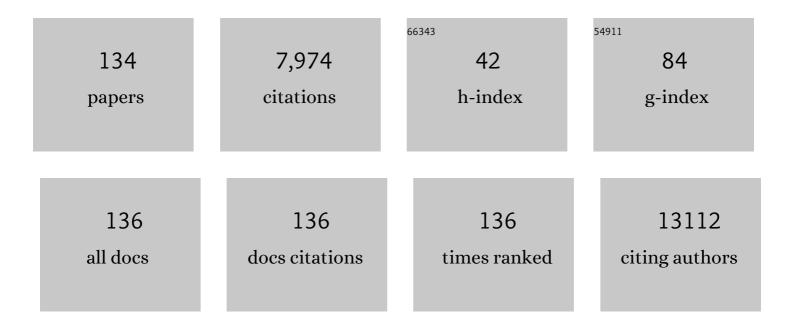
Andreas Tzschach

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	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
2	Deep sequencing reveals 50 novel genes for recessive cognitive disorders. Nature, 2011, 478, 57-63.	27.8	805
3	Mutations in the JARID1C Gene, Which Is Involved in Transcriptional Regulation and Chromatin Remodeling, Cause X-Linked Mental Retardation. American Journal of Human Genetics, 2005, 76, 227-236.	6.2	349
4	<i>STXBP1</i> encephalopathy. Neurology, 2016, 86, 954-962.	1.1	264
5	Mutations in NSUN2 Cause Autosomal- Recessive Intellectual Disability. American Journal of Human Genetics, 2012, 90, 847-855.	6.2	243
6	X-exome sequencing of 405 unresolved families identifies seven novel intellectual disability genes. Molecular Psychiatry, 2016, 21, 133-148.	7.9	243
7	Mutations in the Small GTPase Gene RAB39B Are Responsible for X-linked Mental Retardation Associated with Autism, Epilepsy, and Macrocephaly. American Journal of Human Genetics, 2010, 86, 185-195.	6.2	220
8	Cranioectodermal Dysplasia, Sensenbrenner Syndrome, Is a Ciliopathy Caused by Mutations in the IFT122 Gene. American Journal of Human Genetics, 2010, 86, 949-956.	6.2	180
9	A novel X-linked recessive mental retardation syndrome comprising macrocephaly and ciliary dysfunction is allelic to oral–facial–digital type I syndrome. Human Genetics, 2006, 120, 171-178.	3.8	166
10	Mutations in the polyglutamine binding protein 1 gene cause X-linked mental retardation. Nature Genetics, 2003, 35, 313-315.	21.4	139
11	A Defect in the Ionotropic Glutamate Receptor 6 Gene (GRIK2) Is Associated with Autosomal Recessive Mental Retardation. American Journal of Human Genetics, 2007, 81, 792-798.	6.2	137
12	Mutation Update for Kabuki Syndrome Genes <i>KMT2D</i> and <i>KDM6A</i> and Further Delineation of X-Linked Kabuki Syndrome Subtype 2. Human Mutation, 2016, 37, 847-864.	2.5	134
13	A balanced chromosomal translocation disrupting <i>ARHGEF9</i> is associated with epilepsy, anxiety, aggression, and mental retardation. Human Mutation, 2009, 30, 61-68.	2.5	131
14	Genetics of intellectual disability in consanguineous families. Molecular Psychiatry, 2019, 24, 1027-1039.	7.9	131
15	Structural variation in Xq28: MECP2 duplications in 1% of patients with unexplained XLMR and in 2% of male patients with severe encephalopathy. European Journal of Human Genetics, 2009, 17, 444-453.	2.8	130
16	A Defect in the TUSC3 Gene Is Associated with Autosomal Recessive Mental Retardation. American Journal of Human Genetics, 2008, 82, 1158-1164.	6.2	127
17	NovelJARID1C/SMCX mutations in patients with X-linked mental retardation. Human Mutation, 2006, 27, 389-389.	2.5	120
18	Identification of Mutations in TRAPPC9, which Encodes the NIK- and IKK-β-Binding Protein, in Nonsyndromic Autosomal-Recessive Mental Retardation. American Journal of Human Genetics, 2009, 85, 909-915.	6.2	120

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19	Mapping translocation breakpoints by next-generation sequencing. Genome Research, 2008, 18, 1143-1149.	5.5	118
20	Mutations in autism susceptibility candidate 2 (AUTS2) in patients with mental retardation. Human Genetics, 2007, 121, 501-509.	3.8	116
21	Next-generation sequencing in X-linked intellectual disability. European Journal of Human Genetics, 2015, 23, 1513-1518.	2.8	112
22	High frequency of submicroscopic genomic aberrations detected by tiling path array comparative genome hybridisation in patients with isolated congenital heart disease. Journal of Medical Genetics, 2008, 45, 704-709.	3.2	110
23	Clinical Presentation of a Complex Neurodevelopmental Disorder Caused by Mutations in ADNP. Biological Psychiatry, 2019, 85, 287-297.	1.3	108
