

Anita Rauch

List of Publications by Year in descending order

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255
papers

17,020
citations

17776

65
h-index

22488

117
g-index

270
all docs

270
docs citations

270
times ranked

25224
citing authors

#	ARTICLE	IF	CITATIONS
1	Human COQ4 deficiency: delineating the clinical, metabolic and neuroimaging phenotypes. Journal of Medical Genetics, 2022, 59, 878-887.	1.5	9
2	Response to Cueto-González et al. Genetics in Medicine, 2022, 24, 757.	1.1	0
3	The <i>MAP3K7</i> gene: Further delineation of clinical characteristics and genotype/phenotype correlations. Human Mutation, 2022, 43, 1377-1395.	1.1	5
4	Characterization of SETD1A haploinsufficiency in humans and Drosophila defines a novel neurodevelopmental syndrome. Molecular Psychiatry, 2021, 26, 2013-2024.	4.1	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.	1.1	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.	1.1	23
7	DLG4-related synaptopathy: a new rare brain disorder. Genetics in Medicine, 2021, 23, 888-899.	1.1	16
8	<i>MED27</i> Variants Cause Developmental Delay, Dystonia, and Cerebellar Hypoplasia. Annals of Neurology, 2021, 89, 828-833.	2.8	14
9	SPEN haploinsufficiency causes a neurodevelopmental disorder overlapping proximal 1p36 deletion syndrome with an epistatue of X chromosomes in females. American Journal of Human Genetics, 2021, 108, 502-516.	2.6	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.	1.1	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.	1.8	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.	1.1	24
13	<i>CSNK2B</i> : A broad spectrum of neurodevelopmental disability and epilepsy severity. Epilepsia, 2021, 62, e103-e109.	2.6	13
14	Rare variants in KDR, encoding VEGF Receptor 2, are associated with tetralogy of Fallot. Genetics in Medicine, 2021, 23, 1952-1960.	1.1	7
15	Confirmation of Ogden syndrome as an X-linked recessive fatal disorder due to a recurrent <i>NAA10</i> variant and review of the literature. American Journal of Medical Genetics, Part A, 2021, 185, 2546-2560.	0.7	12
16	Expanding the phenotype: Four new cases and hope for treatment in <i>Bachmann-Bupp</i> syndrome. American Journal of Medical Genetics, Part A, 2021, 185, 3485-3493.	0.7	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.	0.3	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. <i>Ultrasound in Obstetrics and Gynecology</i> , 2020, 55, 691-692.	0.9	2
20	A clinical scoring system for congenital contractural arachnodactyly. <i>Genetics in Medicine</i> , 2020, 22, 124-131.	1.1	17
21	Partial Loss of USP9X Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor β Signaling. <i>Biological Psychiatry</i> , 2020, 87, 100-112.	0.7	42
22	High-resolution chromosomal microarray analysis for copy-number variations in high-functioning autism reveals large aberration typical for intellectual disability. <i>Journal of Neural Transmission</i> , 2020, 127, 81-94.	1.4	5
23	Rare copy number variants in individuals at clinical high risk for psychosis: Enrichment of synaptic/brain-related functional pathways. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2020, 183, 140-151.	1.1	0
24	Severe reaction to radiotherapy provoked by hypomorphic germline mutations in <i>ATM</i> (ataxia-telangiectasia mutated gene). <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1409.	0.6	8
25	Novel morphological and genetic features of fumarate hydratase deficient renal cell carcinoma in <i>HLRCC</i> syndrome patients with a tailored therapeutic approach. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 611-619.	1.5	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. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2020, 252, 19-29.	0.5	7
27	<i>CDK5RAP2</i> primary microcephaly is associated with hypothalamic, retinal and cochlear developmental defects. <i>Journal of Medical Genetics</i> , 2020, 57, 389-399.	1.5	17
28	Insight Into the Ontogeny of GnRH Neurons From Patients Born Without a Nose. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 1538-1551.	1.8	7
29	Bi-allelic Pathogenic Variants in <i>HS2ST1</i> Cause a Syndrome Characterized by Developmental Delay and Corpus Callosum, Skeletal, and Renal Abnormalities. <i>American Journal of Human Genetics</i> , 2020, 107, 1044-1061.	2.6	11
30	Swiss newborn screening for severe T and B cell deficiency with a combined TREC/KREC assay management recommendations. <i>Swiss Medical Weekly</i> , 2020, 150, w20254.	0.8	17
31	Prenatal diagnosis of <i>HNF1B</i> -associated renal cysts: Is there a need to differentiate intragenic variants from 17q12 microdeletion syndrome?. <i>Prenatal Diagnosis</i> , 2019, 39, 1136-1147.	1.1	16
32	De Novo Pathogenic Variants in N-cadherin Cause a Syndromic Neurodevelopmental Disorder with Corpus Callosum, Axon, Cardiac, Ocular, and Genital Defects. <i>American Journal of Human Genetics</i> , 2019, 105, 854-868.	2.6	29
33	Spatially clustering de novo variants in <i>CYFIP2</i> , encoding the cytoplasmic FMRP interacting protein 2, cause intellectual disability and seizures. <i>European Journal of Human Genetics</i> , 2019, 27, 747-759.	1.4	47
34	CUGC for Simpson-Golabi-Behmel syndrome (SGBS). <i>European Journal of Human Genetics</i> , 2019, 27, 663-668.	1.4	9
35	Heterozygous Variants in <i>KMT2E</i> Cause a Spectrum of Neurodevelopmental Disorders and Epilepsy. <i>American Journal of Human Genetics</i> , 2019, 104, 1210-1222.	2.6	56
36	Elucidation of the phenotypic spectrum and genetic landscape in primary and secondary microcephaly. <i>Genetics in Medicine</i> , 2019, 21, 2043-2058.	1.1	57

