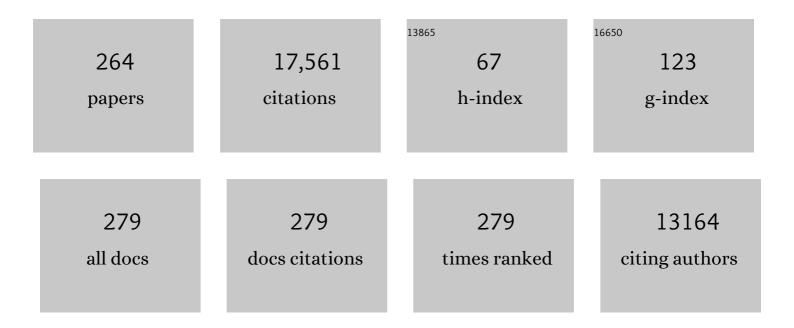
Maximilian Muenke

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Mutations in the human Sonic Hedgehog gene cause holoprosencephaly. Nature Genetics, 1996, 14, 357-360.	21.4	1,075
2	A common mutation in the fibroblast growth factor receptor 1 gene in Pfeiffer syndrome. Nature Genetics, 1994, 8, 269-274.	21.4	615
3	Holoprosencephaly due to mutations in ZIC2, a homologue of Drosophila odd-paired. Nature Genetics, 1998, 20, 180-183.	21.4	448
4	Mutations in the homeodomain of the human SIX3 gene cause holoprosencephaly. Nature Genetics, 1999, 22, 196-198.	21.4	398
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	Overexpression of an Osteogenic Morphogen in Fibrodysplasia Ossificans Progressiva. New England Journal of Medicine, 1996, 335, 555-561.	27.0	364
7	Familial dementia caused by polymerization of mutant neuroserpin. Nature, 1999, 401, 376-379.	27.8	342
8	Opitz G/BBB syndrome, a defect of midline development, is due to mutations in a new RING finger gene on Xp22. Nature Genetics, 1997, 17, 285-291.	21.4	331
9	Loss-of-function mutations in the EGF-CFC gene CFC1 are associated with human left-right laterality defects. Nature Genetics, 2000, 26, 365-369.	21.4	319
10	Loss-of-function mutations in the human <i>GLI2</i> gene are associated with pituitary anomalies and holoprosencephaly-like features. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 13424-13429.	7.1	313
11	Identical mutations in three different fibroblast growth factor receptor genes in autosomal dominant craniosynostosis syndromes. Nature Genetics, 1996, 14, 174-176.	21.4	306
12	Multiple Hits during Early Embryonic Development: Digenic Diseases and Holoprosencephaly. American Journal of Human Genetics, 2002, 71, 1017-1032.	6.2	293
13	Fibroblast-growth-factor receptor mutations in human skeletal disorders. Trends in Genetics, 1995, 11, 308-313.	6.7	292
14	Genetics of ventral forebrain development and holoprosencephaly. Current Opinion in Genetics and Development, 2000, 10, 262-269.	3.3	245
15	Genomic Screening of Fibroblast Growth-Factor Receptor 2 Reveals a Wide Spectrum of Mutations in Patients with Syndromic Craniosynostosis. American Journal of Human Genetics, 2002, 70, 472-486.	6.2	238
16	Mutations in FGFR1 and FGFR2 cause familial and sporadic Pfeiffer syndrome. Human Molecular Genetics, 1995, 4, 323-328.	2.9	221
17	Mutations in PATCHED-1, the receptor for SONIC HEDGEHOG, are associated with holoprosencephaly. Human Genetics, 2002, 110, 297-301.	3.8	220
18	The molecular genetics of holoprosencephaly. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 52-61.	1.6	220

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19	Central Nervous System and Limb Anomalies in Case Reports of First-Trimester Statin Exposure. New England Journal of Medicine, 2004, 350, 1579-1582.	27.0	210
20	Cbfβ interacts with Runx2 and has a critical role in bone development. Nature Genetics, 2002, 32, 639-644.	21.4	207
21	Young adult outcomes in the followâ€up of the multimodal treatment study of attentionâ€deficit/hyperactivity disorder: symptom persistence, source discrepancy, and height suppression. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2017, 58, 663-678.	5.2	207
22	A functional screen for sonic hedgehog regulatory elements across a 1 Mb interval identifies long-range ventral forebrain enhancers. Development (Cambridge), 2006, 133, 761-772.	2.5	198
23	Mutations in holoprosencephaly. Human Mutation, 2000, 16, 99-108.	2.5	194
24	Holoprosencephaly in RSH/Smith-Lemli-Opitz syndrome: Does abnormal cholesterol metabolism affect the function ofsonic hedgehog?. American Journal of Medical Genetics Part A, 1996, 66, 478-484.	2.4	192