24	Mutation frequencies of X-linked mental retardation genes in families from the EuroMRX consortium. Human Mutation, 2007, 28, 207-208.	2.5	103
25	Homozygosity mapping in consanguineous families reveals extreme heterogeneity of non-syndromic autosomal recessive mental retardation and identifies 8 novel gene loci. Human Genetics, 2007, 121, 43-48.	3.8	92
26	ST3GAL3 Mutations Impair the Development of Higher Cognitive Functions. American Journal of Human Genetics, 2011, 89, 407-414.	6.2	89
27	A Noncoding, Regulatory Mutation Implicates HCFC1 in Nonsyndromic Intellectual Disability. American Journal of Human Genetics, 2012, 91, 694-702.	6.2	89
28	A clinical and molecular genetic study of 112 Iranian families with primary microcephaly. Journal of Medical Genetics, 2010, 47, 823-828.	3.2	87
29	Multiple giant cell lesions in patients with Noonan syndrome and cardio-facio-cutaneous syndrome. European Journal of Human Genetics, 2009, 17, 420-425.	2.8	79
30	Mutation of the conserved polyadenosine RNA binding protein, ZC3H14/dNab2, impairs neural function in <i>Drosophila</i> and humans. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12390-12395.	7.1	77
31	Mutations in the Alpha 1,2-Mannosidase Gene, MAN1B1, Cause Autosomal-Recessive Intellectual Disability. American Journal of Human Genetics, 2011, 89, 176-182.	6.2	73
32	SNP array-based homozygosity mapping reveals MCPH1 deletion in family with autosomal recessive mental retardation and mild microcephaly. Human Genetics, 2006, 118, 708-715.	3.8	67
33	Chromosome deletions in 13q33–34: Report of four patients and review of the literature. American Journal of Medical Genetics, Part A, 2008, 146A, 337-342.	1.2	63
34	Breakpoint analysis of balanced chromosome rearrangements by next-generation paired-end sequencing. European Journal of Human Genetics, 2010, 18, 539-543.	2.8	61
35	MCT8 mutation analysis and identification of the first female with Allan–Herndon–Dudley syndrome due to loss of MCT8 expression. European Journal of Human Genetics, 2008, 16, 1029-1037.	2.8	56
36	Characterization of a 5.3 Mb deletion in 15q14 by comparative genomic hybridization using a whole genome "tiling path―BAC array in a girl with heart defect, cleft palate, and developmental delay. American Journal of Medical Genetics, Part A, 2007, 143A, 172-178.	1.2	55

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37	Mutations of the aminoacyl-tRNA-synthetases SARS and WARS2 are implicated in the etiology of autosomal recessive intellectual disability. Human Mutation, 2017, 38, 621-636.	2.5	54
38	Next generation sequencing in a family with autosomal recessive Kahrizi syndrome (OMIM 612713) reveals a homozygous frameshift mutation in SRD5A3. European Journal of Human Genetics, 2011, 19, 115-117.	2.8	52
39	Identification of a nonsense mutation in the very low-density lipoprotein receptor gene (VLDLR) in an Iranian family with dysequilibrium syndrome. European Journal of Human Genetics, 2008, 16, 270-273.	2.8	50
40	Mutation screening in 86 known X-linked mental retardation genes by droplet-based multiplex PCR and massive parallel sequencing. The HUGO Journal, 2009, 3, 41-49.	4.1	48
41	Characterisation of de novo MAPK10/JNK3 truncation mutations associated with cognitive disorders in two unrelated patients. Human Genetics, 2013, 132, 461-471.	3.8	48
42	The molecular and phenotypic spectrum of <i><scp>IQSEC</scp>2</i> â€related epilepsy. Epilepsia, 2016, 57, 1858-1869.	5.1	46
43	Novel missense mutations in the ubiquitinationâ€related gene <i>UBE2A</i> cause a recognizable Xâ€linked mental retardation syndrome. Clinical Genetics, 2010, 77, 541-551.	2.0	45
44	Autosomal recessive mental retardation: homozygosity mapping identifies 27 single linkage intervals, at least 14 novel loci and several mutation hotspots. Human Genetics, 2011, 129, 141-148.	3.8	45
