

Elaine H Zackai

List of Publications by Year in descending order

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263
papers

13,363
citations

34105

52
h-index

31849

101
g-index

275
all docs

275
docs citations

275
times ranked

14075
citing authors

#	ARTICLE	IF	CITATIONS
1	22q11.2 deletion syndrome. Nature Reviews Disease Primers, 2015, 1, 15071.	30.5	954
2	Psychiatric Disorders From Childhood to Adulthood in 22q11.2 Deletion Syndrome: Results From the International Consortium on Brain and Behavior in 22q11.2 Deletion Syndrome. American Journal of Psychiatry, 2014, 171, 627-639.	7.2	645
3	Mutations in the homeodomain of the human SIX3 gene cause holoprosencephaly. Nature Genetics, 1999, 22, 196-198.	21.4	398
4	Deletions and microdeletions of 22q11.2 in velo-cardio-facial syndrome. American Journal of Medical Genetics Part A, 1992, 44, 261-268.	2.4	387
5	Mutations in TGIF cause holoprosencephaly and link NODAL signalling to human neural axis determination. Nature Genetics, 2000, 25, 205-208.	21.4	368
6	Identical mutations in three different fibroblast growth factor receptor genes in autosomal dominant craniosynostosis syndromes. Nature Genetics, 1996, 14, 174-176.	21.4	306
7	Growth Charts for Children With Down Syndrome: 1 Month to 18 Years of Age. Pediatrics, 1988, 81, 102-110.	2.1	298
8	Mutations of ARX are associated with striking pleiotropy and consistent genotype-phenotype correlation. Human Mutation, 2004, 23, 147-159.	2.5	293
9	Analysis of 589,306 genomes identifies individuals resilient to severe Mendelian childhood diseases. Nature Biotechnology, 2016, 34, 531-538.	17.5	273
10	Cognitive and behavior profile of preschool children with chromosome 22q11.2 deletion. American Journal of Medical Genetics Part A, 1999, 85, 127-133.	2.4	263
11	Immunologic features of chromosome 22q11.2 deletion syndrome (DiGeorge) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 342 Td (Cy	1.8	240
12	Optic pathway and hypothalamic/chiasmatic gliomas in children younger than age 5 years with a 6-year follow-up. Cancer, 1995, 75, 1051-1059.	4.1	234
13	Mutations in DDX3X Are a Common Cause of Unexplained Intellectual Disability with Gender-Specific Effects on Wnt Signaling. American Journal of Human Genetics, 2015, 97, 343-352.	6.2	230
14	Neonatal adrenoleukodystrophy: New cases, biochemical studies, and differentiation from Zellweger and related peroxisomal polydystrophy syndromes. American Journal of Medical Genetics Part A, 1986, 23, 869-901.	2.4	216
15	Reanalysis of Clinical Exome Sequencing Data. New England Journal of Medicine, 2019, 380, 2478-2480.	27.0	205
16	<i>GRIN2B</i> encephalopathy: novel findings on phenotype, variant clustering, functional consequences and treatment aspects. Journal of Medical Genetics, 2017, 54, 460-470.	3.2	190
17	Down syndrome congenital heart disease: A narrowed region and a candidate gene. Genetics in Medicine, 2001, 3, 91-101.	2.4	168
18	Site-specific reciprocal translocation, t(11;22) (q23;q11), in several unrelated families with 3:1 meiotic disjunction. American Journal of Medical Genetics Part A, 1980, 7, 507-521.	2.4	165