25	Attention-Deficit/Hyperactivity Disorder in a Population Isolate: Linkage to Loci at 4q13.2, 5q33.3, 11q22, and 17p11. American Journal of Human Genetics, 2004, 75, 998-1014.	6.2	192
26	Adverse Birth Outcome Among Mothers With Low Serum Cholesterol. Pediatrics, 2007, 120, 723-733.	2.1	188
27	CFC1 Mutations in Patients with Transposition of the Great Arteries and Double-Outlet Right Ventricle. American Journal of Human Genetics, 2002, 70, 776-780.	6.2	182
28	Regulation of a remote Shh forebrain enhancer by the Six3 homeoprotein. Nature Genetics, 2008, 40, 1348-1353.	21.4	182
29	Mechanistic and epidemiologic considerations in the evaluation of adverse birth outcomes following gestational exposure to statins. American Journal of Medical Genetics Part A, 2004, 131A, 287-298.	2.4	171
30	Metaâ€analysis of genomeâ€wide linkage scans of attention deficit hyperactivity disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 1392-1398.	1.7	160
31	SHH mutation is associated with solitary median maxillary central incisor: A study of 13 patients and review of the literature. American Journal of Medical Genetics Part A, 2001, 102, 1-10.	2.4	157
32	A previously unidentified amino-terminal domain regulates transcriptional activity of wild-type and disease-associated human GLI2. Human Molecular Genetics, 2005, 14, 2181-2188.	2.9	156
33	Reduced NODAL Signaling Strength via Mutation of Several Pathway Members Including FOXH1 Is Linked to Human Heart Defects and Holoprosencephaly. American Journal of Human Genetics, 2008, 83, 18-29.	6.2	153
34	Human developmental disorders and the Sonic hedgehog pathway. Trends in Molecular Medicine, 1998, 4, 343-349.	2.6	152
35	Localization of DNA sequences required for human centromere function through an analysis of rearranged Y chromosomes. Nature Genetics, 1993, 5, 368-375.	21.4	149
36	Agenesis and dysgenesis of the corpus callosum: Clinical, genetic and neuroimaging findings in a series of 41 patients. American Journal of Medical Genetics, Part A, 2008, 146A, 2501-2511.	1.2	148

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37	Association between conformational mutations in neuroserpin and onset and severity of dementia. Lancet, The, 2002, 359, 2242-2247.	13.7	145
38	Analysis of genotype–phenotype correlations in human holoprosencephaly. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 133-141.	1.6	139
39	Novel mutation in sonic hedgehog in nonâ€syndromic colobomatous microphthalmia. American Journal of Medical Genetics Part A, 2003, 116A, 215-221.	2.4	135
40	Holoprosencephaly: from Homer to Hedgehog. Clinical Genetics, 1998, 53, 155-163.	2.0	130
41	Craniosynostosis Syndromes: From Genes to Premature Fusion of Skull Bones. Molecular Genetics and Metabolism, 1999, 68, 139-151.	1.1	123
42	Molecular Mechanisms of Holoprosencephaly. Molecular Genetics and Metabolism, 1999, 68, 126-138.	1.1	121
43	The human osmoregulatory Na+/myo-inositol cotransporter gene (SLC5A3): molecular cloning and localization to chromosome 21. Genomics, 1995, 25, 507-513.	2.9	119
44	Mutations in CDON, Encoding a Hedgehog Receptor, Result in Holoprosencephaly and Defective Interactions with Other Hedgehog Receptors. American Journal of Human Genetics, 2011, 89, 231-240.	6.2	116
45	Title is missing!. Nature, 1999, 401, 376-379.	27.8	113
46	De Novo Mutations in CHD4 , an ATP-Dependent Chromatin Remodeler Gene, Cause an Intellectual Disability Syndrome with Distinctive Dysmorphisms. American Journal of Human Genetics, 2016, 99, 934-941.	6.2	111
47	Increased Prevalence of ADHD in Turner Syndrome with No Evidence of Imprinting Effects. Journal of Pediatric Psychology, 2006, 31, 945-955.	2.1	110
