

Hİlya Kayserili

List of Publications by Year in descending order

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Version: 2024-02-01

200
papers

15,690
citations

19636

61
h-index

19726

117
g-index

215
all docs

215
docs citations

215
times ranked

22700
citing authors

#	ARTICLE	IF	CITATIONS
1	Expanding the spectrum of syndromic <i>PPP2R3C</i> -related <i>XY</i> gonadal dysgenesis to <i>XX</i> gonadal dysgenesis. <i>Clinical Genetics</i> , 2022, 101, 221-232.	1.0	3
2	Functional loss of ubiquitin-specific protease 14 may lead to a novel distal arthrogryposis phenotype. <i>Clinical Genetics</i> , 2022, 101, 421-428.	1.0	1
3	A New Family with a Novel <i>OTUD6B</i> Mutation: Practicing Whole Exome Sequencing for Antenatal Diagnosis of Tetralogy of Fallot. <i>Molecular Syndromology</i> , 2022, 13, 206-211.	0.3	2
4	Biallelic <i>TERT</i> variant leads to Hoyeraal-Hreidarsson syndrome with additional dyskeratosis congenita findings. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1226-1232.	0.7	2
5	Phenotypic and mutational spectrum of <i>ROR2</i> -related Robinow syndrome. <i>Human Mutation</i> , 2022, 43, 900-918.	1.1	8
6	Evaluation of growth, puberty, osteoporosis, and the response to long-term bisphosphonate therapy in four patients with osteoporosis-pseudoglioma syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2022, , .	0.7	2
7	Human model of <i>IRX5</i> mutations reveals key role for this transcription factor in ventricular conduction. <i>Cardiovascular Research</i> , 2021, 117, 2092-2107.	1.8	17
8	A Micropatterned Human-specific Neuroepithelial Tissue for Modeling Gene and Drug-induced Neurodevelopmental Defects. <i>Advanced Science</i> , 2021, 8, 2001100.	5.6	13
9	Skeletal and molecular findings in 51 Cleidocranial dysplasia patients from Turkey. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2488-2495.	0.7	8
10	Overarching control of autophagy and DNA damage response by CHD6 revealed by modeling a rare human pathology. <i>Nature Communications</i> , 2021, 12, 3014.	5.8	16
11	Loss of <i>C2orf69</i> defines a fatal autoinflammatory syndrome in humans and zebrafish that evokes a glycogen-storage-associated mitochondriopathy. <i>American Journal of Human Genetics</i> , 2021, 108, 1301-1317.	2.6	11
12	ESHG PPC Comments on postmortem use of genetic data for research purposes. <i>European Journal of Human Genetics</i> , 2020, 28, 144-146.	1.4	3
13	A rare cause of chronic hyponatremia in an infant: Questions. <i>Pediatric Nephrology</i> , 2020, 35, 241-242.	0.9	0
14	Heterozygous pathogenic variants in <i>GLI1</i> are a common finding in isolated postaxial polydactyly A/B. <i>Human Mutation</i> , 2020, 41, 265-276.	1.1	6
15	A loss-of-function <i>NUAK2</i> mutation in humans causes anencephaly due to impaired Hippo-YAP signaling. <i>Journal of Experimental Medicine</i> , 2020, 217, .	4.2	25
16	Loss of <i>MTX2</i> causes mandibuloacral dysplasia and links mitochondrial dysfunction to altered nuclear morphology. <i>Nature Communications</i> , 2020, 11, 4589.	5.8	30
17	Characteristic dental pattern with hypodontia and short roots in Fraser syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1681-1689.	0.7	7
18	Clinical exome sequencing in neuromuscular diseases: an experience from Turkey. <i>Neurological Sciences</i> , 2020, 41, 2157-2164.	0.9	10

