

Bertrand Isidor

List of Publications by Year in descending order

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Version: 2024-02-01

209
papers

9,744
citations

44444

50
h-index

64407

83
g-index

222
all docs

222
docs citations

222
times ranked

19024
citing authors

#	ARTICLE	IF	CITATIONS
1	Stankiewicz-Isidor syndrome: expanding the clinical and molecular phenotype. <i>Genetics in Medicine</i> , 2022, 24, 179-191.	1.1	9
2	Rare germline heterozygous missense variants in BRCA1-associated protein 1, BAP1, cause a syndromic neurodevelopmental disorder. <i>American Journal of Human Genetics</i> , 2022, 109, 361-372.	2.6	6
3	Expanding the phenotype of <i>HNRNP1</i> -related neurodevelopmental disorder with emphasis on seizure phenotype and review of literature. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1497-1514.	0.7	6
4	First evidence of <i>SOX2</i> mutations in Peters' anomaly: Lessons from molecular screening of 95 patients. <i>Clinical Genetics</i> , 2022, 101, 494-506.	1.0	9
5	Germline variants in tumor suppressor FBXW7 lead to impaired ubiquitination and a neurodevelopmental syndrome. <i>American Journal of Human Genetics</i> , 2022, 109, 601-617.	2.6	16
6	Possible association of 16p11.2 copy number variation with altered lymphocyte and neutrophil counts. <i>Npj Genomic Medicine</i> , 2022, 7, .	1.7	3
7	<i>SETD1B</i> -associated neurodevelopmental disorder. <i>Journal of Medical Genetics</i> , 2021, 58, 196-204.	1.5	22
8	NEXMIF encephalopathy: an X-linked disorder with male and female phenotypic patterns. <i>Genetics in Medicine</i> , 2021, 23, 363-373.	1.1	28
9	JARID2 haploinsufficiency is associated with a clinically distinct neurodevelopmental syndrome. <i>Genetics in Medicine</i> , 2021, 23, 374-383.	1.1	13
10	Neuropsychological study in 19 French patients with Whitehead-Sutton syndrome and <i>POGZ</i> mutations. <i>Clinical Genetics</i> , 2021, 99, 407-417.	1.0	10
11	Pathogenic variants in THSD4, encoding the ADAMTS-like 6 protein, predispose to inherited thoracic aortic aneurysm. <i>Genetics in Medicine</i> , 2021, 23, 111-122.	1.1	25
12	Comprehensive study of 28 individuals with SIN3A-related disorder underscoring the associated mild cognitive and distinctive facial phenotype. <i>European Journal of Human Genetics</i> , 2021, 29, 625-636.	1.4	17
13	DLG4-related synaptopathy: a new rare brain disorder. <i>Genetics in Medicine</i> , 2021, 23, 888-899.	1.1	16
14	Touch and olfaction/taste differentiate children carrying a 16p11.2 deletion from children with ASD. <i>Molecular Autism</i> , 2021, 12, 8.	2.6	6
15	Letter regarding the article "two girls with short stature, short neck, vertebral anomalies, Sprengel deformity and intellectual disability" (Isidor et al., 2015). <i>European Journal of Medical Genetics</i> , 2021, 64, 104179.	0.7	5
16	Haploinsufficiency of the Sin3/HDAC corepressor complex member SIN3B causes a syndromic intellectual disability/autism spectrum disorder. <i>American Journal of Human Genetics</i> , 2021, 108, 929-941.	2.6	15
17	Neurodevelopmental phenotypes in individuals with pathogenic variants in <i>CHAMP1</i> . <i>Journal of Physical Education and Sports Management</i> , 2021, 7, a006092.	0.5	9
18	Effects of eight neuropsychiatric copy number variants on human brain structure. <i>Translational Psychiatry</i> , 2021, 11, 399.	2.4	18