48	Symptomatology of autism spectrum disorder in a population with neurofibromatosis type 1. Developmental Medicine and Child Neurology, 2013, 55, 131-138.	2.1	109
49	Opitz syndrome is genetically heterogeneous, with one locus on Xp22, and a second locus on 22q11.2. Nature Genetics, 1995, 11, 459-461.	21.4	103
50	22q11.2 deletion syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2017, 173, 879-888.	1.2	103
51	Muenke syndrome (FGFR3â€related craniosynostosis): Expansion of the phenotype and review of the literature. American Journal of Medical Genetics, Part A, 2007, 143A, 3204-3215.	1.2	99
52	Physical mapping of the holoprosencephaly critical region on chromosome 7q36. Nature Genetics, 1993, 3, 247-251.	21.4	95
53	Identification of a genetic cause for isolated unilateral coronal synostosis: A unique mutation in the fibroblast growth factor receptor 3. Journal of Pediatrics, 1998, 132, 714-716.	1.8	95
54	A loss-of-function mutation in the CFC domain of TDGF1 is associated with human forebrain defects. Human Genetics, 2002, 110, 422-428.	3.8	93

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55	Sex differences in neutrophil biology modulate response to type I interferons and immunometabolism. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 16481-16491.	7.1	91
56	Copy-Number Variations Involving the IHH Locus Are Associated with Syndactyly and Craniosynostosis. American Journal of Human Genetics, 2011, 88, 70-75.	6.2	89
57	Fibrodysplasia Ossificans Progressiva, a Heritable Disorder of Severe Heterotopic Ossification, Maps to Human Chromosome 4q27-31*. American Journal of Human Genetics, 2000, 66, 128-135.	6.2	88
58	Opitz G/BBB syndrome: Clinical comparisons of families linked to Xp22 and 22Q and a review of the literature. , 1996, 62, 305-317.		81
59	Compound heterozygosity for mutations in <i>PAX6</i> in a patient with complex brain anomaly, neonatal diabetes mellitus, and microophthalmia. American Journal of Medical Genetics, Part A, 2009, 149A, 2543-2546.	1.2	80
60	The mutational spectrum of holoprosencephaly-associated changes within the <i>SHH</i> gene in humans predicts loss-of-function through either key structural alterations of the ligand or its altered synthesis. Human Mutation, 2009, 30, E921-E935.	2.5	77
61	The decision to continue: The experiences and needs of parents who receive a prenatal diagnosis of holoprosencephaly. American Journal of Medical Genetics Part A, 2002, 112, 369-378.	2.4	75
62	Mutations in ZIC2 in human holoprosencephaly: description of a Novel ZIC2 specific phenotype and comprehensive analysis of 157 individuals. Journal of Medical Genetics, 2010, 47, 513-524.	3.2	75
63	Down syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2017, 173, 42-53.	1.2	75
64	Human Enteric Defensin Genes: Chromosomal Map Position and a Model for Possible Evolutionary Relationships. Genomics, 1996, 31, 95-106.	2.9	74
65	Latent Class Subtyping of Attention-Deficit/Hyperactivity Disorder and Comorbid Conditions. Journal of the American Academy of Child and Adolescent Psychiatry, 2008, 47, 797-807.	0.5	73
66	Phenotype of the fibroblast growth factor receptor 2 Ser351Cys mutation: Pfeiffer syndrome type III. American Journal of Medical Genetics Part A, 1998, 78, 356-360.	2.4	72
67	Midline and laterality defects: Left and right meet in the middleâ€. BioEssays, 2001, 23, 888-900.	2.5	70
68	Impact of genetics on the diagnosis and clinical management of syndromic craniosynostoses. Child's Nervous System, 2012, 28, 1447-1463.	1.1	70
69	The 2019 US medical genetics workforce: a focus on clinical genetics. Genetics in Medicine, 2021, 23, 1458-1464.	2.4	70
70	Genetic approaches to understanding brain development: Holoprosencephaly as a model. , 2000, 6, 15-21.		69
