## Anita Rauch

List of Publications by Year in descending order

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256 papers 17,020 citations

65 h-index 19749 117 g-index

270 all docs

270 docs citations

270 times ranked

23224 citing authors

#	Article	IF	CITATIONS
1	Human COQ4 deficiency: delineating the clinical, metabolic and neuroimaging phenotypes. Journal of Medical Genetics, 2022, 59, 878-887.	3.2	9
2	Response to Cueto-González etÂal. Genetics in Medicine, 2022, 24, 757.	2.4	0
3	The <i>MAP3K7</i> gene: Further delineation of clinical characteristics and genotype/phenotype correlations. Human Mutation, 2022, 43, 1377-1395.	2.5	5
4	Characterization of SETD1A haploinsufficiency in humans and Drosophila defines a novel neurodevelopmental syndrome. Molecular Psychiatry, 2021, 26, 2013-2024.	7.9	43
5	New insights into the clinical and molecular spectrum of the novel CYFIP2-related neurodevelopmental disorder and impairment of the WRC-mediated actin dynamics. Genetics in Medicine, 2021, 23, 543-554.	2.4	32
6	The broad phenotypic spectrum of PPP2R1A-related neurodevelopmental disorders correlates with the degree of biochemical dysfunction. Genetics in Medicine, 2021, 23, 352-362.	2.4	23
7	DLG4-related synaptopathy: a new rare brain disorder. Genetics in Medicine, 2021, 23, 888-899.	2.4	16
8	<scp><i>MED27</i></scp> Variants Cause Developmental Delay, Dystonia, and Cerebellar Hypoplasia. Annals of Neurology, 2021, 89, 828-833.	<b>5.</b> 3	14
9	SPEN haploinsufficiency causes a neurodevelopmental disorder overlapping proximal 1p36 deletion syndrome with an episignature of X chromosomes in females. American Journal of Human Genetics, 2021, 108, 502-516.	6.2	48
10	Disruption of RFX family transcription factors causes autism, attention-deficit/hyperactivity disorder, intellectual disability, and dysregulated behavior. Genetics in Medicine, 2021, 23, 1028-1040.	2.4	34
11	Missense and truncating variants in CHD5 in a dominant neurodevelopmental disorder with intellectual disability, behavioral disturbances, and epilepsy. Human Genetics, 2021, 140, 1109-1120.	3.8	18
12	Loss-of-function and missense variants in NSD2 cause decreased methylation activity and are associated with a distinct developmental phenotype. Genetics in Medicine, 2021, 23, 1474-1483.	2.4	24
13	<i>CSNK2B</i> : A broad spectrum of neurodevelopmental disability and epilepsy severity. Epilepsia, 2021, 62, e103-e109.	5.1	13
14	Rare variants in KDR, encoding VEGF Receptor 2, are associated with tetralogy of Fallot. Genetics in Medicine, 2021, 23, 1952-1960.	2.4	7
15	Confirmation of Ogden syndrome as an Xâ€linked recessive fatal disorder due to a recurrent <scp>NAA10</scp> variant and review of the literature. American Journal of Medical Genetics, Part A, 2021, 185, 2546-2560.	1.2	12
16	Expanding the phenotype: Four new cases and hope for treatment in <scp>Bachmannâ€Bupp</scp> syndrome. American Journal of Medical Genetics, Part A, 2021, 185, 3485-3493.	1.2	10
17	Generation and characterization of an endogenously tagged SPG11-human iPSC line by CRISPR/Cas9 mediated knock-in. Stem Cell Research, 2021, 56, 102520.	0.7	1
18	Novel Biallelic Variants in KIF21A Cause a Novel Phenotype of Fetal Akinesia with Neurodevelopmental Defects., 2021, 52, .		0