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19	Loss of PYCR2 Causes Neurodegeneration by Increasing Cerebral Glycine Levels via SHMT2. <i>Neuron</i> , 2020, 107, 82-94.e6.	3.8	30
20	A novel shoulder disability staging system for scapulothoracic arthrodesis in patients with facioscapulohumeral dystrophy. <i>Orthopaedics and Traumatology: Surgery and Research</i> , 2020, 106, 701-707.	0.9	6
21	Cohesin complex-associated holoprosencephaly. <i>Brain</i> , 2019, 142, 2631-2643.	3.7	43
22	European recommendations integrating genetic testing into multidisciplinary management of sudden cardiac death. <i>European Journal of Human Genetics</i> , 2019, 27, 1763-1773.	1.4	78
23	A homozygous pathogenic missense variant broadens the phenotypic and mutational spectrum of CREB3L1-related osteogenesis imperfecta. <i>Human Molecular Genetics</i> , 2019, 28, 1801-1809.	1.4	21
24	Turkish Ectodermal Dysplasia Cohort: From Phenotype to Genotype in 17 Families. <i>Cytogenetic and Genome Research</i> , 2019, 157, 189-196.	0.6	7
25	Mutation spectrum of 260 dystrophinopathy patients from Turkey and important highlights for genetic counseling. <i>Neuromuscular Disorders</i> , 2019, 29, 601-613.	0.3	18
26	TMX2 Is a Crucial Regulator of Cellular Redox State, and Its Dysfunction Causes Severe Brain Developmental Abnormalities. <i>American Journal of Human Genetics</i> , 2019, 105, 1126-1147.	2.6	25
27	Zoledronate-responsive calcitriol-mediated hypercalcemia in a 5-year-old case with squamous cell carcinoma on the background of xeroderma pigmentosum. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2019, 32, 1403-1406.	0.4	3
28	Terminal osseous dysplasia with pigmentary defects (TODPD) in a Turkish girl with new skin findings. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 123-129.	0.7	4
29	Sclerosing bone dysplasias with hallmarks of dysosteosclerosis in four patients carrying mutations in SLC29A3 and TCIRG1. <i>Bone</i> , 2019, 120, 495-503.	1.4	23
30	The ARID1B spectrum in 143 patients: from nonsyndromic intellectual disability to Coffinâ€“Siris syndrome. <i>Genetics in Medicine</i> , 2019, 21, 1295-1307.	1.1	80
31	Recontacting patients in clinical genetics services: recommendations of the European Society of Human Genetics. <i>European Journal of Human Genetics</i> , 2019, 27, 169-182.	1.4	65
32	MAB21L1 loss of function causes a syndromic neurodevelopmental disorder with distinctive cerebellar, ocular, craniofacial and genital features (COFG) <i>Tj ETQq0 0 0 rgBT 10verlock 20 Tf 50 2</i>	1.0	10
33	Variants affecting diverse domains of MEPE are associated with two distinct bone disorders, a craniofacial bone defect and otosclerosis. <i>Genetics in Medicine</i> , 2019, 21, 1199-1208.	1.1	17
34	CLINICAL CLASSIFICATION OF RADIAL RAY DEFECTS AND RESEARCH INTO ETIOPATHOGENESIS. <i>European Oral Research</i> , 2019, 81, 127-138.	0.5	0
35	APPLICATION OF MLPA (MULTIPLEX LIGATION-DEPENDENT PROBE AMPLIFICATION) IN FETUSES WITH AN ABNORMAL SONOGRAM AND NORMAL KARYOTYPE. <i>Ä°stanbul TÄ±p FakÄ°ltesi Dergisi</i> , 2019, 82, 5-11.	0.1	0
36	A biallelic ANTXR1 variant expands the anthrax toxin receptor associated phenotype to tooth agenesis. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1015-1022.	0.7	11

