal of Medical Genetics</i> , 2004, 41, 916-922.	3.2	106
45	Phenotype-specific effect of chromosome 1q21.1 rearrangements and GJA5 duplications in 2436 congenital heart disease patients and 6760 controls. <i>Human Molecular Genetics</i> , 2012, 21, 1513-1520.	2.9	101
46	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. <i>Genetics in Medicine</i> , 2018, 20, 630-638.	2.4	101
47	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, 1592-1599.	2.8	96
48	A Novel 22q11.2 Microdeletion in DiGeorge Syndrome. <i>American Journal of Human Genetics</i> , 1999, 64, 659-667.	6.2	95
49	Disruption of the histone acetyltransferase MYST4 leads to a Noonan syndrome-like phenotype and hyperactivated MAPK signaling in humans and mice. <i>Journal of Clinical Investigation</i> , 2011, 121, 3479-3491.	8.2	89
50	Gene discovery for Mendelian conditions via social networking: de novo variants in KDM1A cause developmental delay and distinctive facial features. <i>Genetics in Medicine</i> , 2016, 18, 788-795.	2.4	88
51	Further delineation of Pitt-Hopkins syndrome: phenotypic and genotypic description of 16 novel patients. <i>Journal of Medical Genetics</i> , 2008, 45, 738-744.	3.2	86
52	Nephrocalcinosis (Enamel Renal Syndrome) Caused by Autosomal Recessive FAM20A Mutations. <i>Nephron Physiology</i> , 2013, 122, 1-6.	1.2	84
53	Microcephalin and pericentrin regulate mitotic entry via centrosome-associated Chk1. <i>Journal of Cell Biology</i> , 2009, 185, 1149-1157.	5.2	83
54	Disturbed Wnt Signalling due to a Mutation in CCDC88C Causes an Autosomal Recessive Non-Syndromic Hydrocephalus with Medial Diverticulum. <i>Molecular Syndromology</i> , 2010, 1, 99-112.	0.8	82

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55	Genome-wide association study identifies loci on 12q24 and 13q32 associated with Tetralogy of Fallot. <i>Human Molecular Genetics</i> , 2013, 22, 1473-1481.	2.9	82
56	Deletion mapping on chromosome 10p and definition of a critical region for the second DiGeorge syndrome locus (DGS2). <i>European Journal of Human Genetics</i> , 1998, 6, 213-225.	2.8	81
57	A study of ten small supernumerary (marker) chromosomes identified by fluorescence <i>in situ</i> hybridization (FISH). <i>Clinical Genetics</i> , 1992, 42, 84-90.	2.0	81
58	Type and Level of RMRP Functional Impairment Predicts Phenotype in the Cartilage Hair Hypoplasia–Anauxetic Dysplasia Spectrum. <i>American Journal of Human Genetics</i> , 2007, 81, 519-529.	6.2	78
59	Crisponi Syndrome Is Caused by Mutations in the CRLF1 Gene and Is Allelic to Cold-Induced Sweating Syndrome Type 1. <i>American Journal of Human Genetics</i> , 2007, 80, 971-981.	6.2	76
60	Allelic Heterogeneity in the COH1 Gene Explains Clinical Variability in Cohen Syndrome. <i>American Journal of Human Genetics</i> , 2004, 75, 138-145.	6.2	72
61	Klinefelter syndrome and mediastinal germ cell tumors. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 471-481.	1.2	72
62	Molecular karyotyping in patients with mental retardation using 100K single-nucleotide polymorphism arrays. <i>Journal of Medical Genetics</i> , 2007, 44, 629-636.	3.2	72
63	The clinical significance of small copy number variants in neurodevelopmental disorders. <i>Journal of Medical Genetics</i> , 2014, 51, 677-688.	3.2	72
64	De novo missense mutations in the NAA10 gene cause severe non-syndromic developmental delay in males and females. <i>European Journal of Human Genetics</i> , 2015, 23, 602-609.	2.8	72
65	Dosage-Dependent Severity of the Phenotype in Patients with Mental Retardation Due to a Recurrent Copy-Number Gain at Xq28 Mediated by an Unusual Recombination. <i>American Journal of Human Genetics</i> , 2009, 85, 809-822.	6.2	70
66	Need for high-resolution Genetic Analysis in iPSC: Results and Lessons from the ForIPS Consortium. <i>Scientific Reports</i> , 2018, 8, 17201.	3.3	70
67	Biallelic UFM1 and UFC1 mutations expand the essential role of ufmylation in brain development. <i>Brain</i> , 2018, 141, 1934-1945.	7.6	70
68	<i>FOXP2</i> variants in 14 individuals with developmental speech and language disorders broaden the mutational and clinical spectrum. <i>Journal of Medical Genetics</i> , 2017, 54, 64-72.	3.2	67