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19	Fetal tuberous sclerosis and diagnosis of paternal gonadal mosaicism. Ultrasound in Obstetrics and Gynecology, 2020, 55, 691-692.	1.7	2
20	A clinical scoring system for congenital contractural arachnodactyly. Genetics in Medicine, 2020, 22, 124-131.	2.4	17
21	Partial Loss of USP9X Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor $\hat{l}^2$ Signaling. Biological Psychiatry, 2020, 87, 100-112.	1.3	42
22	High-resolution chromosomal microarrayÂanalysis for copy-number variations in high-functioning autism reveals large aberration typical for intellectual disability. Journal of Neural Transmission, 2020, 127, 81-94.	2.8	5
23	Rare copy number variants in individuals at clinical high risk for psychosis: Enrichment of synaptic/brainâ€related functional pathways. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 140-151.	1.7	0
24	Severe reaction to radiotherapy provoked by hypomorphic germline mutations in ⟨i⟩ATM⟨/i⟩ (ataxia–telangiectasia mutated gene). Molecular Genetics & Enomic Medicine, 2020, 8, e1409.	1.2	8
25	Novel morphological and genetic features of fumarate hydratase deficient renal cell carcinoma in <scp>HLRCC</scp> syndrome patients with a tailored therapeutic approach. Genes Chromosomes and Cancer, 2020, 59, 611-619.	2.8	19
26	Genome-wide non-invasive prenatal testing in single- and multiple-pregnancies at any risk: Identification of maternal polymorphisms to reduce the number of unnecessary invasive confirmation testing. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2020, 252, 19-29.	1.1	7
27	<i>CDK5RAP2</i> primary microcephaly is associated with hypothalamic, retinal and cochlear developmental defects. Journal of Medical Genetics, 2020, 57, 389-399.	3.2	17
28	Insight Into the Ontogeny of GnRH Neurons From Patients Born Without a Nose. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 1538-1551.	3.6	7
29	Bi-allelic Pathogenic Variants in HS2ST1 Cause a Syndrome Characterized by Developmental Delay and Corpus Callosum, Skeletal, and Renal Abnormalities. American Journal of Human Genetics, 2020, 107, 1044-1061.	6.2	11
30	Swiss newborn screening for severe T and B cell deficiency with a combined TREC/KREC assay – management recommendations. Swiss Medical Weekly, 2020, 150, w20254.	1.6	17
31	Prenatal diagnosis of <i>HNF1B</i> >â€associated renal cysts: Is there a need to differentiate intragenic variants from 17q12 microdeletion syndrome?. Prenatal Diagnosis, 2019, 39, 1136-1147.	2.3	16
32	De Novo Pathogenic Variants in N-cadherin Cause a Syndromic Neurodevelopmental Disorder with Corpus Callosum, Axon, Cardiac, Ocular, and Genital Defects. American Journal of Human Genetics, 2019, 105, 854-868.	6.2	29
33	Spatially clustering de novo variants in CYFIP2, encoding the cytoplasmic FMRP interacting protein 2, cause intellectual disability and seizures. European Journal of Human Genetics, 2019, 27, 747-759.	2.8	47
34	CUGC for Simpson-Golabi-Behmel syndrome (SGBS). European Journal of Human Genetics, 2019, 27, 663-668.	2.8	9
35	Heterozygous Variants in KMT2E Cause a Spectrum of Neurodevelopmental Disorders and Epilepsy. American Journal of Human Genetics, 2019, 104, 1210-1222.	6.2	56
36	Elucidation of the phenotypic spectrum and genetic landscape in primary and secondary microcephaly. Genetics in Medicine, 2019, 21, 2043-2058.	2.4	57

