

Jeanne Amiel

List of Publications by Year in descending order

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Version: 2024-02-01

200
papers

18,718
citations

14124

69
h-index

15253

130
g-index

205
all docs

205
docs citations

205
times ranked

22175
citing authors

#	ARTICLE	IF	CITATIONS
1	Biallelic alterations in <i>PLXND1</i> cause common arterial trunk and other cardiac malformations in humans. <i>Human Molecular Genetics</i> , 2023, 32, 353-356.	1.4	3
2	Further delineation of auriculocondylar syndrome based on 14 novel cases and reassessment of 25 published cases. <i>Human Mutation</i> , 2022, 43, 582-594.	1.1	6
3	Discovery of a genetic module essential for assigning left-right asymmetry in humans and ancestral vertebrates. <i>Nature Genetics</i> , 2022, 54, 62-72.	9.4	16
4	Suleiman-El-Hattab syndrome: a histone modification disorder caused by TASP1 deficiency. <i>Human Molecular Genetics</i> , 2022, 31, 3083-3094.	1.4	3
5	Heterozygous loss of <i>WBP11</i> function causes multiple congenital defects in humans and mice. <i>Human Molecular Genetics</i> , 2021, 29, 3662-3678.	1.4	14
6	Reply: <i>MN1</i> gene loss-of-function mutation causes cleft palate in a pedigree. <i>Brain</i> , 2021, 144, e19-e19.	3.7	3
7	Impaired eIF5A function causes a Mendelian disorder that is partially rescued in model systems by spermidine. <i>Nature Communications</i> , 2021, 12, 833.	5.8	41
8	Dysregulation of the NRG1/ERBB pathway causes a developmental disorder with gastrointestinal dysmotility in humans. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	24
9	Loss of function mutations in GEMIN5 cause a neurodevelopmental disorder. <i>Nature Communications</i> , 2021, 12, 2558.	5.8	28
10	Heterozygous ANKRD17 loss-of-function variants cause a syndrome with intellectual disability, speech delay, and dysmorphism. <i>American Journal of Human Genetics</i> , 2021, 108, 1138-1150.	2.6	17
11	PRICKLE2 revisited—further evidence implicating PRICKLE2 in neurodevelopmental disorders. <i>European Journal of Human Genetics</i> , 2021, 29, 1235-1244.	1.4	5
12	Size matters: Large copy number losses in Hirschsprung disease patients reveal genes involved in enteric nervous system development. <i>PLoS Genetics</i> , 2021, 17, e1009698.	1.5	14
13	PPA2-associated sudden cardiac death: extending the clinical and allelic spectrum in 20 new families. <i>Genetics in Medicine</i> , 2021, 23, 2415-2425.	1.1	8
14	Genetics of craniofacial malformations. <i>Seminars in Fetal and Neonatal Medicine</i> , 2021, 26, 101290.	1.1	5
15	Quality of life and phonatory and morphological outcomes in cognitively unimpaired adolescents with Pierre Robin sequence: a cross-sectional study of 72 patients. <i>Orphanet Journal of Rare Diseases</i> , 2021, 16, 442.	1.2	6
16	Further delineation of the phenotypic spectrum associated with hemizygous loss-of-function variants in <i>NONO</i> . <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 652-658.	0.7	17
17	Generation of an iPSC line (IMAGINi022-A) from a patient carrying a SOX10 missense mutation and presenting with deafness, depigmentation and progressive neurological impairment. <i>Stem Cell Research</i> , 2020, 48, 101936.	0.3	2
18	Overlapping phenotypes between SHORT and Noonan syndromes in patients with PTPN11 pathogenic variants. <i>Clinical Genetics</i> , 2020, 98, 10-18.	1.0	9