#	ARTICLE	IF	CITATIONS
19	The 22q11.2 Deletion: Screening, Diagnostic Workup, and Outcome of Results; Report on 181 Patients. <i>Genetic Testing and Molecular Biomarkers</i> , 1997, 1, 99-108.	1.7	150
20	Genotype-Phenotype Correlation in NF1: Evidence for a More Severe Phenotype Associated with Missense Mutations Affecting NF1 Codons 844-848. <i>American Journal of Human Genetics</i> , 2018, 102, 69-87.	6.2	144
21	High Incidence of Noonan Syndrome Features Including Short Stature and Pulmonic Stenosis in Patients carrying NF1 Missense Mutations Affecting p.Arg1809: Genotype-Phenotype Correlation. <i>Human Mutation</i> , 2015, 36, 1052-1063.	2.5	143
22	Kabuki syndrome genes <i>KMT2D</i> and <i>KDM6A</i> : functional analyses demonstrate critical roles in craniofacial, heart and brain development. <i>Human Molecular Genetics</i> , 2015, 24, 4443-4453.	2.9	142
23	Aberrant interchromosomal exchanges are the predominant cause of the 22q11.2 deletion. <i>Human Molecular Genetics</i> , 2004, 13, 417-428.	2.9	133
24	T-cell homeostasis in humans with thymic hypoplasia due to chromosome 22q11.2 deletion syndrome. <i>Blood</i> , 2004, 103, 1020-1025.	1.4	119
25	Juvenile rheumatoid arthritis-like polyarthritis in chromosome 22q11.2 deletion syndrome (deGeorge) Tj ETQq1 1 0.784314 rgBT /Overl Rheumatism, 1997, 40, 430-436.	6.7	115
26	Clinical, cytogenetic, and pedigree findings in 18 cases of Aicardi syndrome. <i>American Journal of Medical Genetics Part A</i> , 1989, 32, 461-467.	2.4	113
27	Germline gain-of-function mutations in <i>AFF4</i> cause a developmental syndrome functionally linking the super elongation complex and cohesin. <i>Nature Genetics</i> , 2015, 47, 338-344.	21.4	109
28	Dominant Mutations in <i>KAT6A</i> Cause Intellectual Disability with Recognizable Syndromic Features. <i>American Journal of Human Genetics</i> , 2015, 96, 507-513.	6.2	107
29	What is new with 22q? An update from the 22q and You Center at the Children's Hospital of Philadelphia. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2058-2069.	1.2	106
30	Hemizygous mutations in <i>SNAP29</i> unmask autosomal recessive conditions and contribute to atypical findings in patients with 22q11.2DS. <i>Journal of Medical Genetics</i> , 2013, 50, 80-90.	3.2	104
31	<i>DYRK1A</i> haploinsufficiency causes a new recognizable syndrome with microcephaly, intellectual disability, speech impairment, and distinct facies. <i>European Journal of Human Genetics</i> , 2015, 23, 1473-1481.	2.8	101
32	Oral-facial-digital syndrome type VI (VÃ;radi syndrome): Further clinical delineation. <i>American Journal of Medical Genetics Part A</i> , 1990, 35, 360-369.	2.4	97
33	Deletions of different segments of the long arm of chromosome 4. <i>American Journal of Medical Genetics Part A</i> , 1981, 8, 73-89.	2.4	93
34	Mutations in <i>CDC45</i> , Encoding an Essential Component of the Pre-initiation Complex, Cause Meier-Gorlin Syndrome and Craniosynostosis. <i>American Journal of Human Genetics</i> , 2016, 99, 125-138.	6.2	92
35	Mutations in the human <i>TWIST</i> gene. <i>Human Mutation</i> , 2000, 15, 150-155.	2.5	86
36	Clinical spectrum of individuals with pathogenic <i>NF1</i> missense variants affecting p.Met1149, p.Arg1276, and p.Lys1423: genotype-phenotype study in neurofibromatosis type 1. <i>Human Mutation</i> , 2020, 41, 299-315.	2.5	80

