Marcella Devoto

List of Publications by Year
in descending order

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Version: 2024-02-01

137 papers	11,441 citations	50170 46 h-index	30848 102 g-index
143 all docs	143 docs citations	143 times ranked	14761 citing authors

#	Article	IF	Citations
1	Identification of ALK as a major familial neuroblastoma predisposition gene. Nature, 2008, 455, 930-935.	13.7	1,207
2	A common molecular basis for three inherited kidney stone diseases. Nature, 1996, 379, 445-449.	13.7	694
3	Cornelia de Lange syndrome is caused by mutations in NIPBL, the human homolog of Drosophila melanogaster Nipped-B. Nature Genetics, 2004, 36, 631-635.	9.4	642
4	Polymorphisms of Î±-adducin and salt sensitivity in patients with essential hypertension. Lancet, The, 1997, 349, 1353-1357.	6.3	518
5	A genome-wide association study identifies KIAA0350 as a type 1 diabetes gene. Nature, 2007, 448, 591-594.	13.7	497
6	Charcot-Marie-Tooth type 4B is caused by mutations in the gene encoding myotubularin-related protein-2. Nature Genetics, 2000, 25, 17-19.	9.4	462
7	Genome-wide copy number variation study associates metabotropic glutamate receptor gene networks with attention deficit hyperactivity disorder. Nature Genetics, 2012, 44, 78-84.	9.4	334
8	Copy number variation at 1q21.1 associated with neuroblastoma. Nature, 2009, 459, 987-991.	13.7	329
9	Variants ofDENND1BAssociated with Asthma in Children. New England Journal of Medicine, 2010, 362, 36-44.	13.9	306
10	NIPBL Mutational Analysis in 120 Individuals with Cornelia de Lange Syndrome and Evaluation of Genotype-Phenotype Correlations. American Journal of Human Genetics, 2004, 75, 610-623.	2.6	277
11	Integrative genomics identifies LMO1 as a neuroblastoma oncogene. Nature, 2011, 469, 216-220.	13.7	276
12	Chromosome 6p22 Locus Associated with Clinically Aggressive Neuroblastoma. New England Journal of Medicine, 2008, 358, 2585-2593.	13.9	271
13	Common variations in BARD1 influence susceptibility to high-risk neuroblastoma. Nature Genetics, 2009, 41, 718-723.	9.4	266
14	Common variation at 6q16 within HACE1 and LIN28B influences susceptibility to neuroblastoma. Nature Genetics, 2012, 44, 1126-1130.	9.4	231
15	A common mutation in Sardinian autoimmune polyendocrinopathy-candidiasis-ectodermal dystrophy patients. Human Genetics, 1998, 103, 428-434.	1.8	217
16	New Locus for Autosomal Dominant High Myopia Maps to the Long Arm of Chromosome 17. , 2003, 44, 1830.		160
17	Complex genetic signatures in immune cells underlie autoimmunity and inform therapy. Nature Genetics, 2020, 52, 1036-1045.	9.4	153
18	Meta-Analysis of Genome-Wide Scans Provides Evidence for Sex- and Site-Specific Regulation of Bone Mass. Journal of Bone and Mineral Research, 2006, 22, 173-183.	3.1	144