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19	High rate of hypomorphic variants as the cause of inherited ataxia and related diseases: study of a cohort of 366 families. <i>Genetics in Medicine</i> , 2021, 23, 2160-2170.	1.1	13
20	Integrative approach to interpret DYRK1A variants, leading to a frequent neurodevelopmental disorder. <i>Genetics in Medicine</i> , 2021, 23, 2150-2159.	1.1	21
21	Imatinib, a New Adjuvant Medical Treatment for Multifocal Villonodular Synovitis Associated to Noonan Syndrome: A Case Report and Literature Review. <i>Frontiers in Medicine</i> , 2021, 8, 817873.	1.2	1
22	Increasing knowledge in <i>IGF1R</i> defects: lessons from 35 new patients. <i>Journal of Medical Genetics</i> , 2020, 57, 160-168.	1.5	20
23	Immunopathological manifestations in Kabuki syndrome: a registry study of 177 individuals. <i>Genetics in Medicine</i> , 2020, 22, 181-188.	1.1	30
24	iPSC line derived from a Bloom syndrome patient retains an increased disease-specific sister-chromatid exchange activity.. <i>Stem Cell Research</i> , 2020, 43, 101696.	0.3	4
25	Growth charts in Kabuki syndrome 1. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 446-453.	0.7	7
26	Gain-of-Function MN1 Truncation Variants Cause a Recognizable Syndrome with Craniofacial and Brain Abnormalities. <i>American Journal of Human Genetics</i> , 2020, 106, 13-25.	2.6	25
27	Recurrent arginine substitutions in the <i>ACTG2</i> gene are the primary driver of disease burden and severity in visceral myopathy. <i>Human Mutation</i> , 2020, 41, 641-654.	1.1	27
28	Phenotypic spectrum of <i>TGFB3</i> disease-causing variants in a Dutch-French cohort and first report of a homozygous patient. <i>Clinical Genetics</i> , 2020, 97, 723-730.	1.0	15
29	Genotypic diversity and phenotypic spectrum of infantile liver failure syndrome type 1 due to variants in <i>LARS1</i> . <i>Genetics in Medicine</i> , 2020, 22, 1863-1873.	1.1	19
30	Developmental and epilepsy spectrum of <i>KCNB1</i> encephalopathy with long-term outcome. <i>Epilepsia</i> , 2020, 61, 2461-2473.	2.6	17
31	Ribosomopathies: New Therapeutic Perspectives. <i>Cells</i> , 2020, 9, 2080.	1.8	21
32	RLIM Is a Candidate Dosage-Sensitive Gene for Individuals with Varying Duplications of Xq13, Intellectual Disability, and Distinct Facial Features. <i>American Journal of Human Genetics</i> , 2020, 107, 1157-1169.	2.6	6
33	Next-generation sequencing in a series of 80 fetuses with complex cardiac malformations and/or heterotaxy. <i>Human Mutation</i> , 2020, 41, 2167-2178.	1.1	21
34	Psychosocial Impact of Predictive Genetic Testing in Hereditary Heart Diseases: The PREDICT Study. <i>Journal of Clinical Medicine</i> , 2020, 9, 1365.	1.0	9
35	<i>De novo</i> mutations in the X-linked <i>TFE3</i> gene cause intellectual disability with pigmentary mosaicism and storage disorder-like features. <i>Journal of Medical Genetics</i> , 2020, 57, 808-819.	1.5	11
36	Mutations in the Kinesin-2 Motor <i>KIF3B</i> Cause an Autosomal-Dominant Ciliopathy. <i>American Journal of Human Genetics</i> , 2020, 106, 893-904.	2.6	29