71	Attention-Deficit/Hyperactivity Disorder and Comorbid Disruptive Behavior Disorders: Evidence of Pleiotropy and New Susceptibility Loci. Biological Psychiatry, 2007, 61, 1329-1339.	1.3	69
72	Review: Genetics of Attention Deficit/Hyperactivity Disorder. Journal of Pediatric Psychology, 2008, 33, 1085-1099.	2.1	69

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73	Noonan syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2017, 173, 2323-2334.	1.2	68
74	The interplay of genetic and environmental factors in craniofacial morphogenesis: holoprosencephaly and the role of cholesterol. Congenital Anomalies (discontinued), 2003, 43, 1-21.	0.6	67
75	Cumulative ligand activity of NODAL mutations and modifiers are linked to human heart defects and holoprosencephaly. Molecular Genetics and Metabolism, 2009, 98, 225-234.	1.1	67
76	Genotypic and phenotypic analysis of 396 individuals with mutations in <i>Sonic Hedgehog</i> . Journal of Medical Genetics, 2012, 49, 473-479.	3.2	67
77	Attention deficit/hyperactivity disorder (ADHD): Complex phenotype, simple genotype?. Genetics in Medicine, 2004, 6, 1-15.	2.4	65
78	How a Hedgehog might see holoprosencephaly. Human Molecular Genetics, 2003, 12, 15R-25.	2.9	64
79	Opitz G/BBB Syndrome in Xp22: Mutations in the MID1 Gene Cluster in the Carboxy-Terminal Domain. American Journal of Human Genetics, 1998, 63, 703-710.	6.2	63
80	Functional analysis of mutations in TGIF associated with holoprosencephaly. Molecular Genetics and Metabolism, 2007, 90, 97-111.	1.1	63
81	Current recommendations for the molecular evaluation of newly diagnosed holoprosencephaly patients. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 93-101.	1.6	62
82	Truncating loss-of-function mutations of DISP1 contribute to holoprosencephaly-like microform features in humans. Human Genetics, 2009, 125, 393-400.	3.8	61
83	Molecular Characterization of Breakpoints in Patients with Holoprosencephaly and Definition of the HPE2 Critical Region 2p21. Human Molecular Genetics, 1996, 5, 223-229.	2.9	60
84	A common genetic network underlies substance use disorders and disruptive or externalizing disorders. Human Genetics, 2012, 131, 917-929.	3.8	60
85	REPORTS. Human Molecular Genetics, 1994, 3, 2153-2158.	2.9	59
86	Holoprosencephaly as a genetic model for normal craniofacial development. Seminars in Developmental Biology, 1994, 5, 293-301.	1.3	58
87	Mutations in the human SIX3 gene in holoprosencephaly are loss of function. Human Molecular Genetics, 2008, 17, 3919-3928.	2.9	56
88	The full spectrum of holoprosencephaly-associated mutations within the <i>ZIC2</i> gene in humans predicts loss-of-function as the predominant disease mechanism. Human Mutation, 2009, 30, E541-E554.	2.5	56
89	Holoprosencephaly due to numeric chromosome abnormalities. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 146-148.	1.6	56
90	Williams–Beuren syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2018, 176, 1128-1136.	1.2	55

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91	Pathogenic mutations in <i>GLI2</i> cause a specific phenotype that is distinct from holoprosencephaly. Journal of Medical Genetics, 2014, 51, 413-418.	3.2	53
92	The CHD4-related syndrome: a comprehensive investigation of the clinical spectrum, genotype–phenotype correlations, and molecular basis. Genetics in Medicine, 2020, 22, 389-397.	2.4	53
93	The Topographic Organization of Repetitive DNA in the Human Nucleolus. Genomics, 1993, 15, 123-132.	2.9	52
94	Attention-Deficit/Hyperactivity Disorder and Comorbidities in 18 Paisa Colombian Multigenerational Families. Journal of the American Academy of Child and Adolescent Psychiatry, 2004, 43, 1506-1515.	0.5	52