45	Expanded mutational spectrum in Cohen syndrome, tissue expression, and transcript variants of <i>COH1</i> . Human Mutation, 2009, 30, E404-E420.	2.5	44
46	Musculoskeletal Disease in MDA5â€Related Type I Interferonopathy: A Mendelian Mimic of Jaccoud's Arthropathy. Arthritis and Rheumatology, 2017, 69, 2081-2091.	5.6	44
47	A novel nonsense mutation in <i>TUSC3</i> is responsible for nonâ€syndromic autosomal recessive mental retardation in a consanguineous Iranian family. American Journal of Medical Genetics, Part A, 2011, 155, 1976-1980.	1.2	43
48	<i>KCNC1</i> â€related disorders: new de novo variants expand the phenotypic spectrum. Annals of Clinical and Translational Neurology, 2019, 6, 1319-1326.	3.7	43
49	Variants in <i>CUL4B</i> are Associated with Cerebral Malformations. Human Mutation, 2015, 36, 106-117.	2.5	37
50	Diagnostic value of partial exome sequencing in developmental disorders. PLoS ONE, 2018, 13, e0201041.	2.5	36
51	Characterization of interstitial Xp duplications in two families by tiling path array CGH. American Journal of Medical Genetics, Part A, 2008, 146A, 197-203.	1.2	35
52	A distinctive gene expression fingerprint in mentally retarded male patients reflects disease-causing defects in the histone demethylase KDM5C. PathoGenetics, 2010, 3, 2.	5.7	35
53	Identification of a novel CDKL5 exon and pathogenic mutations in patients with severe mental retardation, early-onset seizures and Rett-like features. Neurogenetics, 2011, 12, 165-167.	1.4	34
54	<i>De novo</i> partial deletion in <i>GRID2</i> presenting with complicated spastic paraplegia. Muscle and Nerve, 2014, 49, 289-292.	2.2	33

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55	<i><scp>PIK3R1</scp></i> mutations in <scp>SHORT</scp> syndrome. Clinical Genetics, 2014, 86, 292-294.	2.0	31
56	Epilepsy is not a mandatory feature of STXBP1 associated ataxia-tremor-retardation syndrome. European Journal of Paediatric Neurology, 2016, 20, 661-665.	1.6	30
57	Radiation-Induced Late Effects in Two Affected Individuals of the Lilo Radiation Accident. Radiation Research, 2007, 167, 615-623.	1.5	29
58	Cohen syndrome diagnosis using whole genome arrays. Journal of Medical Genetics, 2011, 48, 136-140.	3.2	29
59	Novel <i>WDR35</i> mutations in patients with cranioectodermal dysplasia (Sensenbrenner) Tj ETQq1 1 0.7843	14.rgBT /0 2.9	Overlock 10 T
60	Mutations in the histamine <i>N</i> -methyltransferase gene, <i>HNMT</i> , are associated with nonsyndromic autosomal recessive intellectual disability. Human Molecular Genetics, 2015, 24, 5697-5710.	2.9	27
61	Transitioning the Molecular Tumor Board from Proof of Concept to Clinical Routine: A German Single-Center Analysis. Cancers, 2021, 13, 1151.	3.7	27
62	Mutation screening of ASMT, the last enzyme of the melatonin pathway, in a large sample of patients with Intellectual Disability. BMC Medical Genetics, 2011, 12, 17.	2.1	25
63	Clinical and functional characterisation of the combined respiratory chain defect in two sisters due to autosomal recessive mutations in MTFMT. Mitochondrion, 2013, 13, 743-748.	3.4	25
64	Craniosynostosis in a patient with 2q37.3 deletion 5q34 duplication: Association of extra copy of <i>MSX2</i> with craniosynostosis. American Journal of Medical Genetics, Part A, 2009, 149A, 1544-1549.	1.2	24
65	Isolated NIBPL missense mutations that cause Cornelia de Lange syndrome alter MAU2 interaction. European Journal of Human Genetics, 2012, 20, 271-276.	2.8	24
66	Kohlschütter-Tönz Syndrome: Mutations in <i>ROGDI</i> and Evidence of Genetic Heterogeneity. Human Mutation, 2013, 34, 296-300.	2.5	24
67	Novel <scp><i>SLC9A6</i></scp> mutations in two families with Christianson syndrome. Clinical Genetics, 2013, 83, 596-597.	2.0	24