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37	Further corroboration of distinct functional features in SCN2A variants causing intellectual disability or epileptic phenotypes. Molecular Medicine, 2019, 25, 6.	4.4	42
38	Deleterious Variation in BRSK2 Associates with a Neurodevelopmental Disorder. American Journal of Human Genetics, 2019, 104, 701-708.	6.2	19
39	Evolutionary conserved networks of human height identify multiple Mendelian causes of short stature. European Journal of Human Genetics, 2019, 27, 1061-1071.	2.8	11
40	De novo truncating variants in the intronless IRF2BPL are responsible for developmental epileptic encephalopathy. Genetics in Medicine, 2019, 21, 1008-1014.	2.4	34
41	The role of recessive inheritance in early-onset epileptic encephalopathies: a combined whole-exome sequencing and copy number study. European Journal of Human Genetics, 2019, 27, 408-421.	2.8	52
42	Prevalence of genetic susceptibility for breast and ovarian cancer in a non-cancer related study population: secondary germline findings from a Swiss single centre cohort. Swiss Medical Weekly, 2019, 149, w20092.	1.6	1
43	Surprisingly good outcome in antenatal diagnosis of severe hydrocephalus related to CCDC88C deficiency. European Journal of Medical Genetics, 2018, 61, 189-196.	1.3	9
44	Phenotype and genotype of 87 patients with Mowat–Wilson syndrome and recommendations for care. Genetics in Medicine, 2018, 20, 965-975.	2.4	67
45	Clinical and experimental evidence suggest a link between KIF7 and C5orf42-related ciliopathies through Sonic Hedgehog signaling. European Journal of Human Genetics, 2018, 26, 197-209.	2.8	23
46	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. Genetics in Medicine, 2018, 20, 630-638.	2.4	101
47	Novel <i>STRA6</i> null mutations in the original family described with Matthew–Wood syndrome. American Journal of Medical Genetics, Part A, 2018, 176, 134-138.	1.2	10
48	Autosomal recessive primary microcephaly due to <i>ASPM</i> mutations: An update. Human Mutation, 2018, 39, 319-332.	2.5	53
49	Need for high-resolution Genetic Analysis in iPSC: Results and Lessons from the ForIPS Consortium. Scientific Reports, 2018, 8, 17201.	3.3	70
50	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.	6.2	18
51	De Novo Variants in the F-Box Protein FBXO11 in 20 Individuals with a Variable Neurodevelopmental Disorder. American Journal of Human Genetics, 2018, 103, 305-316.	6.2	48
52	Clinical and functional characterization of two novel <i>ZBTB20</i> mutations causing Primrose syndrome. Human Mutation, 2018, 39, 959-964.	2.5	11
53	Biallelic UFM1 and UFC1 mutations expand the essential role of ufmylation in brain development. Brain, 2018, 141, 1934-1945.	7.6	70
54	SMCHD1 mutations associated with a rare muscular dystrophy can also cause isolated arhinia and Bosma arhinia microphthalmia syndrome. Nature Genetics, 2017, 49, 238-248.	21.4	131

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55	Microarrays in prenatal diagnosis. Best Practice and Research in Clinical Obstetrics and Gynaecology, 2017, 42, 53-63.	2.8	43
56	<i>FOXP2</i> variants in 14 individuals with developmental speech and language disorders broaden the mutational and clinical spectrum. Journal of Medical Genetics, 2017, 54, 64-72.	3.2	67
57	<i>STAG1</i> mutations cause a novel cohesinopathy characterised by unspecific syndromic intellectual disability. Journal of Medical Genetics, 2017, 54, 479-488.	3.2	35
58	Confirmation of mutations in <i>PROSC</i> sas a novel cause of vitamin B <sub><sub><sub></sub></sub></sub> -dependent epilepsy. Journal of Medical Genetics, 2017, 54, 809-814.	3.2	66
59	YY1 Haploinsufficiency Causes an Intellectual Disability Syndrome Featuring Transcriptional and Chromatin Dysfunction. American Journal of Human Genetics, 2017, 100, 907-925.	6.2	125
60	The HHID syndrome of hypertrichosis, hyperkeratosis, abnormal corpus callosum, intellectual disability, and minor anomalies is caused by mutations in <i>ARID1B</i> . American Journal of Medical Genetics, Part A, 2017, 173, 1440-1443.	1.2	11
61	Low-Level Chromosomal Mosaicism in Neurodevelopmental Disorders. Molecular Syndromology, 2017, 8, 266-271.	0.8	5
62	Monoallelic BMP2 Variants Predicted to Result in Haploinsufficiency Cause Craniofacial, Skeletal, and Cardiac Features Overlapping Those of 20p12 Deletions. American Journal of Human Genetics, 2017, 101, 985-994.	6.2	44
63	Genotype-phenotype evaluation of MED13L defects in the light of a novel truncating and a recurrent missense mutation. European Journal of Medical Genetics, 2017, 60, 451-464.	1.3	34
64	High resolution chromosomal microarray analysis in paediatric obsessive-compulsive disorder. BMC Medical Genomics, 2017, 10, 68.	1.5	21
65	MCAD-Deficiency with Severe Neonatal Onset, Fatal Outcome and Normal Acylcarnitine Profile. International Journal of Neonatal Screening, 2017, 3, 21.	3.2	0
66	Plasma metabolomics reveals a diagnostic metabolic fingerprint for mitochondrial aconitase (ACO2) deficiency. PLoS ONE, 2017, 12, e0176363.	2.5	40
67	Genetic and neurodevelopmental spectrum of <i>SYNGAP1 </i> epilepsy. Journal of Medical Genetics, 2016, 53, 511-522.	3.2	135
68	Mutations in CRADD Result in Reduced Caspase-2-Mediated Neuronal Apoptosis and Cause Megalencephaly with a Rare Lissencephaly Variant. American Journal of Human Genetics, 2016, 99, 1117-1129.	6.2	50
69	A recurrent germline mutation in the <i>PIGA</i> gene causes Simpsonâ€Golabiâ€Behmel syndrome type 2. American Journal of Medical Genetics, Part A, 2016, 170, 392-402.	1.2	34
70	The value of plasma vitamin B <sub>6</sub> profiles in early onset epileptic encephalopathies. Journal of Inherited Metabolic Disease, 2016, 39, 733-741.	3.6	19
71	Gene discovery for Mendelian conditions via social networking: de novo variants in KDM1A cause developmental delay and distinctive facial features. Genetics in Medicine, 2016, 18, 788-795.	2.4	88
72	Noninvasive prenatal testing: more caution in counseling is needed in high risk pregnancies with ultrasound abnormalities. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2016, 200, 72-75.	1.1	15