69	Phenotype and genotype of 87 patients with Mowat–Wilson syndrome and recommendations for care. <i>Genetics in Medicine</i> , 2018, 20, 965-975.	2.4	67
70	Confirmation of mutations in <i>PROSC</i> as a novel cause of vitamin B ₆ -dependent epilepsy. <i>Journal of Medical Genetics</i> , 2017, 54, 809-814.	3.2	66
71	Incidence and significance of 22q11.2 hemizyosity in patients with interrupted aortic arch. , 1998, 78, 322-331.		62
72	Characterisation of deletions of the ZFX1B region and genotype-phenotype analysis in Mowat-Wilson syndrome. <i>Journal of Medical Genetics</i> , 2003, 40, 601-605.	3.2	61

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73	Systematic survey of variants in TBX1 in non-syndromic tetralogy of Fallot identifies a novel 57 base pair deletion that reduces transcriptional activity but finds no evidence for association with common variants. <i>Heart</i> , 2010, 96, 1651-1655.	2.9	61
74	Rare Copy Number Variants Are a Common Cause of Short Stature. <i>PLoS Genetics</i> , 2013, 9, e1003365.	3.5	60
75	The face of Noonan syndrome: Does phenotype predict genotype. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 1960-1966.	1.2	59
76	Mutations Affecting the SAND Domain of DEAF1 Cause Intellectual Disability with Severe Speech Impairment and Behavioral Problems. <i>American Journal of Human Genetics</i> , 2014, 94, 649-661.	6.2	59
77	Hirschsprung disease, mental retardation, characteristic facial features, and mutation in the gene <i>ZFX1B</i> (<i>SIP1</i>): Confirmation of the Mowat-Wilson syndrome. <i>American Journal of Medical Genetics Part A</i> , 2003, 116A, 385-388.	2.4	58
78	Laterality of the aortic arch and anomalies of the subclavian artery?reliable indicators for 22q11.2 deletion syndromes?. <i>European Journal of Pediatrics</i> , 2004, 163, 642-5.	2.7	58
79	In-Frame Deletion and Missense Mutations of the C-Terminal Helicase Domain of SMARCA2 in Three Patients with Nicolaides-Baraitser Syndrome. <i>Molecular Syndromology</i> , 2011, 2, 237-244.	0.8	58
80	A new quantitative PCR multiplex assay for rapid analysis of chromosome 17p11.2-12 duplications and deletions leading to HMSN/HNPP. <i>European Journal of Human Genetics</i> , 2003, 11, 170-178.	2.8	57
81	Dosage changes of MED13L further delineate its role in congenital heart defects and intellectual disability. <i>European Journal of Human Genetics</i> , 2013, 21, 1100-1104.	2.8	57
82	Elucidation of the phenotypic spectrum and genetic landscape in primary and secondary microcephaly. <i>Genetics in Medicine</i> , 2019, 21, 2043-2058.	2.4	57
83	Highly variable cutis laxa resulting from a dominant splicing mutation of the elastin gene. <i>American Journal of Medical Genetics, Part A</i> , 2008, 146A, 977-983.	1.2	56
84	Heterozygous Variants in KMT2E Cause a Spectrum of Neurodevelopmental Disorders and Epilepsy. <i>American Journal of Human Genetics</i> , 2019, 104, 1210-1222.	6.2	56
85	A Dual Phenotype of Periventricular Nodular Heterotopia and Frontometaphyseal Dysplasia in One Patient Caused by a Single FLNA Mutation Leading to Two Functionally Different Aberrant Transcripts. <i>American Journal of Human Genetics</i> , 2004, 74, 731-737.	6.2	55
86	Independent <i>NF1</i> and <i>PTPN11</i> mutations in a family with neurofibromatosis-Noonan syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 1263-1267.	1.2	55
87	Mutations in <i>CDK5</i> and <i>RAP2</i> cause Seckel syndrome. <i>Molecular Genetics & Genomic Medicine</i> , 2015, 3, 467-480.	1.2	55
88	Clinical and molecular cytogenetic observations in three cases of trisomy 12p syndrome. <i>American Journal of Medical Genetics Part A</i> , 1996, 63, 243-249.	2.4	54
89	A Potential Contributory Role for Ciliary Dysfunction in the 16p11.2 600 kb BP4-BP5 Pathology. <i>American Journal of Human Genetics</i> , 2015, 96, 784-796.	6.2	53
90	Autosomal recessive primary microcephaly due to <i>ASPM</i> mutations: An update. <i>Human Mutation</i> , 2018, 39, 319-332.	2.5	53

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91	The role of recessive inheritance in early-onset epileptic encephalopathies: a combined whole-exome sequencing and copy number study. <i>European Journal of Human Genetics</i> , 2019, 27, 408-421.	2.8	52