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37	Rare Genome-Wide Copy Number Variation and Expression of Schizophrenia in 22q11.2 Deletion Syndrome. <i>American Journal of Psychiatry</i> , 2017, 174, 1054-1063.	7.2	77
38	Skeletal anomalies and deformities in patients with deletions of 22q11. . 1997, 72, 210-215.		75
39	Phenotype of the fibroblast growth factor receptor 2 Ser351Cys mutation: Pfeiffer syndrome type III. <i>American Journal of Medical Genetics Part A</i> , 1998, 78, 356-360.	2.4	72
40	Neurodevelopmental outcomes in preschool survivors of the Fontan procedure. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2014, 147, 1276-1283.e5.	0.8	71
41	Gene domain-specific DNA methylation epigenatures highlight distinct molecular entities of ADNP syndrome. <i>Clinical Epigenetics</i> , 2019, 11, 64.	4.1	71
42	A dyadic approach to the delineation of diagnostic entities in clinical genomics. <i>American Journal of Human Genetics</i> , 2021, 108, 8-15.	6.2	71
43	Maternal uniparental disomy of chromosome 20: a novel imprinting disorder of growth failure. <i>Genetics in Medicine</i> , 2016, 18, 309-315.	2.4	69
44	Longitudinal Analysis of Lymphocyte Function and Numbers in the First Year of Life in Chromosome 22q11.2 Deletion Syndrome (DiGeorge Syndrome/Velocardiofacial Syndrome). <i>Vaccine Journal</i> , 1999, 6, 906-911.	2.6	68
45	De Novo Mutations of RERE Cause a Genetic Syndrome with Features that Overlap Those Associated with Proximal 1p36 Deletions. <i>American Journal of Human Genetics</i> , 2016, 98, 963-970.	6.2	67
46	22q11.2 duplication syndrome: elevated rate of autism spectrum disorder and need for medical screening. <i>Molecular Autism</i> , 2016, 7, 27.	4.9	67
47	Copy-Number Variation of the Glucose Transporter Gene SLC2A3 and Congenital Heart Defects in the 22q11.2 Deletion Syndrome. <i>American Journal of Human Genetics</i> , 2015, 96, 753-764.	6.2	62
48	Expanding the clinical phenotype of individuals with a 3-bp in-frame deletion of the NF1 gene (c.2970_2972del): an update of genotypeâ€“phenotype correlation. <i>Genetics in Medicine</i> , 2019, 21, 867-876.	2.4	62
49	Polytopic anomalies with agenesis of the lower vertebral column. <i>American Journal of Medical Genetics Part A</i> , 1999, 87, 99-114.	2.4	61
50	Aberrant Cortical Morphometry in the 22q11.2 Deletion Syndrome. <i>Biological Psychiatry</i> , 2015, 78, 135-143.	1.3	61
51	22q11.2 Deletion syndrome and obstructive sleep apnea. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2014, 78, 1360-1364.	1.0	58
52	Mouse and Human CRKL Is Dosage Sensitive for Cardiac Outflow Tract Formation. <i>American Journal of Human Genetics</i> , 2015, 96, 235-244.	6.2	58
53	Disrupted auto-regulation of the spliceosomal gene SNRPB causes cerebroâ€“costoâ€“mandibular syndrome. <i>Nature Communications</i> , 2014, 5, 4483.	12.8	57
54	De Novo Truncating Mutations in AHDC1 in Individuals with Syndromic Expressive Language Delay, Hypotonia, and Sleep Apnea. <i>American Journal of Human Genetics</i> , 2014, 94, 784-789.	6.2	57

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55	Congenital heart diseases and cardiovascular abnormalities in 22q11.2 deletion syndrome: From well-established knowledge to new frontiers. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2087-2098.	1.2	57
56	Natural history and genotype-phenotype correlations in 72 individuals with <i>SATB2</i> -associated syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 925-935.	1.2	57
57	Enlarged sylvian fissures in infants with interstitial deletion of chromosome 22q11. <i>American Journal of Medical Genetics Part A</i> , 1997, 74, 538-543.	2.4	56
58	Oculodentodigital dysplasia syndrome associated with abnormal cerebral white matter. <i>American Journal of Medical Genetics Part A</i> , 1991, 41, 18-20.	2.4	55
59	Increasing cumulative exposure to volatile anesthetic agents is associated with poorer neurodevelopmental outcomes in children with hypoplastic left heart syndrome. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2016, 152, 482-489.	0.8	55
60	Burden of potentially pathologic copy number variants is higher in children with isolated congenital heart disease and significantly impairs covariate-adjusted transplant-free survival. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2016, 151, 1147-1151.e4.	0.8	55
61	Gain-of-function mutations in <i>SMAD4</i> cause a distinctive repertoire of cardiovascular phenotypes in patients with Myhre syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2617-2631.	1.2	53
62	The CHD4-related syndrome: a comprehensive investigation of the clinical spectrum, genotype-phenotype correlations, and molecular basis. <i>Genetics in Medicine</i> , 2020, 22, 389-397.	2.4	53
63	Melnick-needles syndrome in males: A lethal multiple congenital anomalies syndrome. <i>American Journal of Medical Genetics Part A</i> , 1987, 27, 159-173.	2.4	52
64	Lateral meningocele syndrome: Three new patients and review of the literature. <i>American Journal of Medical Genetics Part A</i> , 1997, 70, 229-239.	2.4	52
65	Saethre-Chotzen syndrome with familial translocation at chromosome 7p22. <i>American Journal of Medical Genetics Part A</i> , 1993, 47, 637-639.	2.4	51
66	Histone Modifier Genes Alter Conotruncal Heart Phenotypes in 22q11.2 Deletion Syndrome. <i>American Journal of Human Genetics</i> , 2015, 97, 869-877.	6.2	49
67	Association of airway abnormalities with 22q11.2 deletion syndrome. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2017, 96, 11-14.	1.0	49
68	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
69	On lumping and splitting: A fetus with clinical findings of the oral-facial-digital syndrome type VI, the hydrolethalus syndrome, and the Pallister-Hall syndrome. <i>American Journal of Medical Genetics Part A</i> , 1991, 41, 548-556.	2.4	47
70	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.	3.2	47
71	Subthreshold Psychosis in 22q11.2 Deletion Syndrome: Multisite Naturalistic Study. <i>Schizophrenia Bulletin</i> , 2017, 43, 1079-1089.	4.3	47
72	The 22q11.2 deletion syndrome: Cancer predisposition, platelet abnormalities and cytopenias. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2121-2127.	1.2	47