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73	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. JAMA Psychiatry, 2016, 73, 20.	11.0	195
74	N <sup>8</sup> â€acetylspermidine as a potential plasma biomarker for Snyderâ€Robinson syndrome identified by clinical metabolomics. Journal of Inherited Metabolic Disease, 2016, 39, 131-137.	3.6	33
75	Mutations in <i> <scp>CDK</scp> 5 <scp>RAP</scp> 2 </i> cause Seckel syndrome. Molecular Genetics & Genomic Medicine, 2015, 3, 467-480.	1.2	55
76	A severe congenital myasthenic syndrome with "dropped head―caused by novel <i>MUSK</i> mutations. Muscle and Nerve, 2015, 52, 668-673.	2.2	21
77	Intragenic <i>KANSL1</i> mutations and chromosome 17q21.31 deletions: broadening the clinical spectrum and genotype–phenotype correlations in a large cohort of patients. Journal of Medical Genetics, 2015, 52, 804-814.	3.2	47
78	Variants in <i>CUL4B</i> are Associated with Cerebral Malformations. Human Mutation, 2015, 36, 106-117.	2.5	37
79	De novo missense mutations in the NAA10 gene cause severe non-syndromic developmental delay in males and females. European Journal of Human Genetics, 2015, 23, 602-609.	2.8	72
80	PDE3A mutations cause autosomal dominant hypertension with brachydactyly. Nature Genetics, 2015, 47, 647-653.	21.4	146
81	Molecular Diversity and Associated Phenotypic Spectrum of Germline <i>CBL</i> Mutations. Human Mutation, 2015, 36, 787-796.	2.5	36
82	A Potential Contributory Role for Ciliary Dysfunction in the 16p11.2 600 kb BP4-BP5 Pathology. American Journal of Human Genetics, 2015, 96, 784-796.	6.2	53
83	Mutations in XRCC4 cause primary microcephaly, short stature and increased genomic instability. Human Molecular Genetics, 2015, 24, 3708-17.	2.9	26
84	Hydrops, fetal pleural effusions and chylothorax in three patients with <i>CBL</i> mutations. American Journal of Medical Genetics, Part A, 2015, 167, 394-399.	1.2	24
85	Infantile Epileptic Encephalopathy, Transient Choreoathetotic Movements, and Hypersomnia due to a De Novo Missense Mutation in the SCN2A Gene. Neuropediatrics, 2014, 45, 261-264.	0.6	30
86	The clinical significance of small copy number variants in neurodevelopmental disorders. Journal of Medical Genetics, 2014, 51, 677-688.	3.2	72
87	Dysmorphology at a distance: results of a web-based diagnostic service. European Journal of Human Genetics, 2014, 22, 327-332.	2.8	14
88	Exome sequencing in unspecific intellectual disability and rare disorders. Molecular Cytogenetics, 2014, 7, 126.	0.9	1
89	A newly recognized 13q12.3 microdeletion syndrome characterized by intellectual disability, microcephaly, and eczema/atopic dermatitis encompassing the <i>HMGB1</i> and <i>KATNAL1</i> genes. American Journal of Medical Genetics, Part A, 2014, 164, 1277-1283.	1.2	25
90	Microcephalic osteodysplastic primordial dwarfism type II (MOPD II) with multiple vascular complications misdiagnosed as Dubowitz syndrome. European Journal of Pediatrics, 2014, 173, 1253-1256.	2.7	17