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19	<i>EFTUD2</i> missense variants disrupt protein function and splicing in mandibulofacial dysostosis Guion-Almeida type. <i>Human Mutation</i> , 2020, 41, 1372-1382.	1.1	15
20	Bi-allelic Variations of SMO in Humans Cause a Broad Spectrum of Developmental Anomalies Due to Abnormal Hedgehog Signaling. <i>American Journal of Human Genetics</i> , 2020, 106, 779-792.	2.6	25
21	SMCHD1 is involved in <i>de novo</i> methylation of the <i>DUX4</i> -encoding D4Z4 macrosatellite. <i>Nucleic Acids Research</i> , 2019, 47, 2822-2839.	6.5	39
22	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
23	De Novo Mutations Affecting the Catalytic C β Subunit of PP2A, PPP2CA, Cause Syndromic Intellectual Disability Resembling Other PP2A-Related Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2019, 104, 139-156.	2.6	39
24	ZMIZ1 Variants Cause a Syndromic Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2019, 104, 319-330.	2.6	30
25	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
26	Mandibulofacial dysostosis Guion-Almeida type caused by novel <i>EFTUD2</i> splice site variants in two Asian children. <i>Clinical Dysmorphology</i> , 2018, 27, 31-35.	0.1	21
27	Recessive loss of function PIGN alleles, including an intragenic deletion with founder effect in La R�union Island, in patients with Fryns syndrome. <i>European Journal of Human Genetics</i> , 2018, 26, 340-349.	1.4	27
28	<i>MED13L</i> loss-of-function variants in two patients with syndromic Pierre Robin sequence. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 181-186.	0.7	9
29	NONO Detects the Nuclear HIV Capsid to Promote cGAS-Mediated Innate Immune Activation. <i>Cell</i> , 2018, 175, 488-501.e22.	13.5	154
30	Heterozygous Mutations in TBX1 as a Cause of Isolated Hypoparathyroidism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 4023-4032.	1.8	15
31	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
32	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
33	Mutations in the Spliceosome Component CWC27 Cause Retinal Degeneration with or without Additional Developmental Anomalies. <i>American Journal of Human Genetics</i> , 2017, 100, 592-604.	2.6	61
34	Whole exome sequencing coupled with unbiased functional analysis reveals new Hirschsprung disease genes. <i>Genome Biology</i> , 2017, 18, 48.	3.8	72
35	Kaposi sarcoma, oral malformations, mitral dysplasia, and scoliosis associated with 7q34-q36.3 heterozygous terminal deletion. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 1858-1865.	0.7	4
36	Targeted molecular investigation in patients within the clinical spectrum of Auriculocondylar syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 938-945.	0.7	11

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37	De novo mutations in CBL causing early-onset paediatric moyamoya angiopathy. <i>Journal of Medical Genetics</i> , 2017, 54, 550-557.	1.5	33
38	Microduplication of the ARID1A gene causes intellectual disability with recognizable syndromic features. <i>Genetics in Medicine</i> , 2017, 19, 701-710.	1.1	13
39	Mutations in MDH2, Encoding a Krebs Cycle Enzyme, Cause Early-Onset Severe Encephalopathy. <i>American Journal of Human Genetics</i> , 2017, 100, 151-159.	2.6	63
40	Efficiency of prenatal diagnosis in Pierre Robin sequence. <i>Prenatal Diagnosis</i> , 2017, 37, 1169-1175.	1.1	18
41	A Recurrent De Novo Nonsense Variant in ZSWIM6 Results in Severe Intellectual Disability without Frontonasal or Limb Malformations. <i>American Journal of Human Genetics</i> , 2017, 101, 995-1005.	2.6	23
42	Respiratory and gastrointestinal dysfunctions associated with auriculocondylar syndrome and a homozygous PLCB4 loss-of-function mutation. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1471-1478.	0.7	12
43	<i>Trans</i> -ethnic meta-analysis of genome-wide association studies for Hirschsprung disease. <i>Human Molecular Genetics</i> , 2016, 25, dww333.	1.4	38
44	Biallelic PPA2 Mutations Cause Sudden Unexpected Cardiac Arrest in Infancy. <i>American Journal of Human Genetics</i> , 2016, 99, 666-673.	2.6	39
45	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
46	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
47	The expanding spectrum of COL2A1 gene variants IN 136 patients with a skeletal dysplasia phenotype. <i>European Journal of Human Genetics</i> , 2016, 24, 992-1000.	1.4	39
48	<i>RPL10</i> mutation segregating in a family with X-linked syndromic Intellectual Disability. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1908-1912.	0.7	27
49	Functional Assessment of Disease-Associated Regulatory Variants In Vivo Using a Versatile Dual Colour Transgenesis Strategy in Zebrafish. <i>PLoS Genetics</i> , 2015, 11, e1005193.	1.5	31
50	Novel variants in GNAI3 associated with auriculocondylar syndrome strengthen a common dominant negative effect. <i>European Journal of Human Genetics</i> , 2015, 23, 481-485.	1.4	21
51	Functional Loss of Semaphorin 3C and/or Semaphorin 3D and Their Epistatic Interaction with Ret Are Critical to Hirschsprung Disease Liability. <i>American Journal of Human Genetics</i> , 2015, 96, 581-596.	2.6	118
52	Contiguous mutation syndrome in the era of high-throughput sequencing. <i>Molecular Genetics & Genomic Medicine</i> , 2015, 3, 215-220.	0.6	9
53	Mutations in NONO lead to syndromic intellectual disability and inhibitory synaptic defects. <i>Nature Neuroscience</i> , 2015, 18, 1731-1736.	7.1	65
54	Clinical and molecular delineation of Tetrasomy 9p syndrome: Report of 12 new cases and literature review. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1252-1261.	0.7	20