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73	Mandibulofacial Dysostosis with Microcephaly: Mutation and Database Update. <i>Human Mutation</i> , 2016, 37, 148-154.	2.5	45
74	Neurocognitive profile in psychotic versus nonpsychotic individuals with 22q11.2 deletion syndrome. <i>European Neuropsychopharmacology</i> , 2016, 26, 1610-1618.	0.7	45
75	The Psychosis Spectrum in 22q11.2 Deletion Syndrome Is Comparable to That of Nondeleted Youths. <i>Biological Psychiatry</i> , 2017, 82, 17-25.	1.3	45
76	Elucidating the diagnostic odyssey of 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 936-944.	1.2	45
77	Rare copy number variants and congenital heart defects in the 22q11.2 deletion syndrome. <i>Human Genetics</i> , 2016, 135, 273-285.	3.8	43
78	Histone H3.3 beyond cancer: Germline mutations in <i>Histone 3 Family 3A and 3B</i> cause a previously unidentified neurodegenerative disorder in 46 patients. <i>Science Advances</i> , 2020, 6, .	10.3	43
79	Rates of autism and potential risk factors in children with congenital heart defects. <i>Congenital Heart Disease</i> , 2017, 12, 421-429.	0.2	42
80	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.	1.3	42
81	Complete Sequence of the 22q11.2 Allele in 1,053 Subjects with 22q11.2 Deletion Syndrome Reveals Modifiers of Conotruncal Heart Defects. <i>American Journal of Human Genetics</i> , 2020, 106, 26-40.	6.2	42
82	Holoprosencephaly: Association with interstitial deletion of 2p and review of the cytogenetic literature. <i>American Journal of Medical Genetics Part A</i> , 1988, 30, 929-938.	2.4	41
83	Expanding the fetal phenotype: Prenatal sonographic findings and perinatal outcomes in a cohort of patients with a confirmed 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1735-1741.	1.2	41
84	TWIST gene mutation in a patient with radial aplasia and craniosynostosis: Further evidence for heterogeneity of Baller-Gerold syndrome. , 1999, 82, 170-176.		40
85	Mutations in <i>SPECC1L</i> , encoding sperm antigen with calponin homology and coiled-coil domains 1-like, are found in some cases of autosomal dominant Opitz G/BBB syndrome. <i>Journal of Medical Genetics</i> , 2015, 52, 104-110.	3.2	40
86	NF1B Haploinsufficiency Is Associated with Intellectual Disability and Macrocephaly. <i>American Journal of Human Genetics</i> , 2018, 103, 752-768.	6.2	40
87	De novo variants in Myelin regulatory factor (MYRF) as candidates of a new syndrome of cardiac and urogenital anomalies. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 969-972.	1.2	39
88	Craniosynostosis: Another feature of the 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2005, 136A, 358-362.	1.2	38
89	MN1 C-terminal truncation syndrome is a novel neurodevelopmental and craniofacial disorder with partial rhombencephalosynapsis. <i>Brain</i> , 2020, 143, 55-68.	7.6	38
90	Cystic kidney disease in Hajdu-Cheney syndrome. <i>American Journal of Medical Genetics Part A</i> , 1995, 56, 25-30.	2.4	37