92	The MEF2C-Related and 5q14.3q15 Microdeletion Syndrome. <i>Molecular Syndromology</i> , 2011, 2, 164-170.	0.8	51
93	Lethal cutis laxa with contractural arachnodactyly, overgrowth and soft tissue bleeding due to a novel homozygous <i>fibulin4</i> gene mutation. <i>Clinical Genetics</i> , 2009, 76, 276-281.	2.0	50
94	A new face of Borjesonâ€“Forssmanâ€“Lehmann syndrome? De novo mutations in <i>PHF6</i> in seven females with a distinct phenotype. <i>Journal of Medical Genetics</i> , 2013, 50, 838-847.	3.2	50
95	Mutations in CRADD Result in Reduced Caspase-2-Mediated Neuronal Apoptosis and Cause Megalencephaly with a Rare Lissencephaly Variant. <i>American Journal of Human Genetics</i> , 2016, 99, 1117-1129.	6.2	50
96	Partial deletion of the critical 1.5 Mb interval in Williams-Beuren syndrome. <i>Journal of Medical Genetics</i> , 2003, 40, 99e-99.	3.2	49
97	De Novo Variants in the F-Box Protein FBXO11 in 20 Individuals with a Variable Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2018, 103, 305-316.	6.2	48
98	SPEN haploinsufficiency causes a neurodevelopmental disorder overlapping proximal 1p36 deletion syndrome with an epismutation of X chromosomes in females. <i>American Journal of Human Genetics</i> , 2021, 108, 502-516.	6.2	48
99	Intragenic <i>KANSL1</i> mutations and chromosome 17q21.31 deletions: broadening the clinical spectrum and genotypeâ€“phenotype correlations in a large cohort of patients. <i>Journal of Medical Genetics</i> , 2015, 52, 804-814.	3.2	47
100	Spatially clustering de novo variants in CYFIP2, encoding the cytoplasmic FMRP interacting protein 2, cause intellectual disability and seizures. <i>European Journal of Human Genetics</i> , 2019, 27, 747-759.	2.8	47
101	Mosaicism for the Charcot-Marie-Tooth disease type 1A duplication suggests somatic reversion. <i>Human Genetics</i> , 1996, 98, 22-28.	3.8	46
102	A variable combination of features of Noonan syndrome and neurofibromatosis type I are caused by mutations in the <i>NF1</i> gene. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 2749-2756.	1.2	46
103	The molecular basis of the cartilage-hair hypoplasiaâ€“anaxetic dysplasia spectrum. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2011, 25, 131-142.	4.7	46
104	Monosomy 22q11 in patients with pulmonary atresia, ventricular septal defect, and major aortopulmonary collateral arteries. <i>Heart</i> , 1998, 79, 180-185.	2.9	45
105	Monoallelic BMP2 Variants Predicted to Result in Haploinsufficiency Cause Craniofacial, Skeletal, and Cardiac Features Overlapping Those of 20p12 Deletions. <i>American Journal of Human Genetics</i> , 2017, 101, 985-994.	6.2	44
106	The PDAC syndrome (pulmonary hypoplasia/agenesis, diaphragmatic hernia/eventration,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 152 Td (Report of eight cases including a living child and further evidence for autosomal recessive inheritance. <i>American Journal of Medical Genetics, Part A</i> , 2007, 143A, 1268-1281.	1.2	43
107	Microarrays in prenatal diagnosis. <i>Best Practice and Research in Clinical Obstetrics and Gynaecology</i> , 2017, 42, 53-63.	2.8	43
108	Characterization of SETD1A haploinsufficiency in humans and <i>Drosophila</i> defines a novel neurodevelopmental syndrome. <i>Molecular Psychiatry</i> , 2021, 26, 2013-2024.	7.9	43

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109	Mesomelia-Synostoses Syndrome Results from Deletion of SULF1 and SLCO5A1 Genes at 8q13. <i>American Journal of Human Genetics</i> , 2010, 87, 95-100.	6.2	42
110	High-resolution chromosomal microarrays in prenatal diagnosis significantly increase diagnostic power. <i>Prenatal Diagnosis</i> , 2014, 34, 525-533.	2.3	42
111	Further corroboration of distinct functional features in SCN2A variants causing intellectual disability or epileptic phenotypes. <i>Molecular Medicine</i> , 2019, 25, 6.	4.4	42
112	Partial Loss of USP9X Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor β^2 Signaling. <i>Biological Psychiatry</i> , 2020, 87, 100-112.	1.3	42
113	Ichthyosis follicularis, alopecia, and photophobia (IFAP) syndrome: Clinical and neuropathological observations in a 33-year-old man. , 1998, 78, 371-377.		40
114	The shortest of the short: Pericentrin mutations and beyond. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2011, 25, 125-130.	4.7	40