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37	Recontacting or not recontacting? A survey of current practices in clinical genetics centres in Europe. <i>European Journal of Human Genetics</i> , 2018, 26, 946-954.	1.4	33
38	Variants in members of the cadherin-catenin complex, CDH1 and CTNND1, cause blepharocheilodontic syndrome. <i>European Journal of Human Genetics</i> , 2018, 26, 210-219.	1.4	34
39	Whole-Exome Sequencing Identifies Novel Variants for Tooth Agenesis. <i>Journal of Dental Research</i> , 2018, 97, 49-59.	2.5	44
40	The oculoauriculofrontonasal syndrome: Further clinical characterization and additional evidence suggesting a nontraditional mode of inheritance. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2740-2750.	0.7	6
41	Pallister-Killian syndrome: clinical, cytogenetic and molecular findings in 15 cases. <i>Molecular Cytogenetics</i> , 2018, 11, 45.	0.4	18
42	RSPO2 inhibition of RNF43 and ZNRF3 governs limb development independently of LGR4/5/6. <i>Nature</i> , 2018, 557, 564-569.	13.7	141
43	Identification of likely pathogenic and known variants in TSPEAR, LAMB3, BCOR, and WNT10A in four Turkish families with tooth agenesis. <i>Human Genetics</i> , 2018, 137, 689-703.	1.8	24
44	Expanding the phenotypic spectrum of variants in PDE4D/PRKAR1A: from acrodysostosis to acroscaphodysplasia. <i>European Journal of Human Genetics</i> , 2018, 26, 1611-1622.	1.4	18
45	Biallelic loss of human CTNNA2, encoding β -catenin, leads to ARP2/3 complex overactivity and disordered cortical neuronal migration. <i>Nature Genetics</i> , 2018, 50, 1093-1101.	9.4	70
46	A facioscapulohumeralis muscularis dystrophia kezelének multidiszciplináris megközelítése. <i>Ideggyógyászati Szemle</i> , 2018, 71, 337-342.	0.4	0
47	Clinically Distinct Phenotypes of Canavan Disease Correlate with Residual Aspartoacylase Enzyme Activity. <i>Human Mutation</i> , 2017, 38, 524-531.	1.1	18
48	De novo mutations in SMCHD1 cause Bosma arhinia microphthalmia syndrome and abrogate nasal development. <i>Nature Genetics</i> , 2017, 49, 249-255.	9.4	88
49	Biallelic mutations in the 3' exonuclease TOE1 cause pontocerebellar hypoplasia and uncover a role in snRNA processing. <i>Nature Genetics</i> , 2017, 49, 457-464.	9.4	66
50	Homozygous mutation in <i>NUP107</i> leads to microcephaly with steroid-resistant nephrotic condition similar to Galloway-Mowat syndrome. <i>Journal of Medical Genetics</i> , 2017, 54, 399-403.	1.5	62
51	Cleidocranial dysplasia: Clinical, endocrinologic and molecular findings in 15 patients from 11 families. <i>European Journal of Medical Genetics</i> , 2017, 60, 163-168.	0.7	31
52	GLI1 inactivation is associated with developmental phenotypes overlapping with Ellis-van Creveld syndrome. <i>Human Molecular Genetics</i> , 2017, 26, 4556-4571.	1.4	50
53	CDK10 Mutations in Humans and Mice Cause Severe Growth Retardation, Spine Malformations, and Developmental Delays. <i>American Journal of Human Genetics</i> , 2017, 101, 391-403.	2.6	35
54	Loss-of-function mutations in Carboxypeptidase D cause a new syndrome with lymphedema and sensorineural hearing loss. <i>Mechanisms of Development</i> , 2017, 145, S32.	1.7	0