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37	Clinical and Molecular Spectrum of Nonsyndromic Early-Onset Osteoarthritis. <i>Arthritis and Rheumatology</i> , 2020, 72, 1689-1693.	2.9	10
38	De Novo Frameshift Variants in the Neuronal Splicing Factor NOVA2 Result in a Common C-Terminal Extension and Cause a Severe Form of Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2020, 106, 438-452.	2.6	17
39	Pathogenic DDX3X Mutations Impair RNA Metabolism and Neurogenesis during Fetal Cortical Development. <i>Neuron</i> , 2020, 106, 404-420.e8.	3.8	121
40	Primrose syndrome: a phenotypic comparison of patients with a ZBTB20 missense variant versus a 3q13.31 microdeletion including ZBTB20. <i>European Journal of Human Genetics</i> , 2020, 28, 1044-1055.	1.4	4
41	A dominant vimentin variant causes a rare syndrome with premature aging. <i>European Journal of Human Genetics</i> , 2020, 28, 1218-1230.	1.4	23
42	Biological concepts in human sodium channel epilepsies and their relevance in clinical practice. <i>Epilepsia</i> , 2020, 61, 387-399.	2.6	65
43	De Novo SOX6 Variants Cause a Neurodevelopmental Syndrome Associated with ADHD, Craniosynostosis, and Osteochondromas. <i>American Journal of Human Genetics</i> , 2020, 106, 830-845.	2.6	17
44	Abstract 3595: Riboprotein variant and their role in chondrogenic differentiation and osteosarcoma development. , 2020, , .		0
45	A step towards precision medicine in management of severe transient polyhydramnios: <i>MAGED2</i> variant. <i>Journal of Obstetrics and Gynaecology</i> , 2019, 39, 395-397.	0.4	6
46	Oro-dental phenotype in patients with RUNX2 duplication. <i>European Journal of Medical Genetics</i> , 2019, 62, 85-89.	0.7	9
47	Expanding the phenotype of the X-linked BCOR microphthalmia syndromes. <i>Human Genetics</i> , 2019, 138, 1051-1069.	1.8	35
48	Encephalopathies with <i>KCNC1</i> variants: genotype-phenotype-functional correlations. <i>Annals of Clinical and Translational Neurology</i> , 2019, 6, 1263-1272.	1.7	33
49	Genotype/phenotype correlations of childhood-onset congenital sideroblastic anaemia in a European cohort. <i>British Journal of Haematology</i> , 2019, 187, 530-542.	1.2	18
50	<i>SETD2</i> related overgrowth syndrome: Presentation of four new patients and review of the literature. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2019, 181, 509-518.	0.7	24
51	Developmental trajectories of neuroanatomical alterations associated with the 16p11.2 Copy Number Variations. <i>NeuroImage</i> , 2019, 203, 116155.	2.1	9
52	RPL13 Variants Cause Spondyloepimetaphyseal Dysplasia with Severe Short Stature. <i>American Journal of Human Genetics</i> , 2019, 105, 1040-1047.	2.6	17
53	Genetic abnormalities in a large cohort of Coffin-Siris syndrome patients. <i>Journal of Human Genetics</i> , 2019, 64, 1173-1186.	1.1	36
54	Pathogenic variants in <i>USP7</i> cause a neurodevelopmental disorder with speech delays, altered behavior, and neurologic anomalies. <i>Genetics in Medicine</i> , 2019, 21, 1797-1807.	1.1	41