115	Plasma metabolomics reveals a diagnostic metabolic fingerprint for mitochondrial aconitase (ACO2) deficiency. <i>PLoS ONE</i> , 2017, 12, e0176363.	2.5	40
116	Neocentric small supernumerary marker chromosomes (sSMC) â€“ three more cases and review of the literature. <i>Cytogenetic and Genome Research</i> , 2007, 118, 31-37.	1.1	37
117	Analysis of an interstitial deletion in a patient with Kallmann syndrome, X-linked ichthyosis and mental retardation. <i>Clinical Genetics</i> , 1998, 54, 45-51.	2.0	37
118	The smallest teeth in the world are caused by mutations in the <i>PCNT</i> gene. <i>American Journal of Medical Genetics, Part A</i> , 2011, 155, 1398-1403.	1.2	37
119	Variants in <i>CUL4B</i> are Associated with Cerebral Malformations. <i>Human Mutation</i> , 2015, 36, 106-117.	2.5	37
120	Mowat-Wilson syndrome and mutation in the zinc finger homeo box 1B gene: a well defined clinical entity. <i>Journal of Medical Genetics</i> , 2004, 41, 16e-16.	3.2	36
121	Molecular Diversity and Associated Phenotypic Spectrum of Germline <i>CBL</i> Mutations. <i>Human Mutation</i> , 2015, 36, 787-796.	2.5	36
122	Autosomal-Dominant Hypertension With Type E Brachydactyly Is Caused by Rearrangement on the Short Arm of Chromosome 12. <i>Hypertension</i> , 2004, 43, 471-476.	2.7	35
123	Pulmonary hypoplasiaâ€“diaphragmatic herniaâ€“anophthalmiaâ€“cardiac defect (PDAC) syndrome due to <i>STRA6</i> mutationsâ€“What are the minimal criteria?. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 2457-2463.	1.2	35
124	Goltzâ€“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, 1207-1215.	2.8	35
125	<i>STAG1</i> mutations cause a novel cohesinopathy characterised by unspecific syndromic intellectual disability. <i>Journal of Medical Genetics</i> , 2017, 54, 479-488.	3.2	35
126	A recurrent germline mutation in the <i>PIGA</i> gene causes Simpsonâ€“Golabiâ€“Behmel syndrome type 2. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 392-402.	1.2	34

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127	Genotype-phenotype evaluation of MED13L defects in the light of a novel truncating and a recurrent missense mutation. <i>European Journal of Medical Genetics</i> , 2017, 60, 451-464.	1.3	34
128	De novo truncating variants in the intronless IRF2BPL are responsible for developmental epileptic encephalopathy. <i>Genetics in Medicine</i> , 2019, 21, 1008-1014.	2.4	34
129	Disruption of RFX family transcription factors causes autism, attention-deficit/hyperactivity disorder, intellectual disability, and dysregulated behavior. <i>Genetics in Medicine</i> , 2021, 23, 1028-1040.	2.4	34
130	N ⁸ -acetylspermidine as a potential plasma biomarker for Snyder-Robinson syndrome identified by clinical metabolomics. <i>Journal of Inherited Metabolic Disease</i> , 2016, 39, 131-137.	3.6	33
131	A novel 5q35.3 subtelomeric deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2003, 121A, 1-8.	1.2	32
132	Atypical ZFX1B mutation associated with a mild Mowat-Wilson syndrome phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 869-872.	1.2	32
133	Unmasking of a Recessive SCARF2 Mutation by a 22q11.12 de novo Deletion in a Patient with Van den Ende-Gupta Syndrome. <i>Molecular Syndromology</i> , 2010, 1, 239-245.	0.8	32
134	New insights into the clinical and molecular spectrum of the novel CYFIP2-related neurodevelopmental disorder and impairment of the WRC-mediated actin dynamics. <i>Genetics in Medicine</i> , 2021, 23, 543-554.	2.4	32
135	"Mowat-Wilson" syndrome with and without Hirschsprung disease is a distinct, recognizable multiple congenital anomalies-mental retardation syndrome caused by mutations in the zinc finger homeo box 1B gene. <i>American Journal of Medical Genetics Part A</i> , 2002, 108, 177-81.	2.4	32
136	A missense mutation in the ZFX1B gene associated with an atypical Mowat-Wilson syndrome phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 1223-1227.	1.2	31
137	Novel missense, insertion and deletion mutations in the neurotrophic tyrosine kinase receptor type 1 gene (NTRK1) associated with congenital insensitivity to pain with anhidrosis. <i>Neuromuscular Disorders</i> , 2008, 18, 159-166.	0.6	31
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