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37	Further corroboration of distinct functional features in SCN2A variants causing intellectual disability or epileptic phenotypes. <i>Molecular Medicine</i> , 2019, 25, 6.	1.9	42
38	Deleterious Variation in BRSK2 Associates with a Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2019, 104, 701-708.	2.6	19
39	Evolutionary conserved networks of human height identify multiple Mendelian causes of short stature. <i>European Journal of Human Genetics</i> , 2019, 27, 1061-1071.	1.4	11
40	De novo truncating variants in the intronless IRF2BPL are responsible for developmental epileptic encephalopathy. <i>Genetics in Medicine</i> , 2019, 21, 1008-1014.	1.1	34
41	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.	1.4	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. <i>Swiss Medical Weekly</i> , 2019, 149, w20092.	0.8	1
43	Surprisingly good outcome in antenatal diagnosis of severe hydrocephalus related to CCDC88C deficiency. <i>European Journal of Medical Genetics</i> , 2018, 61, 189-196.	0.7	9
44	Phenotype and genotype of 87 patients with Mowatâ€™Wilson syndrome and recommendations for care. <i>Genetics in Medicine</i> , 2018, 20, 965-975.	1.1	67
45	Clinical and experimental evidence suggest a link between KIF7 and C5orf42-related ciliopathies through Sonic Hedgehog signaling. <i>European Journal of Human Genetics</i> , 2018, 26, 197-209.	1.4	23
46	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. <i>Genetics in Medicine</i> , 2018, 20, 630-638.	1.1	101
47	Novel <i>STRA6</i> null mutations in the original family described with Matthewâ€™Wood syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 134-138.	0.7	10
48	Autosomal recessive primary microcephaly due to <i>ASPM</i> mutations: An update. <i>Human Mutation</i> , 2018, 39, 319-332.	1.1	53
49	Need for high-resolution Genetic Analysis in iPSC: Results and Lessons from the ForIPS Consortium. <i>Scientific Reports</i> , 2018, 8, 17201.	1.6	70
50	Bi-allelic TMEM94 Truncating Variants Are Associated with Neurodevelopmental Delay, Congenital Heart Defects, and Distinct Facial Dysmorphism. <i>American Journal of Human Genetics</i> , 2018, 103, 948-967.	2.6	18
51	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.	2.6	48
52	Clinical and functional characterization of two novel <i>ZBTB20</i> mutations causing Primrose syndrome. <i>Human Mutation</i> , 2018, 39, 959-964.	1.1	11
53	Biallelic UFM1 and UFC1 mutations expand the essential role of ufmylation in brain development. <i>Brain</i> , 2018, 141, 1934-1945.	3.7	70
54	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.	9.4	131