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55	Biallelic pathogenic variants in the lanosterol synthase gene LSS involved in the cholesterol biosynthesis cause alopecia with intellectual disability, a rare recessive neuroectodermal syndrome. <i>Genetics in Medicine</i> , 2019, 21, 2025-2035.	1.1	40
56	Identification of mobile retrocopies during genetic testing: Consequences for routine diagnosis. <i>Human Mutation</i> , 2019, 40, 1993-2000.	1.1	4
57	Pycnodysostosis: Natural history and management guidelines from 27 French cases and a literature review. <i>Clinical Genetics</i> , 2019, 96, 309-316.	1.0	31
58	Novel mutations in NLGN3 causing autism spectrum disorder and cognitive impairment. <i>Human Mutation</i> , 2019, 40, 2021-2032.	1.1	42
59	De novo loss-of-function KCNMA1 variants are associated with a new multiple malformation syndrome and a broad spectrum of developmental and neurological phenotypes. <i>Human Molecular Genetics</i> , 2019, 28, 2937-2951.	1.4	76
60	Variants in MED12L, encoding a subunit of the mediator kinase module, are responsible for intellectual disability associated with transcriptional defect. <i>Genetics in Medicine</i> , 2019, 21, 2713-2722.	1.1	28
61	Clinical, Histopathological, and Molecular Diagnostics in Lethal Lung Developmental Disorders. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 200, 1093-1101.	2.5	47
62	Searching for secondary findings: considering actionability and preserving the right not to know. <i>European Journal of Human Genetics</i> , 2019, 27, 1481-1484.	1.4	13
63	Heterozygous Variants in KMT2E Cause a Spectrum of Neurodevelopmental Disorders and Epilepsy. <i>American Journal of Human Genetics</i> , 2019, 104, 1210-1222.	2.6	56
64	Exome sequencing identifies a novel missense variant in CTSC causing nonsyndromic aggressive periodontitis. <i>Journal of Human Genetics</i> , 2019, 64, 689-694.	1.1	8
65	Population genetic screening: current issues in a European country. <i>European Journal of Human Genetics</i> , 2019, 27, 1321-1323.	1.4	2
66	Biallelic MYORG mutation carriers exhibit primary brain calcification with a distinct phenotype. <i>Brain</i> , 2019, 142, 1573-1586.	3.7	49
67	Variable expressivity of syndromic BMP4-related eye, brain, and digital anomalies: A review of the literature and description of three new cases. <i>European Journal of Human Genetics</i> , 2019, 27, 1379-1388.	1.4	8
68	Bilateral retinoblastoma due to a germline mutation of RB1 in a child with down syndrome. <i>Ophthalmic Genetics</i> , 2019, 40, 86-86.	0.5	1
69	Autosomal recessive Treacher Collins syndrome due to <i>POLR1C</i> mutations: Report of a new family and review of the literature. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 1390-1394.	0.7	19
70	Whole genome paired-end sequencing elucidates functional and phenotypic consequences of balanced chromosomal rearrangement in patients with developmental disorders. <i>Journal of Medical Genetics</i> , 2019, 56, 526-535.	1.5	46
71	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.	2.6	30
72	Acanthosis nigricans, hypochondroplasia, and FGFR 3 mutations: Findings with five new patients, and a review of the literature. <i>Pediatric Dermatology</i> , 2019, 36, 242-246.	0.5	6