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91	Critical region within 22q11.2 linked to higher rate of autism spectrum disorder. <i>Molecular Autism</i> , 2017, 8, 58.	4.9	37
92	Mutations in topoisomerase III^2 result in a B cell immunodeficiency. <i>Nature Communications</i> , 2019, 10, 3644.	12.8	37
93	Partial duplication 1q: Report of four patients and review of the literature. <i>American Journal of Medical Genetics Part A</i> , 1990, 36, 137-143.	2.4	36
94	Molecular detection of a Yp/18 translocation in a 45,X holoprosencephalic male. <i>Human Genetics</i> , 1988, 80, 219-223.	3.8	35
95	New finding of Schinzel-Giedion syndrome: A case with a malignant sacrococcygeal teratoma. <i>American Journal of Medical Genetics Part A</i> , 1993, 47, 852-856.	2.4	35
96	Congenital heart disease in supernumerary der(22), t(11;22) syndrome. <i>Clinical Genetics</i> , 1986, 29, 269-275.	2.0	35
97	A Prospective Study of Influenza Vaccination and a Comparison of Immunologic Parameters in Children and Adults with Chromosome 22q11.2 Deletion Syndrome (DiGeorge) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50497 Td IS	3.8	35
98	Identification of 22q11.2 Deletion Syndrome via Newborn Screening for Severe Combined Immunodeficiency. <i>Journal of Clinical Immunology</i> , 2017, 37, 476-485.	3.8	35
99	Prenatal detection of Roberts-SC phocomelia syndrome: Report of 2 sibs with characteristic manifestations. <i>American Journal of Medical Genetics Part A</i> , 1989, 32, 390-394.	2.4	34
100	Expanding the genotypic and phenotypic spectrum in a diverse cohort of 104 individuals with Wiedemann-Steiner syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1649-1665.	1.2	34
101	Variance of IQ is partially dependent on deletion type among 1,427 22q11.2 deletion syndrome subjects. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2172-2181.	1.2	33
102	Ablepharon macrostomia syndrome with associated cutis laxa: Possible localization to 18q. <i>Human Genetics</i> , 1996, 97, 532-536.	3.8	32
103	Neuropathological Findings in Eight Children with Cerebro-oculo-facio-skeletal (COFS) Syndrome. <i>Journal of Neuropathology and Experimental Neurology</i> , 1997, 56, 1147-1157.	1.7	32
104	Further evidence that the Hajdu-Cheney syndrome and the ?serpentine fibula-polycystic kidney syndrome? are a single entity. , 1998, 78, 474-481.		32
105	Blepharo-cheilo-dontic (BCD) syndrome. , 1996, 65, 109-112.		31
106	Bilateral microtia and cleft palate in cousins with Diamond-Blackfan anemia. <i>American Journal of Medical Genetics Part A</i> , 2001, 101, 268-274.	2.4	31
107	Mosaic loss of 15q11q13 in a patient with hypomelanosis of Ito: is there a role for the P gene?. <i>Human Genetics</i> , 1995, 96, 485-9.	3.8	30
108	Further clinical delineation and increased morbidity in males with osteopathia striata with cranial sclerosis: An X-linked disorder?. <i>American Journal of Medical Genetics Part A</i> , 1997, 70, 159-165.	2.4	30

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109	Patient Genotypes Impact Survival After Surgery for Isolated Congenital Heart Disease. <i>Annals of Thoracic Surgery</i> , 2014, 98, 104-111.	1.3	30
110	22q and two: 22q11.2 deletion syndrome and coexisting conditions. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2203-2214.	1.2	30
111	Missense Variants in the Histone Acetyltransferase Complex Component Gene TRRAP Cause Autism and Syndromic Intellectual Disability. <i>American Journal of Human Genetics</i> , 2019, 104, 530-541.	6.2	30
112	Effects of copy number variations on brain structure and risk for psychiatric illness: Large-scale studies from the ENIGMA working groups on CNVs. <i>Human Brain Mapping</i> , 2022, 43, 300-328.	3.6	30
113	Early ultrasound diagnosis of Neuroaxova syndrome. <i>Prenatal Diagnosis</i> , 2001, 21, 575-580.	2.3	29
114	Supernumerary inv dup(15) in a patient with Angelman syndrome and a deletion of 15q11-q13. <i>American Journal of Medical Genetics Part A</i> , 1995, 57, 61-65.	2.4	28
115	Kabuki syndrome is not caused by a microdeletion in the DiGeorge/Velocardiofacial chromosomal region within 22q11.2. , 1996, 65, 101-103.		28
116	The Role of mGluR Copy Number Variation in Genetic and Environmental Forms of Syndromic Autism Spectrum Disorder. <i>Scientific Reports</i> , 2016, 6, 19372.	3.3	28
117	Novel findings with reassessment of exome data: implications for validation testing and interpretation of genomic data. <i>Genetics in Medicine</i> , 2018, 20, 329-336.	2.4	28
118	Tracheal cartilaginous sleeves in children with syndromic craniosynostosis. <i>Genetics in Medicine</i> , 2017, 19, 62-68.	2.4	27
119	Disruption of the blood-brain barrier in 22q11.2 deletion syndrome. <i>Brain</i> , 2021, 144, 1351-1360.	7.6	27
120	Expanding the <i>SPECC1L</i> mutation phenotypic spectrum to include Teebi hypertelorism syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2497-2502.	1.2	26
121	Exome sequencing expands the mechanism of SOX5-associated intellectual disability: A case presentation with review of sox-related disorders. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2548-2554.	1.2	26
122	Utility of genetic evaluation in infants with congenital heart defects admitted to the cardiac intensive care unit. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 3090-3097.	1.2	26
123	Thrombocytopenia with absent radius in a boy and his uncle. <i>American Journal of Medical Genetics Part A</i> , 1987, 28, 117-123.	2.4	25
124	Unusual craniofacial dysmorphism due to prenatal alcohol and cocaine exposure. <i>Teratology</i> , 1994, 50, 160-164.	1.6	25
125	Further delineation of the clinical spectrum of KAT6B disorders and allelic series of pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1338-1347.	2.4	25
126	A second cohort of CHD3 patients expands the molecular mechanisms known to cause Snijders Blok-Campeau syndrome. <i>European Journal of Human Genetics</i> , 2020, 28, 1422-1431.	2.8	25