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55	Mutations in the Endothelin Receptor Type A Cause Mandibulofacial Dysostosis with Alopecia. <i>American Journal of Human Genetics</i> , 2015, 96, 519-531.	2.6	47
56	MMP21 is mutated in human heterotaxy and is required for normal left-right asymmetry in vertebrates. <i>Nature Genetics</i> , 2015, 47, 1260-1263.	9.4	65
57	New insights into genotype-phenotype correlation for GLI3 mutations. <i>European Journal of Human Genetics</i> , 2015, 23, 92-102.	1.4	97
58	Late-onset congenital central hypoventilation syndrome and a rare PHOX2B gene mutation. <i>Sleep and Breathing</i> , 2015, 19, 55-60.	0.9	20
59	Multiple congenital anomalies-intellectual disability (MCA-ID) and neuroblastoma in a patient harboring a de novo 14q23.1q23.3 deletion. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1310-1317.	0.7	9
60	Delineation of EFTUD2 Haploinsufficiency-Related Phenotypes Through a Series of 36 Patients. <i>Human Mutation</i> , 2014, 35, 478-485.	1.1	50
61	Clinical evidence for a mandibular to maxillary transformation in Auriculocondylar syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1850-1853.	0.7	6
62	A Homozygous PDE6D Mutation in Joubert Syndrome Impairs Targeting of Farnesylated INPP5E Protein to the Primary Cilium. <i>Human Mutation</i> , 2014, 35, 137-146.	1.1	113
63	Mutations in the tricarboxylic acid cycle enzyme, aconitase 2, cause either isolated or syndromic optic neuropathy with encephalopathy and cerebellar atrophy. <i>Journal of Medical Genetics</i> , 2014, 51, 834-838.	1.5	80
64	Rubinstein-Taybi syndrome predisposing to non-WNT, non-SHH, group 3 medulloblastoma. <i>Pediatric Blood and Cancer</i> , 2014, 61, 383-386.	0.8	33
65	A syndromic form of Pierre Robin sequence is caused by 5q23 deletions encompassing FBN2 and PHAX. <i>European Journal of Medical Genetics</i> , 2014, 57, 587-595.	0.7	18
66	An unusual cause of fetal hypomobility: congenital central hypoventilation syndrome associated with hirschsprung disease. <i>European Journal of Pediatrics</i> , 2014, 173, 1607-1609.	1.3	3
67	Identification of Novel Craniofacial Regulatory Domains Located far Upstream of SOX9 and Disrupted in Pierre Robin Sequence. <i>Human Mutation</i> , 2014, 35, 1011-1020.	1.1	69
68	Blepharophimosis, short humeri, developmental delay and hirschsprung disease: Expanding the phenotypic spectrum of MED12 mutations. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1821-1825.	0.7	19
69	PIK3R1 Mutations Cause Syndromic Insulin Resistance with Lipoatrophy. <i>American Journal of Human Genetics</i> , 2013, 93, 141-149.	2.6	162
70	Mutations in KCTD1 Cause Scalp-Ear-Nipple Syndrome. <i>American Journal of Human Genetics</i> , 2013, 92, 621-626.	2.6	65
71	Contribution of rare and common variants determine complex diseases-Hirschsprung disease as a model. <i>Developmental Biology</i> , 2013, 382, 320-329.	0.9	119
72	Autonomic dysfunction of glucose homeostasis in congenital central hypoventilation syndrome. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 2013, 102, e178-80.	0.7	15