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73	Estimating the effect size of the 15Q11.2 BP1â€“BP2 deletion and its contribution to neurodevelopmental symptoms: recommendations for practice. <i>Journal of Medical Genetics</i> , 2019, 56, 701-710.	1.5	43
74	Rare variants in the genetic background modulate cognitive and developmental phenotypes in individuals carrying disease-associated variants. <i>Genetics in Medicine</i> , 2019, 21, 816-825.	1.1	127
75	Clinical and functional characterization of recurrent missense variants implicated in <i>THOC6</i> -related intellectual disability. <i>Human Molecular Genetics</i> , 2019, 28, 952-960.	1.4	23
76	A de novo 2q37.2 deletion encompassing <i>AGAP1</i> and <i>SH3BP4</i> in a patient with autism and intellectual disability. <i>European Journal of Medical Genetics</i> , 2019, 62, 103586.	0.7	12
77	<i>ZMIZ1</i> Variants Cause a Syndromic Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2019, 104, 319-330.	2.6	30
78	Complex Compound Inheritance of Lethal Lung Developmental Disorders Due to Disruption of the <i>TBX-FGF</i> Pathway. <i>American Journal of Human Genetics</i> , 2019, 104, 213-228.	2.6	90
79	Heterozygous loss-of-function variants of <i>MEIS2</i> cause a triad of palatal defects, congenital heart defects, and intellectual disability. <i>European Journal of Human Genetics</i> , 2019, 27, 278-290.	1.4	30
80	<i>KAT6A</i> Syndrome: genotypeâ€“phenotype correlation in 76 patients with pathogenic <i>KAT6A</i> variants. <i>Genetics in Medicine</i> , 2019, 21, 850-860.	1.1	68
81	Clinical Presentation of a Complex Neurodevelopmental Disorder Caused by Mutations in <i>ADNP</i> . <i>Biological Psychiatry</i> , 2019, 85, 287-297.	0.7	108
82	Duplications at 19q13.33 in patients with neurodevelopmental disorders. <i>Neurology: Genetics</i> , 2018, 4, e210.	0.9	4
83	Diagnostic strategy in segmentation defect of the vertebrae: a retrospective study of 73 patients. <i>Journal of Medical Genetics</i> , 2018, 55, 422.2-429.	1.5	14
84	Quantifying the Effects of 16p11.2 Copy Number Variants on Brain Structure: A Multisite Genetic-First Study. <i>Biological Psychiatry</i> , 2018, 84, 253-264.	0.7	56
85	Further delineation of the <i>MECP2</i> duplication syndrome phenotype in 59 French male patients, with a particular focus on morphological and neurological features. <i>Journal of Medical Genetics</i> , 2018, 55, 359-371.	1.5	45
86	A novel mutation in the transmembrane 6 domain of <i>GABBR2</i> leads to a Rettâ€“like phenotype. <i>Annals of Neurology</i> , 2018, 83, 437-439.	2.8	19
87	Rare Coding Variants in <i>ANGPTL6</i> Are Associated with Familial Forms of Intracranial Aneurysm. <i>American Journal of Human Genetics</i> , 2018, 102, 133-141.	2.6	37
88	Dual Molecular Effects of Dominant <i>RORA</i> Mutations Cause Two Variants of Syndromic Intellectual Disability with Either Autism or Cerebellar Ataxia. <i>American Journal of Human Genetics</i> , 2018, 102, 744-759.	2.6	51
89	Familial autosomal dominant severe ankyloglossia with tooth abnormalities. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1614-1617.	0.7	10
90	Familial deep endometriosis: A rare monogenic disease?. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2018, 221, 190-193.	0.5	5

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91	Delineating <i>FOXG1</i> syndrome. <i>Neurology: Genetics</i> , 2018, 4, e281.	0.9	51
92	<i>NBEA</i> : Developmental disease gene with early generalized epilepsy phenotypes. <i>Annals of Neurology</i> , 2018, 84, 788-795.	2.8	44
93	Chromosome 14q32.2 Imprinted Region Disruption as an Alternative Molecular Diagnosis of Silver-Russell Syndrome. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 2436-2446.	1.8	48
94	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
95	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
96	Novel <i>SUZ12</i> mutations in Weaver-like syndrome. <i>Clinical Genetics</i> , 2018, 94, 461-466.	1.0	36
97	Natural history of GATA2 deficiency in a survey of 79 French and Belgian patients. <i>Haematologica</i> , 2018, 103, 1278-1287.	1.7	129
98	New splicing pathogenic variant in EBP causing extreme familial variability of Conradi-Häpple Syndrome. <i>European Journal of Human Genetics</i> , 2018, 26, 1784-1790.	1.4	7
99	Microduplication in the 2p16.1p15 chromosomal region linked to developmental delay and intellectual disability. <i>Molecular Cytogenetics</i> , 2018, 11, 39.	0.4	4
100	De Novo and Inherited Loss-of-Function Variants in TLK2: Clinical and Genotype-Phenotype Evaluation of a Distinct Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2018, 102, 1195-1203.	2.6	37
101	Molecular diagnosis of PIK3CA-related overgrowth spectrum (PROS) in 162 patients and recommendations for genetic testing. <i>Genetics in Medicine</i> , 2017, 19, 989-997.	1.1	90
102	De Novo Disruption of the Proteasome Regulatory Subunit PSMD12 Causes a Syndromic Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2017, 100, 352-363.	2.6	86
103	Barber-Say Syndrome and Ablepharon-Macrostomia Syndrome: A Patient's View. <i>Molecular Syndromology</i> , 2017, 8, 172-178.	0.3	6
104	Understanding the Pathophysiology of Intracranial Aneurysm: The ICAN Project. <i>Neurosurgery</i> , 2017, 80, 621-626.	0.6	22
105	Sex chromosome aneuploidies and copy-number variants: a further explanation for neurodevelopmental prognosis variability?. <i>European Journal of Human Genetics</i> , 2017, 25, 930-934.	1.4	19
106	Biallelic Variants in OTUD6B Cause an Intellectual Disability Syndrome Associated with Seizures and Dysmorphic Features. <i>American Journal of Human Genetics</i> , 2017, 100, 676-688.	2.6	54
107	Genetic and phenotypic dissection of 1q43q44 microdeletion syndrome and neurodevelopmental phenotypes associated with mutations in ZBTB18 and HNRNPU. <i>Human Genetics</i> , 2017, 136, 463-479.	1.8	66
108	Mutations in EBF3 Disturb Transcriptional Profiles and Cause Intellectual Disability, Ataxia, and Facial Dysmorphism. <i>American Journal of Human Genetics</i> , 2017, 100, 117-127.	2.6	62