95	Missense substitutions in the GAS1 protein present in holoprosencephaly patients reduce the affinity for its ligand, SHH. Human Genetics, 2012, 131, 301-310.	3.8	52
96	Heterozygous mutations in SIX3 and SHH are associated with schizencephaly and further expand the clinical spectrum of holoprosencephaly. Human Genetics, 2010, 127, 555-561.	3.8	51
97	Holoprosencephaly: A guide to diagnosis and clinical management. Indian Pediatrics, 2011, 48, 457-466.	0.4	51
98	Screening of human <i>LPHN3</i> for variants with a potential impact on ADHD susceptibility. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 11-18.	1.7	49
99	Utilizing prospective sequence analysis of SHH, ZIC2, SIX3 and TGIF in holoprosencephaly probands to describe the parameters limiting the observed frequency of mutant gene×gene interactions. Molecular Genetics and Metabolism, 2012, 105, 658-664.	1.1	48
100	An Ultraconserved Brain-Specific Enhancer Within ADGRL3 (LPHN3) Underpins Attention-Deficit/Hyperactivity Disorder Susceptibility. Biological Psychiatry, 2016, 80, 943-954.	1.3	48
101	Noonan syndrome. American Family Physician, 2014, 89, 37-43.	0.1	48
102	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. American Journal of Medical Genetics Part A, 1991, 41, 548-556.	2.4	47
103	Identification of novel mutations in SHH and ZIC2 in a South American (ECLAMC) population with holoprosencephaly. Human Genetics, 2001, 109, 1-6.	3.8	46
104	Dominant-negative kinase domain mutations in <i>FGFR1</i> can explain the clinical severity of Hartsfield syndrome. Human Molecular Genetics, 2016, 25, 1912-1922.	2.9	46
105	Syndromes associated with holoprosencephaly. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2018, 178, 229-237.	1.6	45
106	Holoprosencephaly: recommendations for diagnosis and management. Current Opinion in Pediatrics, 2010, 22, 687-695.	2.0	44
107	An electronic atlas of human malformation syndromes in diverse populations. Genetics in Medicine, 2016, 18, 1085-1087.	2.4	44

Holoprosencephaly: Molecular study of a California Population. , 2000, 90, 315-319.

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109	Cohesin complex-associated holoprosencephaly. Brain, 2019, 142, 2631-2643.	7.6	43
110	ADHD latent class clusters: DSM-IV subtypes and comorbidity. Psychiatry Research, 2009, 170, 192-198.	3.3	42
111	Cyclopia (synophthalmia) in Smith–Lemli–Opitz syndrome: First reported case and consideration of mechanism. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 142-145.	1.6	40
112	Mutations inSPECC1L, encoding sperm antigen with calponin homology and coiled-coil domains 1-like, are found in some cases of autosomal dominant Opitz G/BBB syndrome. Journal of Medical Genetics, 2015, 52, 104-110.	3.2	40
113	Holoprosencephaly in the genomics era. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2018, 178, 165-174.	1.6	40
114	Cornelia de Lange syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2019, 179, 150-158.	1.2	40
115	SONIC HEDGEHOG mutations causing human holoprosencephaly impair neural patterning activity. Human Genetics, 2003, 113, 170-177.	3.8	38
116	Craniosynostosis and Noonan syndrome with <i>KRAS</i> mutations: Expanding the phenotype with a case report and review of the literature. American Journal of Medical Genetics, Part A, 2015, 167, 2657-2663.	1.2	38
117	Muenke syndrome: An international multicenter natural history study. American Journal of Medical Genetics, Part A, 2016, 170, 918-929.	1.2	37
118	FISH diagnosis of the common 57-kb deletion in CTNS causing cystinosis. Human Genetics, 2004, 115, 510-514.	3.8	36
119	Toward a better understanding of ADHD: LPHN3 gene variants and the susceptibility to develop ADHD. ADHD Attention Deficit and Hyperactivity Disorders, 2010, 2, 139-147.	1.7	36