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55	PYCR2 Protects from Neurodegeneration by Controlling Oligodendrocyte Maturation and Glycinemia through SHMT2. <i>Mechanisms of Development</i> , 2017, 145, S116-S117.	1.7	0
56	Clinical delineation of a subtype of frontonasal dysplasia with creased nasal ridge and upper limb anomalies: Report of six unrelated patients. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 3136-3142.	0.7	9
57	Teratogenicity of Antiepileptic Drugs. <i>Clinical Psychopharmacology and Neuroscience</i> , 2017, 15, 19-27.	0.9	30
58	Overlapping SETBP1 gain-of-function mutations in Schinzel-Giedion syndrome and hematologic malignancies. <i>PLoS Genetics</i> , 2017, 13, e1006683.	1.5	35
59	Homozygous, and compound heterozygous mutation in 3 Turkish family with Jervell and Lange-Nielsen syndrome: case reports. <i>BMC Medical Genetics</i> , 2017, 18, 114.	2.1	0
60	Mutations in ACTRT1 and its enhancer RNA elements lead to aberrant activation of Hedgehog signaling in inherited and sporadic basal cell carcinomas. <i>Nature Medicine</i> , 2017, 23, 1226-1233.	15.2	59
61	Imatinib response of gastrointestinal stromal tumor patients with germline mutation on KIT exon 13: A family report. <i>World Journal of Radiology</i> , 2017, 9, 365-370.	0.5	3
62	Mutation Update for Kabuki Syndrome Genes <i>KMT2D</i> and <i>KDM6A</i> and Further Delineation of X-Linked Kabuki Syndrome Subtype 2. <i>Human Mutation</i> , 2016, 37, 847-864.	1.1	134
63	Ungual squamous cell carcinoma in a patient with Mal de Meleda. <i>JDDG - Journal of the German Society of Dermatology</i> , 2016, 14, 514-516.	0.4	1
64	485 Mutations in ACTRT1 and its transcribed non-coding elements lead to aberrant activation of the Hedgehog signaling pathway in inherited and sporadic basal cell carcinomas. <i>Journal of Investigative Dermatology</i> , 2016, 136, S243.	0.3	0
65	Characteristic calcaneal ossification: an additional early radiographic finding in infants with fibrodysplasia ossificans progressiva. <i>Pediatric Radiology</i> , 2016, 46, 1568-1572.	1.1	7
66	An unusual presentation of Kabuki syndrome with orbital cysts, microphthalmia, and cholestasis with bile duct paucity. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 3282-3288.	0.7	16
67	Holt-Oram syndrome because of the novel TBX5 mutation c.481A>C. <i>Clinical Dysmorphology</i> , 2016, 25, 192-194.	0.1	1
68	Mutations in <i>CEP120</i> cause Joubert syndrome as well as complex ciliopathy phenotypes. <i>Journal of Medical Genetics</i> , 2016, 53, 608-615.	1.5	55
69	<i>ALX4</i> related parietal foramina mimicking encephalocele in prenatal period. <i>Prenatal Diagnosis</i> , 2016, 36, 591-593.	1.1	4
70	Microcephaly, dysmorphic features, corneal dystrophy, hairy nipples, underdeveloped labioscrotal folds, and small cerebellum in four patients. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1391-1399.	0.7	5
71	Natural History of Congenital Generalized Lipodystrophy: A Nationwide Study From Turkey. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2016, 101, 2759-2767.	1.8	67
72	Loss of Iroquois homeobox transcription factors 3 and 5 in osteoblasts disrupts cranial mineralization. <i>Bone Reports</i> , 2016, 5, 86-95.	0.2	21