68	The power of the Mediator complex-Expanding the genetic architecture and phenotypic spectrum of MED12 -related disorders. Clinical Genetics, 2018, 94, 450-456.	2.0	24
69	Chromosome aberrations involving 10q22: report of three overlapping interstitial deletions and a balanced translocation disrupting C10orf11. European Journal of Human Genetics, 2010, 18, 291-295.	2.8	22
70	X-linked intellectual disability type Nascimento is a clinically distinct, probably underdiagnosed entity. Orphanet Journal of Rare Diseases, 2013, 8, 146.	2.7	22
71	Hybridisation-based resequencing of 17 X-linked intellectual disability genes in 135 patients reveals novel mutations in ATRX, SLC6A8 and PQBP1. European Journal of Human Genetics, 2011, 19, 717-720.	2.8	21
72	Sema3a plays a role in the pathogenesis of CHARGE syndrome. Human Molecular Genetics, 2018, 27, 1343-1352.	2.9	20

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73	X-linked mental retardation: a comprehensive molecular screen of 47 candidate genes from a 7.4 Mb interval in Xp11. European Journal of Human Genetics, 2007, 15, 68-75.	2.8	19
74	Xq22.3–q23 deletion including <i>ACSL4</i> in a patient with intellectual disability. American Journal of Medical Genetics, Part A, 2013, 161, 860-864.	1.2	19
75	Next-generation panel sequencing identifies NF1 germline mutations in three patients with pheochromocytoma but no clinical diagnosis of neurofibromatosis type 1. European Journal of Endocrinology, 2018, 178, K1-K9.	3.7	19
76	Czech dysplasia: Report of a large family and further delineation of the phenotype. American Journal of Medical Genetics, Part A, 2008, 146A, 1859-1864.	1.2	18
77	Characterization of an interstitial 4q32 deletion in a patient with mental retardation and a complex chromosome rearrangement. American Journal of Medical Genetics, Part A, 2010, 152A, 1008-1012.	1.2	18
78	Floatingâ€Harbor syndrome: <i><scp>SRCAP</scp></i> mutations are not restricted to exon 34. Clinical Genetics, 2014, 85, 498-499.	2.0	18
79	Variable clinical phenotype in two siblings with Aicardi-Goutières syndrome type 6 and a novel mutation in the ADAR gene. European Journal of Paediatric Neurology, 2018, 22, 186-189.	1.6	18
80	BOD1 Is Required for Cognitive Function in Humans and Drosophila. PLoS Genetics, 2016, 12, e1006022.	3.5	18
81	An autosomal recessive syndrome of severe mental retardation, cataract, coloboma and kyphosis maps to the pericentromeric region of chromosome 4. European Journal of Human Genetics, 2009, 17, 125-128.	2.8	17
82	Heterotaxy and cardiac defect in a girl with chromosome translocation t(X;1)(q26;p13.1) and involvement of ZIC3. European Journal of Human Genetics, 2006, 14, 1317-1320.	2.8	16
83	Novel <i>GDI1</i> mutation in a large family with nonsyndromic Xâ€linked intellectual disability. American Journal of Medical Genetics, Part A, 2011, 155, 3067-3070.	1.2	16
84	A novel <i>ALDH5A1</i> mutation is associated with succinic semialdehyde dehydrogenase deficiency and severe intellectual disability in an Iranian family. American Journal of Medical Genetics, Part A, 2013, 161, 1915-1922.	1.2	16
85	Deletions in 14q24.1q24.3 are associated with congenital heart defects, brachydactyly, and mild intellectual disability. American Journal of Medical Genetics, Part A, 2014, 164, 620-626.	1.2	16
86	Characterization of a 16 Mb interstitial chromosome 7q21 deletion by tiling path array CGH. American Journal of Medical Genetics, Part A, 2007, 143A, 333-337.	1.2	15
87	Ready to clone: CNV detection and breakpoint fine-mapping in breast and ovarian cancer susceptibility genes by high-resolution array CGH. Breast Cancer Research and Treatment, 2016, 159, 585-590.	2.5	15
88	A mosaic maternal splice donor mutation in the EHMT1 gene leads to aberrant transcripts and to Kleefstra syndrome in the offspring. European Journal of Human Genetics, 2013, 21, 887-890.	2.8	14