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73	Further characterization of atypical features in auriculocondylar syndrome caused by recessive <i>PLCB4</i> mutations. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2339-2346.	0.7	22
74	Mutations in Endothelin 1 Cause Recessive Auriculocondylar Syndrome and Dominant Isolated Question-Mark Ears. <i>American Journal of Human Genetics</i> , 2013, 93, 1118-1125.	2.6	59
75	Clinical and Molecular Spectrum of Renal Malformations in Kabuki Syndrome. <i>Journal of Pediatrics</i> , 2013, 163, 742-746.	0.9	27
76	Phenotypic Spectrum of Simpson-Golabi-Behmel Syndrome in a Series of 42 Cases With a Mutation in <i>GPC3</i> and Review of the Literature. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2013, 163, 92-105.	0.7	78
77	Understanding the basis of auriculocondylar syndrome: Insights from human, mouse and zebrafish genetic studies. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2013, 163, 306-317.	0.7	48
78	Finger creases lend a hand in Kabuki syndrome. <i>European Journal of Medical Genetics</i> , 2013, 56, 556-560.	0.7	11
79	NF- κ B signalling requirement for brain myelin formation is shown by genotype/MRI phenotype correlations in patients with Xq28 duplications. <i>European Journal of Human Genetics</i> , 2013, 21, 195-199.	1.4	21
80	FOX gene cluster defects in alveolar capillary dysplasia associated with congenital heart disease. <i>Cardiology in the Young</i> , 2013, 23, 697-704.	0.4	12
81	ZEB2 zinc-finger missense mutations lead to hypomorphic alleles and a mild Mowat-Wilson syndrome. <i>Human Molecular Genetics</i> , 2013, 22, 2652-2661.	1.4	51
82	Heterogeneity of mutational mechanisms and modes of inheritance in auriculocondylar syndrome. <i>Journal of Medical Genetics</i> , 2013, 50, 174-186.	1.5	44
83	Congenital Heart Defects in Patients with Deletions Upstream of <i>SOX9</i> . <i>Human Mutation</i> , 2013, 34, 1628-1631.	1.1	33
84	Contiguous gene deletion of TBX5 and TBX3 leads to a variable phenotype with combined features of Holt-Oram and ulnar-mammary syndromes. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 1797-1802.	0.7	16
85	Contributions of PHOX2B in the Pathogenesis of Hirschsprung Disease. <i>PLoS ONE</i> , 2013, 8, e54043.	1.1	30
86	Chromosome 21 Scan in Down Syndrome Reveals DSCAM as a Predisposing Locus in Hirschsprung Disease. <i>PLoS ONE</i> , 2013, 8, e62519.	1.1	22
87	Male and female differential reproductive rate could explain parental transmission asymmetry of mutation origin in Hirschsprung disease. <i>European Journal of Human Genetics</i> , 2012, 20, 917-920.	1.4	8
88	<i>EFTUD2</i> haploinsufficiency leads to syndromic oesophageal atresia. <i>Journal of Medical Genetics</i> , 2012, 49, 737-746.	1.5	89
89	Intragenic <i>CAMTA1</i> rearrangements cause non-progressive congenital ataxia with or without intellectual disability. <i>Journal of Medical Genetics</i> , 2012, 49, 400-408.	1.5	39
90	Mutations of TSEN and CASK genes are prevalent in pontocerebellar hypoplasias type 2 and 4. <i>Brain</i> , 2012, 135, e199-e199.	3.7	18