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73	A novel ICK mutation causes ciliary disruption and lethal endocrine-cerebro-osteodysplasia syndrome. <i>Cilia</i> , 2016, 5, 8.	1.8	37
74	Novel <i>DDR2</i> mutation identified by whole exome sequencing in a Moroccan patient with spondyloepiphyseal dysplasia, short limb abnormal calcification type. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 460-465.	0.7	12
75	Responsible implementation of expanded carrier screening. <i>European Journal of Human Genetics</i> , 2016, 24, e1-e12.	1.4	240
76	PIK3CA-associated developmental disorders exhibit distinct classes of mutations with variable expression and tissue distribution. <i>JCI Insight</i> , 2016, 1, .	2.3	134
77	Prenatal ultrasonographic diagnosis of generalized arterial calcification of infancy. <i>Journal of Clinical Ultrasound</i> , 2015, 43, 50-54.	0.4	10
78	Mutations in <i>CDK5</i> & <i>RAP2</i> cause Seckel syndrome. <i>Molecular Genetics & Genomic Medicine</i> , 2015, 3, 467-480.	0.6	55
79	Novel <i>MASP1</i> mutations are associated with an expanded phenotype in 3MC1 syndrome. <i>Orphanet Journal of Rare Diseases</i> , 2015, 10, 128.	1.2	46
80	Sclerosteosis (craniotubular hyperostosis-syndactyly) with complex hyperphalangy of the index finger. <i>Pediatric Radiology</i> , 2015, 45, 1239-1243.	1.1	7
81	De novo mutations in <i>PLXND1</i> and <i>REV3L</i> cause M ^u bius syndrome. <i>Nature Communications</i> , 2015, 6, 7199.	5.8	76
82	A severe collodion phenotype in the newborn period associated with a homozygous missense mutation in <i>ALOX12B</i> . <i>British Journal of Dermatology</i> , 2015, 173, 285-287.	1.4	2
83	A new hereditary congenital facial palsy case supports <i>arg5</i> in HOX-DNA binding domain as possible hot spot for mutations. <i>European Journal of Medical Genetics</i> , 2015, 58, 358-363.	0.7	9
84	<i>DOCK6</i> Mutations Are Responsible for a Distinct Autosomal-Recessive Variant of Adams-Oliver Syndrome Associated with Brain and Eye Anomalies. <i>Human Mutation</i> , 2015, 36, 593-598.	1.1	32
85	Disruptions of Topological Chromatin Domains Cause Pathogenic Rewiring of Gene-Enhancer Interactions. <i>Cell</i> , 2015, 161, 1012-1025.	13.5	1,725
86	<i>DVL1</i> Frameshift Mutations Clustering in the Penultimate Exon Cause Autosomal-Dominant Robinow Syndrome. <i>American Journal of Human Genetics</i> , 2015, 96, 612-622.	2.6	110
87	Biallelic mutations in <i>SNX14</i> cause a syndromic form of cerebellar atrophy and lysosome-autophagosome dysfunction. <i>Nature Genetics</i> , 2015, 47, 528-534.	9.4	111
88	<i>TCTEX1D2</i> mutations underlie Jeune asphyxiating thoracic dystrophy with impaired retrograde intraflagellar transport. <i>Nature Communications</i> , 2015, 6, 7074.	5.8	51
89	Determining the genome-wide kinship coefficient seems unhelpful in distinguishing consanguineous couples with a high versus low risk for adverse reproductive outcome. <i>BMC Medical Genetics</i> , 2015, 16, 50.	2.1	1
90	Points to consider for prioritizing clinical genetic testing services: a European consensus process oriented at accountability for reasonableness. <i>European Journal of Human Genetics</i> , 2015, 23, 729-735.	1.4	26