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91	Mutations Affecting the SAND Domain of DEAF1 Cause Intellectual Disability with Severe Speech Impairment and Behavioral Problems. American Journal of Human Genetics, 2014, 94, 649-661.	6.2	59
92	<i>LETM1</i> haploinsufficiency causes mitochondrial defects in Wolf-Hirschhorn syndrome patient cells: implications for dissecting underlying pathomechanisms in this condition. DMM Disease Models and Mechanisms, 2014, 7, 535-45.	2.4	26
93	Highâ€resolution chromosomal microarrays in prenatal diagnosis significantly increase diagnostic power. Prenatal Diagnosis, 2014, 34, 525-533.	2.3	42
94	Nephrocalcinosis (Enamel Renal Syndrome) Caused by Autosomal Recessive FAM20A Mutations. Nephron Physiology, 2013, 122, 1-6.	1.2	84
95	Further delineation of genotype–phenotype correlation in homozygous 2p21 deletion syndromes: First description of patients without cystinuria. American Journal of Medical Genetics, Part A, 2013, 161, 1853-1859.	1.2	22
96	Platelet defects in congenital variant of Rett syndrome patients with FOXG1 mutations or reduced expression due to a position effect at 14q12. European Journal of Human Genetics, 2013, 21, 1349-1355.	2.8	13
97	Somatic mosaicism in a mother of two children with Pitt–Hopkins syndrome. Clinical Genetics, 2013, 83, 73-77.	2.0	31
98	Dosage changes of MED13L further delineate its role in congenital heart defects and intellectual disability. European Journal of Human Genetics, 2013, 21, 1100-1104.	2.8	57
99	De Novo Mutations in the Genome Organizer CTCF Cause Intellectual Disability. American Journal of Human Genetics, 2013, 93, 124-131.	6.2	151
100	A comprehensive molecular study on Coffin–Siris and Nicolaides–Baraitser syndromes identifies a broad molecular and clinical spectrum converging on altered chromatin remodeling. Human Molecular Genetics, 2013, 22, 5121-5135.	2.9	190
101	A new face of Borjeson–Forssman–Lehmann syndrome? De novo mutations in <i>PHF6</i> in seven females with a distinct phenotype. Journal of Medical Genetics, 2013, 50, 838-847.	3.2	50
102	Rare Copy Number Variants Are a Common Cause of Short Stature. PLoS Genetics, 2013, 9, e1003365.	3.5	60
103	Association Between C677T Polymorphism of Methylene Tetrahydrofolate Reductase and Congenital Heart Disease. Circulation: Cardiovascular Genetics, 2013, 6, 347-353.	5.1	31
104	Biallelic <i>SEMA3A</i> defects cause a novel type of syndromic short stature. American Journal of Medical Genetics, Part A, 2013, 161, 2880-2889.	1.2	9
105	Genome-wide association study identifies loci on 12q24 and 13q32 associated with Tetralogy of Fallot. Human Molecular Genetics, 2013, 22, 1473-1481.	2.9	82
106	Phenotype-specific effect of chromosome 1q21.1 rearrangements and GJA5 duplications in 2436 congenital heart disease patients and 6760 controls. Human Molecular Genetics, 2012, 21, 1513-1520.	2.9	101
107	Range of genetic mutations associated with severe non-syndromic sporadic intellectual disability: an exome sequencing study. Lancet, The, 2012, 380, 1674-1682.	13.7	940
108	Contribution of Global Rare Copy-Number Variants to the Risk of Sporadic Congenital Heart Disease. American Journal of Human Genetics, 2012, 91, 489-501.	6.2	272