120	Holoprosencephaly and agnathia spectrum: Presentation of two new patients and review of the literature. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 158-169.	1.6	36
121	A novel <i>SIX3</i> mutation segregates with holoprosencephaly in a large family. American Journal of Medical Genetics, Part A, 2009, 149A, 919-925.	1.2	35
122	Abnormal sterol metabolism in holoprosencephaly. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 102-108.	1.6	35
123	A broad range of ophthalmologic anomalies is part of the holoprosencephaly spectrum. American Journal of Medical Genetics, Part A, 2011, 155, 2713-2720.	1.2	35
124	Influence of a Latrophilin 3 (LPHN3) risk haplotype on event-related potential measures of cognitive response control in attention-deficit hyperactivity disorder (ADHD). European Neuropsychopharmacology, 2013, 23, 458-468.	0.7	35
125	<i> <scp>ADGRL</scp> 3 (<scp>LPHN</scp> 3) </i> variants are associated with a refined phenotype of <scp>ADHD</scp> in the <scp>MTA</scp> study. Molecular Genetics & Genomic Medicine, 2016, 4, 540-547.	1.2	35
126	Additional <i>EFNB1</i> mutations in craniofrontonasal syndrome. American Journal of Medical Genetics, Part A, 2008, 146A, 2008-2012.	1.2	34

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127	Structure of the Human Gene Encoding the Associated Microfibrillar Protein (MFAP1) and Localization to Chromosome 15q15-q21. Genomics, 1994, 23, 443-449.	2.9	33
128	Haploinsufficiency of ZNF462 is associated with craniofacial anomalies, corpus callosum dysgenesis, ptosis, and developmental delay. European Journal of Human Genetics, 2017, 25, 946-951.	2.8	33
129	Reciprocal Mouse and Human Limb Phenotypes Caused by Gain- and Loss-of-Function Mutations Affecting <i>Lmbr1</i> . Genetics, 2001, 159, 715-726.	2.9	32
130	A call for global action for rare diseases in Africa. Nature Genetics, 2020, 52, 21-26.	21.4	31
131	The genetics of addiction. Human Genetics, 2012, 131, 773-777.	3.8	30
132	A CCR4-NOT Transcription Complex, Subunit 1, CNOT1, Variant Associated with Holoprosencephaly. American Journal of Human Genetics, 2019, 104, 990-993.	6.2	30
133	Loss-of-Function Variants in PPP1R12A: From Isolated Sex Reversal to Holoprosencephaly Spectrum and Urogenital Malformations. American Journal of Human Genetics, 2020, 106, 121-128.	6.2	30
134	ARP3β, the gene encoding a new human actin-related protein, is alternatively spliced and predominantly expressed in brain neuronal cells. FEBS Journal, 2000, 267, 2921-2928.	0.2	29
135	Clinical Characteristics of Patients with Unicoronal Synostosis and Mutations of Fibroblast Growth Factor Receptor 3: A Preliminary Report. Plastic and Reconstructive Surgery, 2001, 108, 1849-1854.	1.4	29
136	From the black widow spider to human behavior: Latrophilins, a relatively unknown class of G proteinâ€coupled receptors, are implicated in psychiatric disorders. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2011, 156, 1-10.	1.7	29
137	ADGRL3 (LPHN3) variants predict substance use disorder. Translational Psychiatry, 2019, 9, 42.	4.8	29
138	Fibrodysplasia ossificans progressiva in two half-sisters: Evidence for maternal mosaicism. , 1996, 61, 320-324.		28
139	Mutational analysis of the <i>Sonic Hedgehog</i> gene in 220 newborns with oral clefts in a South American (ECLAMC) populationâ€. American Journal of Medical Genetics Part A, 2002, 108, 12-15.	2.4	28
140	VACTERL Association With High Prenatal Lead Exposure: Similarities to Animal Models of Lead Teratogenicity. Pediatrics, 1991, 87, 390-392.	2.1	28
141	Towards a more representative morphology: clinical and ethical considerations for including diverse populations in diagnostic genetic atlases. Genetics in Medicine, 2016, 18, 1069-1074.	2.4	27