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127	Association of Mitochondrial Biogenesis With Variable Penetrance of Schizophrenia. <i>JAMA Psychiatry</i> , 2021, 78, 911.	11.0	25
128	B cell development in chromosome 22q11.2 deletion syndrome. <i>Clinical Immunology</i> , 2016, 163, 1-9.	3.2	24
129	Phenotypic spectrum associated with SPECC1L pathogenic variants: new families and critical review of the nosology of Teebi, Opitz GBBB, and Baraitser-Winter syndromes. <i>European Journal of Medical Genetics</i> , 2019, 62, 103588.	1.3	24
130	Ocular albinism in a male with del (6)(q13-q15): Candidate region for autosomal recessive ocular albinism?. <i>American Journal of Medical Genetics Part A</i> , 1992, 42, 700-705.	2.4	23
131	Tricho-Rhino-Phalangeal syndrome type II (Langer-Giedion) with persistent cloaca and prune belly sequence in a girl with 8q interstitial deletion. <i>American Journal of Medical Genetics Part A</i> , 1992, 44, 790-794.	2.4	23
132	Mosaicism for a chromosome 8â€derived minute marker chromosome in a patient with manifestations of trisomy 8 mosaicism. <i>American Journal of Medical Genetics Part A</i> , 1995, 56, 22-24.	2.4	23
133	Phenotype delineation of <i>ZNF462</i> related syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 2075-2082.	1.2	23
134	Diaphragmatic hernia-exomphalos-hypertelorism syndrome: A new case and further evidence of autosomal recessive inheritance. , 1997, 68, 441-444.		22
135	A diagnostic approach to identifying submicroscopic 7p21 deletions in Saethre-Chotzen syndrome: Fluorescence in situ hybridization and dosage-sensitive Southern blot analysis. <i>Genetics in Medicine</i> , 2001, 3, 102-108.	2.4	22
136	Boy with bilateral retinoblastoma due to an unusual ring chromosome 13 with activation of a latent centromere. <i>American Journal of Medical Genetics Part A</i> , 2001, 99, 21-28.	2.4	22
137	Performance on a computerized neurocognitive battery in 22q11.2 deletion syndrome: A comparison between US and Israeli cohorts. <i>Brain and Cognition</i> , 2016, 106, 33-41.	1.8	22
138	Genome-Wide Association Study to Find Modifiers for Tetralogy of Fallot in the 22q11.2 Deletion Syndrome Identifies Variants in the <i>GPR98</i> Locus on 5q14.3. <i>Circulation: Cardiovascular Genetics</i> , 2017, 10, .	5.1	22
139	Deletion size analysis of 1680 22q11.2DS subjects identifies a new recombination hotspot on chromosome 22q11.2. <i>Human Molecular Genetics</i> , 2018, 27, 1150-1163.	2.9	22
140	Anatomic Malformations of the Middle and Inner Ear in 22q11.2 Deletion Syndrome: Case Series and Literature Review. <i>American Journal of Neuroradiology</i> , 2018, 39, 928-934.	2.4	22
141	Cerebroâ€costoâ€mandibular syndrome: Clinical, radiological, and genetic findings. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1115-1126.	1.2	21
142	Congenital diaphragmatic hernia in 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 135-142.	1.2	21
143	<i>De novo</i> missense variants in <i>MEIS2</i> recapitulate the microdeletion phenotype of cardiac and palate abnormalities, developmental delay, intellectual disability and dysmorphic features. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1845-1851.	1.2	21
144	Nonreentrant atrial tachycardia occurs independently of hypertrophic cardiomyopathy in RASopathy patients. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1711-1722.	1.2	21