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109	Macrocerebellum: Significance and Pathogenic Considerations. Cerebellum, 2012, 11, 1026-1036.	2.5	25
110	Novel <i>KIF7 &lt; /i&gt; mutations extend the phenotypic spectrum of acrocallosal syndrome. Journal of Medical Genetics, 2012, 49, 713-720.</i>	3.2	28
111	Wachstumsstörungen als Leitsymptom. Medizinische Genetik, 2012, 24, 123-137.	0.2	0
112	Haploinsufficiency of ARID1B, a Member of the SWI/SNF-A Chromatin-Remodeling Complex, Is a Frequent Cause of Intellectual Disability. American Journal of Human Genetics, 2012, 90, 565-572.	6.2	225
113	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. Nature, 2011, 478, 97-102.	27.8	394
114	Microdeletions of chromosome 7p21, including TWIST1, associated with significant microcephaly, facial dysmorphism, and short stature. European Journal of Medical Genetics, 2011, 54, 256-261.	1.3	9
115	7ÂMb de novo deletion within 8q21 in a patient with distal arthrogryposis type 2B (DA2B). European Journal of Medical Genetics, 2011, 54, e495-e500.	1.3	10
116	Familial short stature due to a 5q22.1–q23.2 duplication refines the 5q duplication spectrum. European Journal of Medical Genetics, 2011, 54, e521-e524.	1.3	5
117	The molecular basis of the cartilage-hair hypoplasia–anauxetic dysplasia spectrum. Best Practice and Research in Clinical Endocrinology and Metabolism, 2011, 25, 131-142.	4.7	46
118	The shortest of the short: Pericentrin mutations and beyond. Best Practice and Research in Clinical Endocrinology and Metabolism, 2011, 25, 125-130.	4.7	40
119	De novo nonsense mutations in ASXL1 cause Bohring-Opitz syndrome. Nature Genetics, 2011, 43, 729-731.	21.4	236
120	Genome-wide copy number profiling using a 100K SNP array reveals novel disease-related genes BORIS and TSHZ1 in juvenile angiofibroma. International Journal of Oncology, 2011, 39, 1143-51.	3.3	12
121	CEP152 is a genome maintenance protein disrupted in Seckel syndrome. Nature Genetics, 2011, 43, 23-26.	21.4	201
122	Clinical utility gene card for: Mowat–Wilson syndrome. European Journal of Human Genetics, 2011, 19, 4-4.	2.8	5
123	The MEF2C-Related and 5q14.3q15 Microdeletion Syndrome. Molecular Syndromology, 2011, 2, 164-170.	0.8	51
124	The core FOXG1 syndrome phenotype consists of postnatal microcephaly, severe mental retardation, absent language, dyskinesia, and corpus callosum hypogenesis. Journal of Medical Genetics, 2011, 48, 396-406.	3.2	220
125	NEK1 Mutations Cause Short-Rib Polydactyly Syndrome Type Majewski. American Journal of Human Genetics, 2011, 88, 106-114.	6.2	151
126	Expanding the clinical spectrum associated with defects in CNTNAP2 and NRXN1. BMC Medical Genetics, 2011, 12, 106.	2.1	109