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91	<i>De novo</i> WNT5A-associated autosomal dominant Robinow syndrome suggests specificity of genotype and phenotype. <i>Clinical Genetics</i> , 2015, 87, 34-41.	1.0	56
92	RAP1-mediated MEK/ERK pathway defects in Kabuki syndrome. <i>Journal of Clinical Investigation</i> , 2015, 125, 3585-3599.	3.9	69
93	Genetic heterogeneity in Cornelia de Lange syndrome (CdLS) and CdLS-like phenotypes with observed and predicted levels of mosaicism. <i>Journal of Medical Genetics</i> , 2014, 51, 659-668.	1.5	141
94	Newly described clinical features in two siblings with MACS syndrome and a novel mutation in RIN2. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 484-489.	0.7	11
95	Skull defects, alopecia, hypertelorism, and notched alae nasi caused by homozygous ALX4 gene mutation. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1322-1327.	0.7	17
96	Extreme Growth Failure is a Common Presentation of Ligase IV Deficiency. <i>Human Mutation</i> , 2014, 35, 76-85.	1.1	74
97	Enamel "Renal" Gingival syndrome, hypodontia, and a novel FAM20A mutation. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 2124-2128.	0.7	21
98	Mild nasal clefting may be predictive for ALX4 heterozygotes. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 2054-2058.	0.7	9
99	A patient with a de-novo deletion 3p25.3 and features overlapping with Rubinstein-Taybi syndrome. <i>Clinical Dysmorphology</i> , 2014, 23, 67-70.	0.1	3
100	Clinicogenetic Study of Turkish Patients With Syndromic Craniosynostosis and Literature Review. <i>Pediatric Neurology</i> , 2014, 50, 482-490.	1.0	14
101	C5orf42 is the major gene responsible for OFD syndrome type VI. <i>Human Genetics</i> , 2014, 133, 367-377.	1.8	71
102	Exome Sequencing Links Corticospinal Motor Neuron Disease to Common Neurodegenerative Disorders. <i>Science</i> , 2014, 343, 506-511.	6.0	466
103	CLP1 Founder Mutation Links tRNA Splicing and Maturation to Cerebellar Development and Neurodegeneration. <i>Cell</i> , 2014, 157, 651-663.	13.5	228
104	Clinical manifestations of 17 patients affected with mucopolysaccharidosis type VI and eight novel ARSB mutations. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1443-1453.	0.7	20
105	Neu-Laxova Syndrome Is a Heterogeneous Metabolic Disorder Caused by Defects in Enzymes of the L-Serine Biosynthesis Pathway. <i>American Journal of Human Genetics</i> , 2014, 95, 285-293.	2.6	110
106	Oral manifestations of 17 patients affected with mucopolysaccharidosis type VI. <i>Journal of Inherited Metabolic Disease</i> , 2014, 37, 263-268.	1.7	32
107	Twins with hereditary sensory and autonomic neuropathy type IV with preserved periodontal sensation. <i>European Journal of Medical Genetics</i> , 2014, 57, 240-246.	0.7	9
108	Mutations of CEP83 Cause Infantile Nephronophthisis and Intellectual Disability. <i>American Journal of Human Genetics</i> , 2014, 94, 905-914.	2.6	90

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109	Evaluation of Clinical Manifestations in Patients with Severe Lymphedema with and without CCBE1 Mutations. <i>Molecular Syndromology</i> , 2013, 4, 107-113.	0.3	45
110	Clinical and mutation data in 12 patients with the clinical diagnosis of Nager syndrome. <i>Human Genetics</i> , 2013, 132, 885-898.	1.8	77
111	Mutations in the Gene Encoding IFT Dynein Complex Component WDR34 Cause Jeune Asphyxiating Thoracic Dystrophy. <i>American Journal of Human Genetics</i> , 2013, 93, 932-944.	2.6	108
112	Genotype-phenotype spectrum of PYCR1-related autosomal recessive cutis laxa. <i>Molecular Genetics and Metabolism</i> , 2013, 110, 352-361.	0.5	57
113	Deficiency for the ER-stress transducer OASIS causes severe recessive osteogenesis imperfecta in humans. <i>Orphanet Journal of Rare Diseases</i> , 2013, 8, 154.	1.2	98
114	Defects in the IFT-B Component IFT172 Cause Jeune and Mainzer-Saldino Syndromes in Humans. <i>American Journal of Human Genetics</i> , 2013, 93, 915-925.	2.6	196
115	Multiple synostoses syndrome in three members of a family displaying a novel mutation in NOGGIN gene. <i>Journal of Plastic, Reconstructive and Aesthetic Surgery</i> , 2013, 66, e287-e289.	0.5	3
116	Mutations in WNT1 Cause Different Forms of Bone Fragility. <i>American Journal of Human Genetics</i> , 2013, 92, 565-574.	2.6	240
117	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
118	Phenotypic spectrum and prevalence of INPP5E mutations in Joubert Syndrome and related disorders. <i>European Journal of Human Genetics</i> , 2013, 21, 1074-1078.	1.4	64
119	Exome sequencing identifies <i>DYNC2H1</i> mutations as a common cause of asphyxiating thoracic dystrophy (Jeune syndrome) without major polydactyly, renal or retinal involvement. <i>Journal of Medical Genetics</i> , 2013, 50, 309-323.	1.5	127
120	Prenatal diagnosis of frontonasal dysplasia with anterior encephalocele. <i>Journal of the Turkish German Gynecology Association</i> , 2013, 14, 50-52.	0.2	7
121	Clinical and Radiographic Features of the Autosomal Recessive form of Brachyolmia Caused by <i>PAPSS2</i> Mutations. <i>Human Mutation</i> , 2013, 34, 1381-1386.	1.1	29
122	Genotypic and phenotypic analysis of 396 individuals with mutations in <i>Sonic Hedgehog</i> . <i>Journal of Medical Genetics</i> , 2012, 49, 473-479.	1.5	67
123	A large duplication involving the IHH locus mimics acrocallosal syndrome. <i>European Journal of Human Genetics</i> , 2012, 20, 639-644.	1.4	14
124	Mutations in ISPD cause Walker-Warburg syndrome and defective glycosylation of Î±-dystroglycan. <i>Nature Genetics</i> , 2012, 44, 581-585.	9.4	191
125	<i>OTX2</i> mutations contribute to the otocephaly-dysgnathia complex. <i>Journal of Medical Genetics</i> , 2012, 49, 373-379.	1.5	58
126	Cardiovascular abnormalities in Williams syndrome: 20 years' experience in Istanbul. <i>Acta Cardiologica</i> , 2012, 67, 649-655.	0.3	12