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55	Microarrays in prenatal diagnosis. Best Practice and Research in Clinical Obstetrics and Gynaecology, 2017, 42, 53-63.	1.4	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.	1.5	67
57	<i>STAG1</i> mutations cause a novel cohesinopathy characterised by unspecific syndromic intellectual disability. Journal of Medical Genetics, 2017, 54, 479-488.	1.5	35
58	Confirmation of mutations in <i>PROSC</i> as a novel cause of vitamin B₆-dependent epilepsy. Journal of Medical Genetics, 2017, 54, 809-814.	1.5	66
59	YY1 Haploinsufficiency Causes an Intellectual Disability Syndrome Featuring Transcriptional and Chromatin Dysfunction. American Journal of Human Genetics, 2017, 100, 907-925.	2.6	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.	0.7	11
61	Low-Level Chromosomal Mosaicism in Neurodevelopmental Disorders. Molecular Syndromology, 2017, 8, 266-271.	0.3	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.	2.6	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.	0.7	34
64	High resolution chromosomal microarray analysis in paediatric obsessive-compulsive disorder. BMC Medical Genomics, 2017, 10, 68.	0.7	21
65	MCAD-Deficiency with Severe Neonatal Onset, Fatal Outcome and Normal Acylcarnitine Profile. International Journal of Neonatal Screening, 2017, 3, 21.	1.2	0
66	Plasma metabolomics reveals a diagnostic metabolic fingerprint for mitochondrial aconitase (ACO2) deficiency. PLoS ONE, 2017, 12, e0176363.	1.1	40
67	Genetic and neurodevelopmental spectrum of <i>SYNGAP1</i>-associated intellectual disability and epilepsy. Journal of Medical Genetics, 2016, 53, 511-522.	1.5	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.	2.6	50
69	A recurrent germline mutation in the <i>PIGA</i> gene causes Simpsonâ€™Colabiâ€™Behmel syndrome type 2. American Journal of Medical Genetics, Part A, 2016, 170, 392-402.	0.7	34
70	The value of plasma vitamin B₆ profiles in early onset epileptic encephalopathies. Journal of Inherited Metabolic Disease, 2016, 39, 733-741.	1.7	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.	1.1	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.	0.5	15

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73	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. <i>JAMA Psychiatry</i> , 2016, 73, 20.	6.0	195
74	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.	1.7	33
75	Mutations in <i>CDK5</i> & <i>RAP2</i> cause Seckel syndrome. <i>Molecular Genetics & Genomic Medicine</i> , 2015, 3, 467-480.	0.6	55
76	A severe congenital myasthenic syndrome with "dropped head" caused by novel <i>MUSK</i> mutations. <i>Muscle and Nerve</i> , 2015, 52, 668-673.	1.0	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. <i>Journal of Medical Genetics</i> , 2015, 52, 804-814.	1.5	47
78	Variants in <i>CUL4B</i> are Associated with Cerebral Malformations. <i>Human Mutation</i> , 2015, 36, 106-117.	1.1	37
79	De novo missense mutations in the <i>NAA10</i> gene cause severe non-syndromic developmental delay in males and females. <i>European Journal of Human Genetics</i> , 2015, 23, 602-609.	1.4	72
80	<i>PDE3A</i> mutations cause autosomal dominant hypertension with brachydactyly. <i>Nature Genetics</i> , 2015, 47, 647-653.	9.4	146
81	Molecular Diversity and Associated Phenotypic Spectrum of Germline <i>CBL</i> Mutations. <i>Human Mutation</i> , 2015, 36, 787-796.	1.1	36
82	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.	2.6	53
83	Mutations in <i>XRCC4</i> cause primary microcephaly, short stature and increased genomic instability. <i>Human Molecular Genetics</i> , 2015, 24, 3708-17.	1.4	26
84	Hydrops, fetal pleural effusions and chylothorax in three patients with <i>CBL</i> mutations. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 394-399.	0.7	24
85	Infantile Epileptic Encephalopathy, Transient Choreoathetotic Movements, and Hypersomnia due to a De Novo Missense Mutation in the <i>SCN2A</i> Gene. <i>Neuropediatrics</i> , 2014, 45, 261-264.	0.3	30
86	The clinical significance of small copy number variants in neurodevelopmental disorders. <i>Journal of Medical Genetics</i> , 2014, 51, 677-688.	1.5	72
87	Dysmorphology at a distance: results of a web-based diagnostic service. <i>European Journal of Human Genetics</i> , 2014, 22, 327-332.	1.4	14
88	Exome sequencing in unspecific intellectual disability and rare disorders. <i>Molecular Cytogenetics</i> , 2014, 7, 126.	0.4	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. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1277-1283.	0.7	25
90	Microcephalic osteodysplastic primordial dwarfism type II (MOPD II) with multiple vascular complications misdiagnosed as Dubowitz syndrome. <i>European Journal of Pediatrics</i> , 2014, 173, 1253-1256.	1.3	17