89	Congenital CLN disease in two siblings. Wiener Medizinische Wochenschrift, 2015, 165, 210-213.	1.1	14
90	Interstitial deletion 2p11.2–p12: Report of a patient with mental retardation and review of the literature. American Journal of Medical Genetics, Part A, 2009, 149A, 242-245.	1.2	13

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91	Christianson syndrome in a patient with an interstitial Xq26.3 deletion. American Journal of Medical Genetics, Part A, 2011, 155, 2771-2774.	1.2	13
92	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
93	Molecular cytogenetic analysis of a de novo interstitial deletion of 5q23.3q31.2 and its phenotypic consequences. American Journal of Medical Genetics, Part A, 2006, 140A, 496-502.	1.2	12
94	Pierpont syndrome: report of a new patient. Clinical Dysmorphology, 2017, 26, 205-208.	0.3	12
95	Skewed Xâ€inactivation in a family with <i>DLG3â€</i> associated Xâ€linked intellectual disability. American Journal of Medical Genetics, Part A, 2017, 173, 2545-2550.	1.2	12
96	ABSENCE OF Yq MICRODELETIONS IN INFERTILE MEN. Archives of Andrology, 2001, 47, 167-171.	1.0	11
97	Interstitial 3p25.3–p26.1 deletion in a patient with intellectual disability. American Journal of Medical Genetics, Part A, 2012, 158A, 2587-2590.	1.2	11
98	Angelman syndrome and severe infections in a patient with de novo 15q11.2–q13.1 deletion and maternally inherited 2q21.3 microdeletion. Gene, 2013, 512, 453-455.	2.2	11
99	Tentative clinical diagnosis of Lujanâ€Fryns syndrome—A conglomeration of different genetic entities?. American Journal of Medical Genetics, Part A, 2016, 170, 94-102.	1.2	11
100	Novel PRPS1 gain-of-function mutation in a patient with congenital hyperuricemia and facial anomalies. , 2017, 173, 2736-2742.		11
101	Marfan Syndrome Caused by Disruption of the FBN1 Gene due to A Reciprocal Chromosome Translocation. Genes, 2021, 12, 1836.	2.4	11
102	Autosomal dominant inheritance in a large family with focal facial dermal dysplasia (Brauer–Setleis) Tj ETQq0 (0 rgBT /0 1.2	Dverlock 10 Tf
103	Interstitial 9q34.11–q34.13 deletion in a patient with severe intellectual disability, hydrocephalus, and cleft lip/palate. American Journal of Medical Genetics, Part A, 2012, 158A, 1709-1712.	1.2	10
104	A Novel SLC6A8 Mutation in a Large Family with X-Linked Intellectual Disability: Clinical and Proton Magnetic Resonance Spectroscopy Data of Both Hemizygous Males and Heterozygous Females. JIMD Reports, 2013, 13, 91-99.	1.5	10
105	Novel truncating PPM1D mutation in a patient with intellectual disability. European Journal of Medical Genetics, 2019, 62, 70-72.	1.3	10
106	Skeletal abnormalities are common features in Ayméâ€Gripp syndrome. Clinical Genetics, 2020, 97, 362-369.	2.0	10

107	New Cav1.2 Channelopathy with High-Functioning Autism, Affective Disorder, Severe Dental Enamel Defects, a Short QT Interval, and a Novel CACNA1C Loss-of-Function Mutation. International Journal	4.1	10
107	of Molecular Sciences, 2020, 21, 8611.		10
108	A Novel MGST2 Non-Synonymous Mutation in a Chinese Pedigree with Psoriasis Vulgaris. Journal of	0.7	9
	Investigative Dermatology, 2006, 126, 1003-1005.		-

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109	Molecular cytogenetic analysis of a de novo interstitial chromosome 10q22 deletion. American Journal of Medical Genetics, Part A, 2006, 140A, 1108-1110.	1.2	9
110	Mirror-Image Asymmetry in Monozygotic Twins with Kabuki Syndrome. Molecular Syndromology, 2012, 3, 94-97.	0.8	9
111	Parental Origin of de novo Cytogenetically Balanced Reciprocal Non-Robertsonian Translocations. Cytogenetic and Genome Research, 2012, 136, 242-245.	1.1	9