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109	Novel KCNB1 mutation associated with non-syndromic intellectual disability. <i>Journal of Human Genetics</i> , 2017, 62, 569-573.	1.1	28
110	De Novo Mutations in Protein Kinase Genes CAMK2A and CAMK2B Cause Intellectual Disability. <i>American Journal of Human Genetics</i> , 2017, 101, 768-788.	2.6	136
111	Mutations in GREB1L Cause Bilateral Kidney Agenesis in Humans and Mice. <i>American Journal of Human Genetics</i> , 2017, 101, 803-814.	2.6	76
112	Two novel variants in CNTNAP1 in two siblings presenting with congenital hypotonia and hypomyelinating neuropathy. <i>European Journal of Human Genetics</i> , 2017, 25, 150-152.	1.4	13
113	A framework to identify contributing genes in patients with Phelan-McDermid syndrome. <i>Npj Genomic Medicine</i> , 2017, 2, 32.	1.7	58
114	Novel promoters and coding first exons in DLG2 linked to developmental disorders and intellectual disability. <i>Genome Medicine</i> , 2017, 9, 67.	3.6	29
115	Mutations in signal recognition particle SRP54 cause syndromic neutropenia with Shwachman-Diamond-like features. <i>Journal of Clinical Investigation</i> , 2017, 127, 4090-4103.	3.9	126
116	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
117	Protein-altering MYH3 variants are associated with a spectrum of phenotypes extending to spondylocarpotarsal synostosis syndrome. <i>European Journal of Human Genetics</i> , 2016, 24, 1746-1751.	1.4	21
118	De novo MEIS2 mutation causes syndromic developmental delay with persistent gastro-esophageal reflux. <i>Journal of Human Genetics</i> , 2016, 61, 835-838.	1.1	27
119	Germline De Novo Mutations in GNB1 Cause Severe Neurodevelopmental Disability, Hypotonia, and Seizures. <i>American Journal of Human Genetics</i> , 2016, 98, 1001-1010.	2.6	102
120	Mutations in the HECT domain of NEDD4L lead to AKT-mTOR pathway deregulation and cause periventricular nodular heterotopia. <i>Nature Genetics</i> , 2016, 48, 1349-1358.	9.4	101
121	Large national series of patients with Xq28 duplication involving <i>MECP2</i> : Delineation of brain MRI abnormalities in 30 affected patients. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 116-129.	0.7	19
122	Oncologic Phenotype of Peripheral Neuroblastic Tumors Associated With <i>PHOX2B</i> Non-Polyalanine Repeat Expansion Mutations. <i>Pediatric Blood and Cancer</i> , 2016, 63, 71-77.	0.8	14
123	Mandibular dysostosis without microphthalmia caused by <i>OTX2</i> deletion. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2466-2470.	0.7	4
124	De Novo Truncating Variants in SON Cause Intellectual Disability, Congenital Malformations, and Failure to Thrive. <i>American Journal of Human Genetics</i> , 2016, 99, 720-727.	2.6	45
125	Autosomal-Recessive Mutations in AP3B2, Adaptor-Related Protein Complex 3 Beta 2 Subunit, Cause an Early-Onset Epileptic Encephalopathy with Optic Atrophy. <i>American Journal of Human Genetics</i> , 2016, 99, 1368-1376.	2.6	46
126	Contactin-Associated Protein 1 (<i>CNTNAP1</i>) Mutations Induce Characteristic Lesions of the Paranodal Region. <i>Journal of Neuropathology and Experimental Neurology</i> , 2016, 75, 1155-1159.	0.9	31