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91	MicroRNAs in Genetic Disease: Rethinking the Dosage. <i>Current Gene Therapy</i> , 2012, 12, 292-300.	0.9	10
92	Mutations in the chromatin modifier gene KANSL1 cause the 17q21.31 microdeletion syndrome. <i>Nature Genetics</i> , 2012, 44, 639-641.	9.4	194
93	Two independent de novo mutations as a cause for neurofibromatosis type 1 and Noonan syndrome in a single family. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 2290-2291.	0.7	4
94	Constitutional <i>NRAS</i> mutations are rare among patients with Noonan syndrome or juvenile myelomonocytic leukemia. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 2407-2411.	0.7	20
95	miRNA, Development and Disease. <i>Advances in Genetics</i> , 2012, 80, 1-36.	0.8	60
96	A Human Homeotic Transformation Resulting from Mutations in <i>PLCB4</i> and <i>GNAI3</i> Causes Auriculocondylar Syndrome. <i>American Journal of Human Genetics</i> , 2012, 91, 397.	2.6	1
97	In-Frame Mutations in Exon 1 of <i>SKI</i> Cause Dominant Shprintzen-Goldberg Syndrome. <i>American Journal of Human Genetics</i> , 2012, 91, 950-957.	2.6	95
98	A Human Homeotic Transformation Resulting from Mutations in <i>PLCB4</i> and <i>GNAI3</i> Causes Auriculocondylar Syndrome. <i>American Journal of Human Genetics</i> , 2012, 90, 907-914.	2.6	75
99	Novel comprehensive diagnostic strategy in Pitt-Hopkins syndrome: Clinical score and further delineation of the <i>TCF4</i> mutational spectrum. <i>Human Mutation</i> , 2012, 33, 64-72.	1.1	102
100	Autonomic neurocristopathy-associated mutations in <i>PHOX2B</i> dysregulate <i>Sox10</i> expression. <i>Journal of Clinical Investigation</i> , 2012, 122, 3145-3158.	3.9	89
101	Cis-Regulatory Disruption at the <i>SOX9</i> Locus as a Cause of Pierre Robin Sequence. , 2012, , 123-136.		0
102	<i>RET</i> and <i>GDNF</i> mutations are rare in fetuses with renal agenesis or other severe kidney development defects. <i>Journal of Medical Genetics</i> , 2011, 48, 497-504.	1.5	60
103	Genetic Factors in Isolated and Syndromic Esophageal Atresia. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2011, 52, S6-8.	0.9	22
104	Dissection of the <i>MYCN</i> locus in Feingold syndrome and isolated oesophageal atresia. <i>European Journal of Human Genetics</i> , 2011, 19, 602-606.	1.4	24
105	Only four genes (<i>EDA1</i> , <i>EDAR</i> , <i>EDARADD</i> , and <i>WNT10A</i>) account for 90% of hypohidrotic/anhidrotic ectodermal dysplasia cases. <i>Human Mutation</i> , 2011, 32, 70-72.	1.1	240
106	Germline gain-of-function mutations of <i>ALK</i> disrupt central nervous system development. <i>Human Mutation</i> , 2011, 32, 272-276.	1.1	38
107	Pregnancy in women heterozygous for <i>MCT8</i> mutations: risk of maternal hypothyroxinemia and fetal care. <i>European Journal of Endocrinology</i> , 2011, 164, 309-314.	1.9	19
108	Germline deletion of the miR-17 [~] 1492 cluster causes skeletal and growth defects in humans. <i>Nature Genetics</i> , 2011, 43, 1026-1030.	9.4	275

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127	A human mutation in Phox2b causes lack of CO ₂ chemosensitivity, fatal central apnea, and specific loss of parafacial neurons. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 1067-1072.	3.3	271
128	<i>PHOX2B</i> Germline and Somatic Mutations in Late-Onset Central Hypoventilation Syndrome. American Journal of Respiratory and Critical Care Medicine, 2008, 177, 906-911.	2.5	85
129	Delineation of Late Onset Hypoventilation Associated with Hypothalamic Dysfunction Syndrome. Pediatric Research, 2008, 64, 689-694.	1.1	63
130	Hirschsprung disease, associated syndromes and genetics: a review. Journal of Medical Genetics, 2007, 45, 1-14.	1.5	848
131	Methylation-associated PHOX2B gene silencing is a rare event in human neuroblastoma. European Journal of Cancer, 2007, 43, 2366-2372.	1.3	20
132	Mutations in TCF4, Encoding a Class I Basic Helix-Loop-Helix Transcription Factor, Are Responsible for Pitt-Hopkins Syndrome, a Severe Epileptic Encephalopathy Associated with Autonomic Dysfunction. American Journal of Human Genetics, 2007, 80, 988-993.	2.6	264
133	Deletions at the SOX10 Gene Locus Cause Waardenburg Syndrome Types 2 and 4. American Journal of Human Genetics, 2007, 81, 1169-1185.	2.6	216
134	Polyalanine expansions might not result from unequal crossing-over. Human Mutation, 2007, 28, 1043-1044.	1.1	19
135	Homozygous silencing of T-box transcription factor EOMES leads to microcephaly with polymicrogyria and corpus callosum agenesis. Nature Genetics, 2007, 39, 454-456.	9.4	181
136	An overview of isolated and syndromic oesophageal atresia. Clinical Genetics, 2007, 71, 392-399.	1.0	70
137	Polymorphic length of FOXE1 alanine stretch: evidence for genetic susceptibility to thyroid dysgenesis. Human Genetics, 2007, 122, 467-476.	1.8	61
138	Congenital pontocerebellar atrophy and telencephalic defects in three siblings: a new subtype. Acta Neuropathologica, 2007, 114, 387-399.	3.9	7
139	Loss-of-Function Mutations in Euchromatin Histone Methyl Transferase 1 (EHMT1) Cause the 9q34 Subtelomeric Deletion Syndrome. American Journal of Human Genetics, 2006, 79, 370-377.	2.6	343
140	Array-based comparative genomic hybridisation identifies high frequency of cryptic chromosomal rearrangements in patients with syndromic autism spectrum disorders. Journal of Medical Genetics, 2006, 43, 843-849.	1.5	267
141	Molecular Bases of Human Neurocristopathies. , 2006, 589, 213-234.		79
142	The French Congenital Central Hypoventilation Syndrome Registry. Chest, 2005, 127, 72-79.	0.4	199
143	Autonomic Function in Children With Congenital Central Hypoventilation Syndrome and Their Families. Chest, 2005, 128, 2478-2484.	0.4	42
144	Failure to detect an 8p22-8p23.1 duplication in patients with Kabuki (Niikawa-Kuroki) syndrome. European Journal of Human Genetics, 2005, 13, 690-693.	1.4	28