142	Loss of function in <i>ROBO1</i> is associated with tetralogy of Fallot and septal defects. Journal of Medical Genetics, 2017, 54, 825-829.	3.2	27
143	<i>BOC</i> is a modifier gene in holoprosencephaly. Human Mutation, 2017, 38, 1464-1470.	2.5	27
144	Hearing loss in syndromic craniosynostoses: Otologic manifestations and clinical findings. International Journal of Pediatric Otorhinolaryngology, 2014, 78, 2037-2047.	1.0	26

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145	Whole-Exome Sequencing for Diagnosis of Turner Syndrome: Toward Next-Generation Sequencing and Newborn Screening. Journal of Clinical Endocrinology and Metabolism, 2017, 102, 1529-1537.	3.6	26
146	SIX3 mutations with holoprosencephaly. American Journal of Medical Genetics, Part A, 2006, 140A, 2577-2583.	1.2	25
147	Palatal and Oral Manifestations of Muenke Syndrome (FGFR3-Related Craniosynostosis). Journal of Craniofacial Surgery, 2012, 23, 664-668.	0.7	25
148	Common genetic causes of holoprosencephaly are limited to a small set of evolutionarily conserved driver genes of midline development coordinated by TGF-β, hedgehog, and FGF signaling. Human Mutation, 2018, 39, 1416-1427.	2.5	25
149	Human germline hedgehog pathway mutations predispose to fatty liver. Journal of Hepatology, 2017, 67, 809-817.	3.7	24
150	Holoprosencephaly flashcards: A summary for the clinician. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2010, 154C, 3-7.	1.6	23
151	Phenotype delineation of <i>ZNF462</i> related syndrome. American Journal of Medical Genetics, Part A, 2019, 179, 2075-2082.	1.2	23
152	Molecular testing in holoprosencephaly. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2018, 178, 187-193.	1.6	22
153	When to suspect a genetic syndrome. American Family Physician, 2012, 86, 826-33.	0.1	21
154	Craniosynostosis, Philadelphia type: A new autosomal dominant syndrome with sagittal craniosynostosis and syndactyly of the fingers and toes. , 1996, 62, 184-191.		20
155	Phenotype profile of a genetic mouse model for Muenke syndrome. Child's Nervous System, 2012, 28, 1483-1493.	1.1	20
156	Epilepsy in Muenke Syndrome: FGFR3-Related Craniosynostosis. Pediatric Neurology, 2012, 47, 355-361.	2.1	20
157	Analysis of brain metabolism by proton magnetic resonance spectroscopy (1H-MRS) in attention-deficit/hyperactivity disorder suggests a generalized differential ontogenic pattern from controls. ADHD Attention Deficit and Hyperactivity Disorders, 2012, 4, 205-212.	1.7	20
158	Prenatal exposure to pesticides and risk for holoprosencephaly: a case-control study. Environmental Health, 2020, 19, 65.	4.0	20
159	Polymorphisms in the neural nicotinic acetylcholine receptor α4 subunit (CHRNA4) are associated with ADHD in a genetic isolate. ADHD Attention Deficit and Hyperactivity Disorders, 2009, 1, 19-24.	1.7	19
160	Genomics and Epigenomics of Congenital Heart Defects: Expert Review and Lessons Learned in Africa. OMICS A Journal of Integrative Biology, 2018, 22, 301-321.	2.0	18
161	The pit, the cleft and the web. Nature Genetics, 2002, 32, 219-220.	21.4	17
162	Potential cognitive endophenotypes in multigenerational families: segregating ADHD from a genetic isolate. ADHD Attention Deficit and Hyperactivity Disorders, 2011, 3, 291-299.	1.7	17

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163	Limb body wall complex, amniotic band sequence, or new syndrome caused by mutation in <i>IQ Motif containing K</i> (<i>IQCK</i>)?. Molecular Genetics & Genomic Medicine, 2015, 3, 424-432.	1.2	17
164	Diversity and dysmorphology. Current Opinion in Pediatrics, 2019, 31, 702-707.	2.0	17
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