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146	Novel truncating mutations in <i>CTNND1</i> cause a dominant craniofacial and cardiac syndrome. <i>Human Molecular Genetics</i> , 2020, 29, 1900-1921.	2.9	21
147	<i>ANKRD11</i> variants: <i>KBG</i> syndrome and beyond. <i>Clinical Genetics</i> , 2021, 100, 187-200.	2.0	21
148	Aglossia with congenital absence of the mandibular rami and other craniofacial abnormalities. <i>American Journal of Medical Genetics Part A</i> , 1988, 31, 161-166.	2.4	20
149	A human case of <i>SLC35A3</i> -related skeletal dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2758-2762.	1.2	20
150	Variable Clinical Manifestations of Xia-Gibbs syndrome: Findings of Consecutively Identified Cases at a Single Children's Hospital. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1890-1896.	1.2	20
151	<i>PCDH19</i> -related epilepsy in a male with Klinefelter syndrome: Additional evidence supporting <i>PCDH19</i> cellular interference disease mechanism. <i>Epilepsy Research</i> , 2018, 145, 89-92.	1.6	20
152	Optical mapping of the 22q11.2DS region reveals complex repeat structures and preferred locations for non-allelic homologous recombination (NAHR). <i>Scientific Reports</i> , 2020, 10, 12235.	3.3	20
153	Recurrence rate for de novo 21q21q translocation Down syndrome: A study of 112 families. <i>American Journal of Medical Genetics Part A</i> , 1984, 17, 523-530.	2.4	19
154	Nasal dimple as part of the 22q11.2 deletion syndrome. , 1997, 69, 290-292.		19
155	Attention deficit hyperactivity disorder symptoms as antecedents of later psychotic outcomes in 22q11.2 deletion syndrome. <i>Schizophrenia Research</i> , 2019, 204, 320-325.	2.0	19
156	Heterotaxia in a fetus with campomelia, cervical lymphocele, polysplenia, and multicystic dysplastic kidneys: Expanding the phenotype of Cumming syndrome. , 1997, 73, 419-424.		18
157	Impairment of different protein domains causes variable clinical presentation within Pitt-Hopkins syndrome and suggests intragenic molecular syndromology of <i>TCF4</i> . <i>European Journal of Medical Genetics</i> , 2017, 60, 565-571.	1.3	18
158	Bi-allelic Loss-of-Function Variants in <i>NUP188</i> Cause a Recognizable Syndrome Characterized by Neurologic, Ocular, and Cardiac Abnormalities. <i>American Journal of Human Genetics</i> , 2020, 106, 623-631.	6.2	18
159	46, XX, 15p+ documented as dup (17p) by fluorescence in situ hybridization. <i>American Journal of Medical Genetics Part A</i> , 1993, 46, 95-97.	2.4	17
160	Aphallia as part of urorectal septum malformation sequence in an infant of a diabetic mother. <i>American Journal of Medical Genetics Part A</i> , 1999, 82, 363-367.	2.4	17
161	Limb body wall complex, amniotic band sequence, or new syndrome caused by mutation in <i>IQ Motif</i> containing <i>K</i> (<i>IQCK</i>)?. <i>Molecular Genetics & Genomic Medicine</i> , 2015, 3, 424-432.	1.2	17
162	White matter microstructural deficits in 22q11.2 deletion syndrome. <i>Psychiatry Research - Neuroimaging</i> , 2017, 268, 35-44.	1.8	17

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164	Neurologic challenges in 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2140-2145.	1.2	17
165	Pathogenic variants in CDC45 on the remaining allele in patients with a chromosome 22q11.2 deletion result in a novel autosomal recessive condition. <i>Genetics in Medicine</i> , 2020, 22, 326-335.	2.4	17
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168	Hyperinsulinemic hypoglycemia in seven patients with de novo <i>NSD1</i> mutations. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 542-551.	1.2	16
169	Anomalies of the genitourinary tract in children with 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 381-385.	1.2	16
170	EP300-related Rubinstein-Taybi syndrome: Highlighted rare phenotypic findings and a genotype-phenotype meta-analysis of 74 patients. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2926-2938.	1.2	16
171	De novo loss-of-function variants in X-linked MED12 are associated with Hardikar syndrome in females. <i>Genetics in Medicine</i> , 2021, 23, 637-644.	2.4	16
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175	Autosomal dominant mannose-binding lectin deficiency is associated with worse neurodevelopmental outcomes after cardiac surgery in infants. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2018, 155, 1139-1147.e2.	0.8	15
176	Phenotypic predictors and final diagnoses in patients referred for RASopathy testing by targeted next-generation sequencing. <i>Genetics in Medicine</i> , 2017, 19, 715-718.	2.4	14
177	Hearing Loss after Cardiac Surgery in Infancy: An Unintended Consequence of Life-Saving Care. <i>Journal of Pediatrics</i> , 2018, 192, 144-151.e1.	1.8	14
178	The dimensional structure of psychopathology in 22q11.2 Deletion Syndrome. <i>Journal of Psychiatric Research</i> , 2017, 92, 124-131.	3.1	13
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180	Novel variants in <i>CDH2</i> are associated with a new syndrome including Peters anomaly. <i>Clinical Genetics</i> , 2020, 97, 502-508.	2.0	13