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127	Expanding the Phenotype Associated with NAA10-Related N-Terminal Acetylation Deficiency. <i>Human Mutation</i> , 2016, 37, 755-764.	1.1	70
128	De Novo Truncating Mutations in the Kinetochores-Microtubules Attachment Gene <i>CHAMP1</i> Cause Syndromic Intellectual Disability. <i>Human Mutation</i> , 2016, 37, 354-358.	1.1	40
129	The Number of Genomic Copies at the 16p11.2 Locus Modulates Language, Verbal Memory, and Inhibition. <i>Biological Psychiatry</i> , 2016, 80, 129-139.	0.7	78
130	Mutations in <i>RIT1</i> cause Noonan syndrome with possible juvenile myelomonocytic leukemia but are not involved in acute lymphoblastic leukemia. <i>European Journal of Human Genetics</i> , 2016, 24, 1124-1131.	1.4	23
131	The expanding spectrum of <i>COL2A1</i> gene variants IN 136 patients with a skeletal dysplasia phenotype. <i>European Journal of Human Genetics</i> , 2016, 24, 992-1000.	1.4	39
132	Defining the Effect of the 16p11.2 Duplication on Cognition, Behavior, and Medical Comorbidities. <i>JAMA Psychiatry</i> , 2016, 73, 20.	6.0	195
133	Abnormal spindle-like microcephaly-associated (<i>ASPM</i>) mutations strongly disrupt neocortical structure but spare the hippocampus and long-term memory. <i>Cortex</i> , 2016, 74, 158-176.	1.1	32
134	Treacher Collins syndrome: a clinical and molecular study based on a large series of patients. <i>Genetics in Medicine</i> , 2016, 18, 49-56.	1.1	125
135	Axial Spondylometaphyseal Dysplasia Is Caused by <i>C21orf2</i> Mutations. <i>PLoS ONE</i> , 2016, 11, e0150555.	1.1	32
136	Neonatal Marfan Syndrome: Report of a Case with an Inherited Splicing Mutation outside the Neonatal Domain. <i>Molecular Syndromology</i> , 2015, 6, 281-286.	0.3	5
137	Patients with isolated oligo/hypodontia caused by <i>RUNX2</i> duplication. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1386-1390.	0.7	10
138	A <i>de novo</i> <i>ADCY5</i> mutation causes early-onset autosomal dominant chorea and dystonia. <i>Movement Disorders</i> , 2015, 30, 423-427.	2.2	131
139	Epileptic patients with <i>de novo</i> <i>STXBP1</i> mutations: Key clinical features based on 24 cases. <i>Epilepsia</i> , 2015, 56, 1931-1940.	2.6	44
140	Muscle magnetic resonance imaging abnormalities in X-linked myopathy with excessive autophagy. <i>Muscle and Nerve</i> , 2015, 52, 673-680.	1.0	6
141	Single amino acid charge switch defines clinically distinct proline-serine-threonine phosphatase-interacting protein 1 (<i>PSTPIP1</i>)-associated inflammatory diseases. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 1337-1345.	1.5	103
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