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91	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.	2.6	59
92	<i>LETM1</i> haploinsufficiency causes mitochondrial defects in Wolf-Hirschhorn syndrome patient cells: implications for dissecting underlying pathomechanisms in this condition. <i>DMM Disease Models and Mechanisms</i> , 2014, 7, 535-45.	1.2	26
93	High-resolution chromosomal microarrays in prenatal diagnosis significantly increase diagnostic power. <i>Prenatal Diagnosis</i> , 2014, 34, 525-533.	1.1	42
94	Nephrocalcinosis (Enamel Renal Syndrome) Caused by Autosomal Recessive FAM20A Mutations. <i>Nephron Physiology</i> , 2013, 122, 1-6.	1.5	84
95	Further delineation of genotype-phenotype correlation in homozygous 2p21 deletion syndromes: First description of patients without cystinuria. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1853-1859.	0.7	22
96	Platelet defects in congenital variant of Rett syndrome patients with FOXP1 mutations or reduced expression due to a position effect at 14q12. <i>European Journal of Human Genetics</i> , 2013, 21, 1349-1355.	1.4	13
97	Somatic mosaicism in a mother of two children with Pitt-Hopkins syndrome. <i>Clinical Genetics</i> , 2013, 83, 73-77.	1.0	31
98	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.	1.4	57
99	De Novo Mutations in the Genome Organizer CTCF Cause Intellectual Disability. <i>American Journal of Human Genetics</i> , 2013, 93, 124-131.	2.6	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. <i>Human Molecular Genetics</i> , 2013, 22, 5121-5135.	1.4	190
101	A new face of Borjeson-Forssman-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.	1.5	50
102	Rare Copy Number Variants Are a Common Cause of Short Stature. <i>PLoS Genetics</i> , 2013, 9, e1003365.	1.5	60
103	Association Between C677T Polymorphism of Methylene Tetrahydrofolate Reductase and Congenital Heart Disease. <i>Circulation: Cardiovascular Genetics</i> , 2013, 6, 347-353.	5.1	31
104	Biallelic <i>SEMA3A</i> defects cause a novel type of syndromic short stature. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2880-2889.	0.7	9
105	Genome-wide association study identifies loci on 12q24 and 13q32 associated with Tetralogy of Fallot. <i>Human Molecular Genetics</i> , 2013, 22, 1473-1481.	1.4	82
106	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.	1.4	101
107	Range of genetic mutations associated with severe non-syndromic sporadic intellectual disability: an exome sequencing study. <i>Lancet, The</i> , 2012, 380, 1674-1682.	6.3	940
108	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.	2.6	272