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127	Further characterization of ATP6V0A2-related autosomal recessive cutis laxa. <i>Human Genetics</i> , 2012, 131, 1761-1773.	1.8	73
128	Down Syndrome Diagnosis Based on Gabor Wavelet Transform. <i>Journal of Medical Systems</i> , 2012, 36, 3205-3213.	2.2	39
129	Mutations in IRX5 impair craniofacial development and germ cell migration via SDF1. <i>Nature Genetics</i> , 2012, 44, 709-713.	9.4	68
130	Mutations in <i>BCKD-kinase</i> Lead to a Potentially Treatable Form of Autism with Epilepsy. <i>Science</i> , 2012, 338, 394-397.	6.0	272
131	A novel c.1255G>T (p.D419Y) mutation in SH3BP2 gene causes cherubism in a Turkish family. <i>Oral Surgery, Oral Medicine, Oral Pathology and Oral Radiology</i> , 2012, 114, e42-e46.	0.2	8
132	Candidate locus analysis for PHACE syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 1363-1367.	0.7	24
133	Multiple supernumerary molars, anterior openbite, and large ear lobules in mucopolysaccharidosis type VI patient. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 1798-1800.	0.7	5
134	Mutations in RIPK4 Cause the Autosomal-Recessive Form of Popliteal Pterygium Syndrome. <i>American Journal of Human Genetics</i> , 2012, 90, 76-85.	2.6	80
135	CantÃ© Syndrome Is Caused by Mutations in ABCC9. <i>American Journal of Human Genetics</i> , 2012, 90, 1094-1101.	2.6	141
136	Mild nasal malformations and parietal foramina caused by homozygous <i>ALX4</i> mutations. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 236-244.	0.7	29
137	Evaluation of coronary artery abnormalities in Williams syndrome patients using myocardial perfusion scintigraphy and CT angiography. <i>Cardiology Journal</i> , 2012, 19, 301-308.	0.5	13
138	CEP152 is a genome maintenance protein disrupted in Seckel syndrome. <i>Nature Genetics</i> , 2011, 43, 23-26.	9.4	201
139	TTC21B contributes both causal and modifying alleles across the ciliopathy spectrum. <i>Nature Genetics</i> , 2011, 43, 189-196.	9.4	326
140	A novel homozygous <i>COL11A2</i> deletion causes a C-terminal protein truncation with incomplete mRNA decay in a Turkish patient. <i>American Journal of Medical Genetics, Part A</i> , 2011, 155, 180-185.	0.7	3
141	Harlequin Ichthyosis. <i>Archives of Dermatology</i> , 2011, 147, 681.	1.7	145
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