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183	Fetal akinesia deformation sequence due to a congenital disorder of glycosylation. American Journal of Medical Genetics, Part A, 2015, 167, 2411-2417.	1.2	12
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187	Frontonasal malformation and cloacal exstrophy: A previously unreported association. , 1996, 61, 75-78.		11
188	Arthritis associated with deletion of 22q11.2: More common than previously suspected. , 1997, 71, 488-488.		11
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190	Kabuki syndrome as a cause of non-immune fetal hydrops/ascites. American Journal of Medical Genetics, Part A, 2016, 170, 3333-3337.	1.2	11
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195	Study of carrier frequency of Warsaw breakage syndrome in the Ashkenazi Jewish population and presentation of two cases. American Journal of Medical Genetics, Part A, 2019, 179, 2144-2151.	1.2	10
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200	Non-immune hydrops fetalis associated with impaired fetal movement: A case report and review. <i>American Journal of Medical Genetics Part A</i> , 1994, 53, 251-254.	2.4	9
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204	Bi-allelic variants in the ER quality-control mannosidase gene EDEM3 cause a congenital disorder of glycosylation. <i>American Journal of Human Genetics</i> , 2021, 108, 1342-1349.	6.2	9
205	The Genomics of Congenital Diaphragmatic Hernia: A 10-Year Retrospective Review. <i>Journal of Pediatrics</i> , 2022, 248, 108-113.e2.	1.8	9
206	Genetics etiologies and genotype phenotype correlations in a cohort of individuals with central conducting lymphatic anomaly. <i>European Journal of Human Genetics</i> , 2022, 30, 1022-1028.	2.8	9
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219	Molecular Diagnostic Outcomes from 700 Cases. <i>Journal of Molecular Diagnostics</i> , 2022, 24, 274-286.	2.8	7
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222	A Novel Approach to Dysmorphology to Enhance the Phenotypic Classification of Autism Spectrum Disorder in the Study to Explore Early Development. <i>Journal of Autism and Developmental Disorders</i> , 2019, 49, 2184-2202.	2.7	6
223	Aortic Root Dilation in Patients with 22q11.2 Deletion Syndrome Without Intracardiac Anomalies. <i>Pediatric Cardiology</i> , 2021, 42, 1594-1600.	1.3	6
224	Prenatal diagnosis of mosaicism 46, XX/46, XX, ~21, +t(21q21q). <i>Prenatal Diagnosis</i> , 1984, 4, 73-77.	2.3	5
225	Primary lymphedema and other lymphatic anomalies are associated with 22q11.2 deletion syndrome. <i>European Journal of Medical Genetics</i> , 2018, 61, 411-415.	1.3	5
226	Musical auditory processing, cognition, and psychopathology in 22q11.2 deletion syndrome. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 765-773.	1.7	5
227	Perinatal distress in 1p36 deletion syndrome can mimic hypoxic ischemic encephalopathy. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 1543-1546.	1.2	5
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232	Compound heterozygote <i>CDK5RAP2</i> mutations in a Guatemalan/Honduran child with autosomal recessive primary microcephaly, failure to thrive and speech delay. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1414-1417.	1.2	4
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234	Magnetic resonance angiography (MRA) in preoperative planning for patients with 22q11.2 deletion syndrome undergoing craniofacial and otorhinolaryngologic procedures. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2020, 138, 110236.	1.0	4

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237	Molecular Mechanisms Contributing to the Etiology of Congenital Diaphragmatic Hernia: A Review and Novel Cases. <i>Journal of Pediatrics</i> , 2022, 246, 251-265.e2.	1.8	4
238	Phenotypic modifications of patients with full chromosome aneuploidies and concurrent suspected or confirmed second diagnoses. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2168-2175.	1.2	3
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241	Congenital polyvalvular disease expands the cardiac phenotype of the <i>RAS</i> opathies. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1486-1493.	1.2	3
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247	Increased T-cell counts in patients with 22q11.2 deletion syndrome who have anxiety. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1815-1818.	1.2	2
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252	Cleft palate morphology, genetic etiology, and risk of mortality in infants with Robin sequence. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3694-3700.	1.2	1

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254	Altered functional brain dynamics in chromosome 22q11.2 deletion syndrome during facial affect processing. Molecular Psychiatry, 2022, 27, 1158-1166.	7.9	1
255	The microdeletion syndromes. , 2005, , .		0
256	Cover Image, Volume 170A, Number 5, May 2016. , 2016, 170, i-i.		0
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