112	A newly recognized autosomal recessive syndrome affecting neurologic function and vision. American Journal of Medical Genetics, Part A, 2013, 161, 1207-1213.	1.2	9
113	Molecular characterization of a balanced chromosome translocation in psoriasis vulgaris. Clinical Genetics, 2005, 69, 189-193.	2.0	8
114	Novel VPS33B mutation in a patient with autosomal recessive keratodermaâ€ichthyosisâ€deafness syndrome. American Journal of Medical Genetics, Part A, 2018, 176, 2862-2866.	1.2	8
115	Interstitial 1q23.3q24.1 deletion in a patient with renal malformation, congenital heart disease, and mild intellectual disability. American Journal of Medical Genetics, Part A, 2016, 170, 2394-2399.	1.2	7
116	PUF60-SCRIB fusion transcript in a patient with 8q24.3 microdeletion and atypical Verheij syndrome. European Journal of Medical Genetics, 2019, 62, 103587.	1.3	7
117	Blepharophimosis-Ptosis-Epicanthus Inversus Syndrome in a Girl with Chromosome Translocation t(2;3)(q33;q23). Ophthalmic Genetics, 2008, 29, 37-40.	1.2	6
118	Interstitial 1p32.1p32.3 deletion in a patient with multiple congenital anomalies. American Journal of Medical Genetics, Part A, 2015, 167, 2406-2410.	1.2	6
119	11q14.1â€11q22.1 deletion in a 1â€yearâ€old male with minor dysmorphic features. American Journal of Medical Genetics, Part A, 2010, 152A, 2651-2655.	1.2	5
120	Novel ADAMTSL2-mutations in a patient with geleophysic dysplasia type I. Clinical Dysmorphology, 2016, 25, 106-109.	0.3	5
121	Alopeciamental retardation syndrome: clinical and molecular characterization of four patients. British Journal of Dermatology, 2008, 159, ???-???.	1.5	4
122	Interstitial duplication of chromosome region 1q25.1q25.3: Report of a patient with mild cognitive deficits, tall stature and facial dysmorphisms. American Journal of Medical Genetics, Part A, 2015, 167, 653-656.	1.2	4
123	Posterior amorphous corneal dystrophy in a patient with 12q21.33 deletion. Ophthalmic Genetics, 2018, 39, 645-647.	1.2	4
124	<i>PIGN</i> encephalopathy: Characterizing the epileptology. Epilepsia, 2022, 63, 974-991.	5.1	4
125	Molecular breakpoint analysis and relevance of variable mosaicism in a woman with short stature, primary amenorrhea, unilateral gonadoblastoma, and a 46,X,del(Y)(q11)/45,X karyotype. American Journal of Medical Genetics Part A, 2002, 112, 51-55.	2.4	3
126	Brachyphalangy, polydactyly and tibial aplasia/hypoplasia syndrome (OMIM 609945): case report and review of the literature. European Journal of Pediatrics, 2010, 169, 1535-1539.	2.7	3

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127	Hypergonadotropic hypogonadism in a patient with inv ins (2;4). Journal of Developmental and Physical Disabilities, 2009, 32, 226-230.	3.6	2
128	12q24.33 deletion: Report of a patient with intellectual disability and review of the literature. American Journal of Medical Genetics, Part A, 2013, 161, 1409-1413.	1.2	2
129	The Importance of Extended Analysis Using Current Molecular Genetic Methods Based on the Example of a Cohort of 228 Patients with Hereditary Breast and Ovarian Cancer Syndrome. Genes, 2021, 12, 1483.	2.4	1
130	Genetik der nichtsyndromalen geistigen Behinderung. Medizinische Genetik, 2009, 21, 231-236.	0.2	0
131	Reply to 5q35 duplication and Hunter–McAlpine syndrome: Missing the link. American Journal of Medical Genetics, Part A, 2010, 152A, 804-804.	1.2	0
132	Chromosome aberration associated with hippocampal impairment. Psychiatry Research - Neuroimaging, 2016, 254, 1-2.	1.8	0
133	Abstract LB-044: Germline mutations in patients with hereditary breast and ovarian cancer establish ERCC2 as a cancer susceptibility gene. , 2015, , .		Ο
134	Abstract 3391: Next generation sequencing paves the way for personalized medicine in pheochromocytoma and paraganglioma patients and their families. , 2017, , .		0