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145	Adams-Oliver syndrome and hepatoportal sclerosis: Occasional association or common mechanism?. American Journal of Medical Genetics, Part A, 2005, 135A, 186-189.	0.7	52
146	Molecular consequences of PHOX2B missense, frameshift and alanine expansion mutations leading to autonomic dysfunction. Human Molecular Genetics, 2005, 14, 3697-3708.	1.4	135
147	CHARGE Syndrome Includes Hypogonadotropic Hypogonadism and Abnormal Olfactory Bulb Development. Journal of Clinical Endocrinology and Metabolism, 2005, 90, 5621-5626.	1.8	142
148	PHOX2B Genotype Allows for Prediction of Tumor Risk in Congenital Central Hypoventilation Syndrome. American Journal of Human Genetics, 2005, 76, 421-426.	2.6	222
149	Germline mutations of the paired-like homeobox 2B (PHOX2B) gene in neuroblastoma. Cancer Letters, 2005, 228, 51-58.	3.2	63
150	Phenotypic spectrum of CHARGE syndrome in fetuses with CHD7 truncating mutations correlates with expression during human development. Journal of Medical Genetics, 2005, 43, 211-317.	1.5	199
151	Genetics and Early Disturbances of Breathing Control: The Genetics of Childhood Disease and Development: A Series of Review Articles. Pediatric Research, 2004, 55, 729-733.	1.1	64
152	Polyalanine expansions in human. Human Molecular Genetics, 2004, 13, R235-R243.	1.4	131
153	Molecular screening of the ZFX1B gene in prenatally diagnosed isolated agenesis of the corpus callosum. Prenatal Diagnosis, 2004, 24, 298-301.	1.1	9
154	PHOX2B gene mutation in a patient with late-onset central hypoventilation. Pediatric Pulmonology, 2004, 38, 349-351.	1.0	50
155	Germline Mutations of the Paired-Like Homeobox 2B (PHOX2B) Gene in Neuroblastoma. American Journal of Human Genetics, 2004, 74, 761-764.	2.6	288
156	Lamin A Truncation in Hutchinson-Gilford Progeria. Science, 2003, 300, 2055-2055.	6.0	1,247
157	Exclusion of RNX as a major gene in congenital central hypoventilation syndrome (CCHS, Ondine's curse). Tj ETQq1 1 0.784314 rgBT / Overlo 2.4 16	2.4	16
158	Polyalanine expansion and frameshift mutations of the paired-like homeobox gene PHOX2B in congenital central hypoventilation syndrome. Nature Genetics, 2003, 33, 459-461.	9.4	771
159	Antenatal manifestations of mitochondrial respiratory chain deficiency. Journal of Pediatrics, 2003, 143, 208-212.	0.9	129
160	Noradrenergic neuronal development is impaired by mutation of the proneural HASH-1 gene in congenital central hypoventilation syndrome (Ondine's curse). Human Molecular Genetics, 2003, 12, 3173-3180.	1.4	72
161	Spectrum of NSD1 mutations in Sotos and Weaver syndromes. Journal of Medical Genetics, 2003, 40, 436-440.	1.5	116
162	A genomic rearrangement resulting in a tandem duplication is associated with split hand-split foot malformation 3 (SHFM3) at 10q24. Human Molecular Genetics, 2003, 12, 1959-1971.	1.4	88

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163	Expression of the SMADIP1 gene during early human development. <i>Mechanisms of Development</i> , 2002, 114, 187-191.	1.7	49
164	Facial appearance in persistent hyperinsulinemic hypoglycemia. <i>American Journal of Medical Genetics Part A</i> , 2002, 111, 130-133.	2.4	29
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