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19	IRAK-M Is Involved in the Pathogenesis of Early-Onset Persistent Asthma. American Journal of Human Genetics, 2007, 80, 1103-1114.	2.6	144
20	Phenotype Restricted Genome-Wide Association Study Using a Gene-Centric Approach Identifies Three Low-Risk Neuroblastoma Susceptibility Loci. PLoS Genetics, 2011, 7, e1002026.	1.5	141
21	Mapping a gene for familial situs abnormalities to human chromosome Xq24-q27.1. Nature Genetics, 1993, 5, 403-407.	9.4	140
22	A Novel Susceptibility Locus for Type 1 Diabetes on Chr12q13 Identified by a Genome-Wide Association Study. Diabetes, 2008, 57, 1143-1146.	0.3	137
23	Evidence From Human and Zebrafish That GPC1 Is a Biliary Atresia Susceptibility Gene. Gastroenterology, 2013, 144, 1107-1115.e3.	0.6	125
24	Identification of a Novel Locus on 2q for Autosomal Dominant High-Grade Myopia. , 2005, 46, 2300.		112
25	Genetic Mapping to 10q23.3-q24.2, in a Large Italian Pedigree, of a New Syndrome Showing Bilateral Cataracts, Gastroesophageal Reflux, and Spastic Paraparesis with Amyotrophy. American Journal of Human Genetics, 1999, 64, 586-593.	2.6	108
26	Recessive and dominant mutations in COL12A1 cause a novel EDS/myopathy overlap syndrome in humans and mice. Human Molecular Genetics, 2014, 23, 2339-2352.	1.4	107
27	Rare Variants in TP53 and Susceptibility to Neuroblastoma. Journal of the National Cancer Institute, 2014, 106, dju047.	3.0	100
28	Exome Sequencing Analysis Reveals Variants in Primary Immunodeficiency Genes in Patients With Very Early Onset Inflammatory Bowel Disease. Gastroenterology, 2015, 149, 1415-1424.	0.6	99
29	Why is the cystic fibrosis gene so frequent?. Human Genetics, 1989, 84, 1-5.	1.8	97
30	Common Variation at BARD1 Results in the Expression of an Oncogenic Isoform That Influences Neuroblastoma Susceptibility and Oncogenicity. Cancer Research, 2012, 72, 2068-2078.	0.4	97
31	Replication of GWAS-identified neuroblastoma risk loci strengthens the role of BARD1 and affirms the cumulative effect of genetic variations on disease susceptibility. Carcinogenesis, 2013, 34, 605-611.	1.3	95
32	Close linkage with the RET protooncogene and boundaries of deletion mutations in autosomal dominant Hirschsprung disease. Human Molecular Genetics, 1993, 2, 1803-1808.	1.4	88
33	Precocious sister chromatid separation (PSCS) in Cornelia de Lange syndrome. American Journal of Medical Genetics, Part A, 2005, 138A, 27-31.	0.7	82
34	Genetic analysis in Italian families with inflammatory bowel disease supports linkage to the IBD1 locus â€“ A GISC study. European Journal of Human Genetics, 1999, 7, 567-573.	1.4	81
35	Common Genetic Variants in NEFL Influence Gene Expression and Neuroblastoma Risk. Cancer Research, 2014, 74, 6913-6924.	0.4	74
36	Search for linkage to schizophrenia on the X and Y chromosomes. American Journal of Medical Genetics Part A, 1994, 54, 113-121.	2.4	66