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109	Macrocerbellum: Significance and Pathogenic Considerations. <i>Cerebellum</i> , 2012, 11, 1026-1036.	1.4	25
110	Novel <i>KIF7</i> mutations extend the phenotypic spectrum of acrocallosal syndrome. <i>Journal of Medical Genetics</i> , 2012, 49, 713-720.	1.5	28
111	Wachstumsstörungen als Leitsymptom. <i>Medizinische Genetik</i> , 2012, 24, 123-137.	0.1	0
112	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.	2.6	225
113	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. <i>Nature</i> , 2011, 478, 97-102.	13.7	394
114	Microdeletions of chromosome 7p21, including TWIST1, associated with significant microcephaly, facial dysmorphism, and short stature. <i>European Journal of Medical Genetics</i> , 2011, 54, 256-261.	0.7	9
115	7 Mb de novo deletion within 8q21 in a patient with distal arthrogyriposis type 2B (DA2B). <i>European Journal of Medical Genetics</i> , 2011, 54, e495-e500.	0.7	10
116	Familial short stature due to a 5q22.1-q23.2 duplication refines the 5q duplication spectrum. <i>European Journal of Medical Genetics</i> , 2011, 54, e521-e524.	0.7	5
117	The molecular basis of the cartilage-hair hypoplasia-anaxetic dysplasia spectrum. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2011, 25, 131-142.	2.2	46
118	The shortest of the short: Pericentrin mutations and beyond. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2011, 25, 125-130.	2.2	40
119	De novo nonsense mutations in ASXL1 cause Bohring-Opitz syndrome. <i>Nature Genetics</i> , 2011, 43, 729-731.	9.4	236
120	Genome-wide copy number profiling using a 100K SNP array reveals novel disease-related genes BORIS and TSHZ1 in juvenile angiofibroma. <i>International Journal of Oncology</i> , 2011, 39, 1143-51.	1.4	12
121	CEP152 is a genome maintenance protein disrupted in Seckel syndrome. <i>Nature Genetics</i> , 2011, 43, 23-26.	9.4	201
122	Clinical utility gene card for: Mowat-Wilson syndrome. <i>European Journal of Human Genetics</i> , 2011, 19, 4-4.	1.4	5
123	The MEF2C-Related and 5q14.3q15 Microdeletion Syndrome. <i>Molecular Syndromology</i> , 2011, 2, 164-170.	0.3	51
124	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.	1.5	220
125	NEK1 Mutations Cause Short-Rib Polydactyly Syndrome Type Majewski. <i>American Journal of Human Genetics</i> , 2011, 88, 106-114.	2.6	151
126	Expanding the clinical spectrum associated with defects in CNTNAP2 and NRXN1. <i>BMC Medical Genetics</i> , 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.	0.7	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.	1.0	178
129	In-Frame Deletion and Missense Mutations of the C-Terminal Helicase Domain of <i>SMARCA2</i> in Three Patients with Nicolaides-Baraitser Syndrome. Molecular Syndromology, 2011, 2, 237-244.	0.3	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.	3.9	89
131	Disturbed Wnt Signalling due to a Mutation in <i>CCDC88C</i> Causes an Autosomal Recessive Non-Syndromic Hydrocephalus with Medial Diverticulum. Molecular Syndromology, 2010, 1, 99-112.	0.3	82
132	Mesomelia-Synostoses Syndrome Results from Deletion of <i>SULF1</i> and <i>SLCO5A1</i> Genes at 8q13. American Journal of Human Genetics, 2010, 87, 95-100.	2.6	42
133	Severe clinical course of Hirschsprung disease in a Mowat-Wilson syndrome patient. Journal of Applied Genetics, 2010, 51, 111-113.	1.0	14
134	Mutations in <i>MEF2C</i> from the 5q14.3q15 microdeletion syndrome region are a frequent cause of severe mental retardation and diminish <i>MECP2</i> and <i>CDKL5</i> expression. Human Mutation, 2010, 31, 722-733.	1.1	163
135	The face of Noonan syndrome: Does phenotype predict genotype. American Journal of Medical Genetics, Part A, 2010, 152A, 1960-1966.	0.7	59
136	Altered TGF β 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.	1.4	132
137	Clinical utility gene card for: DiGeorge syndrome, velocardiofacial syndrome, Shprintzen syndrome, chromosome 22q11.2 deletion syndrome (22q11.2, <i>TBX1</i>). European Journal of Human Genetics, 2010, 18, 1071-1071.	1.4	6
138	Mutations in <i>GRIN2A</i> and <i>GRIN2B</i> encoding regulatory subunits of NMDA receptors cause variable neurodevelopmental phenotypes. Nature Genetics, 2010, 42, 1021-1026.	9.4	431
139	Disruption of <i>ST5</i> is associated with mental retardation and multiple congenital anomalies. Journal of Medical Genetics, 2010, 47, 91-98.	1.5	12
140	Unmasking of a Recessive <i>SCARF2</i> Mutation by a 22q11.12 de novo Deletion in a Patient with Van den Ende-Gupta Syndrome. Molecular Syndromology, 2010, 1, 239-245.	0.3	32
141	Systematic survey of variants in <i>TBX1</i> 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.	1.2	61
142	Comprehensive genotype-phenotype analysis in 230 patients with tetralogy of Fallot. Journal of Medical Genetics, 2010, 47, 321-331.	1.5	126
143	Microcephalin and pericentrin regulate mitotic entry via centrosome-associated <i>Chk1</i> . Journal of Cell Biology, 2009, 185, 1149-1157.	2.3	83
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