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127	The smallest teeth in the world are caused by mutations in the <i>PCNT</i> gene. American Journal of Medical Genetics, Part A, 2011, 155, 1398-1403.	1.2	37
128	Identification of a novel loss-of-function calcium channel gene mutation in short QT syndrome (SQTS6). European Heart Journal, 2011, 32, 1077-1088.	2.2	178
129	In-Frame Deletion and Missense Mutations of the C-Terminal Helicase Domain of <b><i>SMARCA2</i></b> in Three Patients with Nicolaides-Baraitser Syndrome. Molecular Syndromology, 2011, 2, 237-244.	0.8	58
130	Disruption of the histone acetyltransferase MYST4 leads to a Noonan syndrome–like phenotype and hyperactivated MAPK signaling in humans and mice. Journal of Clinical Investigation, 2011, 121, 3479-3491.	8.2	89
131	Disturbed Wnt Signalling due to a Mutation in CCDC88C Causes an Autosomal Recessive Non-Syndromic Hydrocephalus with Medial Diverticulum. Molecular Syndromology, 2010, 1, 99-112.	0.8	82
132	Mesomelia-Synostoses Syndrome Results from Deletion of SULF1 and SLCO5A1 Genes at 8q13. American Journal of Human Genetics, 2010, 87, 95-100.	6.2	42
133	Severe clinical course of Hirschsprung disease in a Mowat-Wilson syndrome patient. Journal of Applied Genetics, 2010, 51, 111-113.	1.9	14
134	Mutations in MEF2C from the 5q14.3q15 microdeletion syndrome region are a frequent cause of severe mental retardation and diminish MECP2 and CDKL5 expression. Human Mutation, 2010, 31, 722-733.	2.5	163
135	The face of Noonan syndrome: Does phenotype predict genotype. American Journal of Medical Genetics, Part A, 2010, 152A, 1960-1966.	1.2	59
136	Altered $TGF\hat{l}^2$ signaling and cardiovascular manifestations in patients with autosomal recessive cutis laxa type I caused by fibulin-4 deficiency. European Journal of Human Genetics, 2010, 18, 895-901.	2.8	132
137	Clinical utility gene card for: DiGeorge syndrome, velocardiofacial syndrome, Shprintzen syndrome, chromosome 22q11.2 deletion syndrome (22q11.2, TBX1). European Journal of Human Genetics, 2010, 18, 1071-1071.	2.8	6
138	Mutations in GRIN2A and GRIN2B encoding regulatory subunits of NMDA receptors cause variable neurodevelopmental phenotypes. Nature Genetics, 2010, 42, 1021-1026.	21.4	431
139	Disruption of ST5 is associated with mental retardation and multiple congenital anomalies. Journal of Medical Genetics, 2010, 47, 91-98.	3.2	12
140	Unmasking of a Recessive SCARF2 Mutation by a 22q11.12 de novo Deletion in a Patient with Van den Ende-Gupta Syndrome. Molecular Syndromology, 2010, 1, 239-245.	0.8	32
141	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. Heart, 2010, 96, 1651-1655.	2.9	61
142	Comprehensive genotype-phenotype analysis in 230 patients with tetralogy of Fallot. Journal of Medical Genetics, 2010, 47, 321-331.	3.2	126
143	Microcephalin and pericentrin regulate mitotic entry via centrosome-associated Chk1. Journal of Cell Biology, 2009, 185, 1149-1157.	5.2	83
144	Independent <i>NF1</i> and <i>PTPN11</i> mutations in a family with neurofibromatosisâ€Noonan syndrome. American Journal of Medical Genetics, Part A, 2009, 149A, 1263-1267.	1.2	55

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145	Pulmonary hypoplasia–diaphragmatic hernia–anophthalmia–cardiac defect (PDAC) syndrome due to <i>&gt;STRA6</i> mutations—What are the minimal criteria?. American Journal of Medical Genetics, Part A, 2009, 149A, 2457-2463.	1.2	35
146	Chromosomale Ursachen der geistigen Behinderung. Medizinische Genetik, 2009, 21, 237-245.	0.2	0
147	Goltz–Gorlin (focal dermal hypoplasia) and the microphthalmia with linear skin defects (MLS) syndrome: no evidence of genetic overlap. European Journal of Human Genetics, 2009, 17, 1207-1215.	2.8	35
148	A novel microdeletion syndrome involving 5q14.3-q15: clinical and molecular cytogenetic characterization of three patients. European Journal of Human Genetics, 2009, 17, 1592-1599.	2.8	96
149	Lethal cutis laxa with contractural arachnodactyly, overgrowth and soft tissue bleeding due to a novel homozygous <i>fibulinâ€4</i> gene mutation. Clinical Genetics, 2009, 76, 276-281.	2.0	50
150	CNTNAP2 and NRXN1 Are Mutated in Autosomal-Recessive Pitt-Hopkins-like Mental Retardation and Determine the Level of a Common Synaptic Protein in Drosophila. American Journal of Human Genetics, 2009, 85, 655-666.	6.2	573
151	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. American Journal of Human Genetics, 2009, 85, 809-822.	6.2	70
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