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37	Genetic modifiers of Â-thalassemia and clinical severity as assessed by age at first transfusion. Haematologica, 2012, 97, 989-993.	1.7	64
38	Genetic linkage study of high-grade myopia in a Hutterite population from South Dakota. Molecular Vision, 2007, 13, 229-36.	1.1	63
39	Copy-Number Variation of the Glucose Transporter Gene SLC2A3 and Congenital Heart Defects in the 22q11.2 Deletion Syndrome. American Journal of Human Genetics, 2015, 96, 753-764.	2.6	62
40	Common variants upstream of MLF1 at 3q25 and within CPZ at 4p16 associated with neuroblastoma. PLoS Genetics, 2017, 13, e1006787.	1.5	62
41	A linkage study of schizophrenia to markers within Xp11 near the MAOB gene. Psychiatry Research, 1997, 70, 131-143.	1.7	60
42	Association of the T300A non-synonymous variant of the ATG16L1 gene with susceptibility to paediatric Crohn's disease. Gut, 2007, 56, 1171-1173.	6.1	60
43	Replication of a GWAS signal in a Caucasian population implicates ADD3 in susceptibility to biliary atresia. Human Genetics, 2014, 133, 235-243.	1.8	59
44	Genotype and cardiovascular phenotype correlations with TBX1 in 1,022 velo-cardio-facial/digeorge/22q11.2 deletion syndrome patients. Human Mutation, 2011, 32, 1278-1289.	1.1	57
45	A Genotype-Phenotype Correlation with Gender-Effect for Hearing Impairment Caused by TECTA Mutations. Cellular Physiology and Biochemistry, 2004, 14, 369-376.	1.1	56
46	Association of Variants of the Interleukin-23 Receptor Gene With Susceptibility to Pediatric Crohnâ€™s Disease. Clinical Gastroenterology and Hepatology, 2007, 5, 972-976.	2.4	56
47	Replication of Neuroblastoma SNP Association at the BARD1 Locus in African-Americans. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 658-663.	1.1	54
48	Fine mapping of 2q35 highâ€risk neuroblastoma locus reveals independent functional risk variants and suggests fullâ€length BARD1 as tumorâ€suppressor. International Journal of Cancer, 2018, 143, 2828-2837.	2.3	54
49	A genome-wide association study identifies a susceptibility locus for biliary atresia on 2p16.1 within the gene EFEMP1. PLoS Genetics, 2018, 14, e1007532.	1.5	51
50	Rare copy number variants in over 100,000 European ancestry subjects reveal multiple disease associations. Nature Communications, 2020, 11, 255.	5.8	48
51	Nocturnal Enuresis: A Suggestive Endophenotype Marker for a Subgroup of Inattentive Attention-Deficit/Hyperactivity Disorder. Journal of Pediatrics, 2009, 155, 239-244.e5.	0.9	47
52	The Otto Aufranc Award: Identification of a 4 Mb Region on Chromosome 17q21 Linked to Developmental Dysplasia of the Hip in One 18-member, Multigeneration Family. Clinical Orthopaedics and Related Research, 2010, 468, 337-344.	0.7	47
53	Developmental Dysplasia of the Hip: Linkage Mapping and Whole Exome Sequencing Identify a Shared Variant in CX 3 CR 1 in All Affected Members of a Large Multigeneration Family. Journal of Bone and Mineral Research, 2013, 28, 2540-2549.	3.1	47
54	A common haplotype at the 5â€² end of the RET protoâ€oncogene, overrepresented in Hirschsprung patients, is associated with reduced gene expression. Human Mutation, 2005, 25, 189-195.	1.1	46

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55	A single-nucleotide polymorphic variant of the RET proto-oncogene is underrepresented in sporadic Hirschsprung disease. European Journal of Human Genetics, 2000, 8, 721-724.	1.4	45
56	Four new mutations of the CFTR gene (541delC, R347H, R352Q, E585X) detected by DGGE analysis in Italian CF patients, associated with different clinical phenotypes. Human Mutation, 1992, 1, 314-319.	1.1	44
57	Gender-dependent disease severity in autosomal polycystic kidney disease of rats. Kidney International, 1995, 48, 496-500.	2.6	44
58	Heterozygous Deletion ofFOXA2Segregates with Disease in a Family with Heterotaxy, Panhypopituitarism, and Biliary Atresia. Human Mutation, 2015, 36, 631-637.	1.1	43
59	Variants of the ACTG2 gene correlate with degree of severity and presence of megacystis in chronic intestinal pseudo-obstruction. European Journal of Human Genetics, 2016, 24, 1211-1215.	1.4	43
60	Rare copy number variants and congenital heart defects in the 22q11.2 deletion syndrome. Human Genetics, 2016, 135, 273-285.	1.8	43
61	A Rare Haplotype of the RET Proto-Oncogene Is a Risk-Modifying Allele in Hirschsprung Disease. American Journal of Human Genetics, 2002, 71, 969-974.	2.6	41
62	Weak linkage at 4p16 to predisposition for human neuroblastoma. Oncogene, 2002, 21, 8356-8360.	2.6	40
63	Univariate and bivariate variance component linkage analysis of a whole-genome scan for loci contributing to bone mineral density. European Journal of Human Genetics, 2005, 13, 781-788.	1.4	38
64	A de novo whole gene deletion of XIAP detected by exome sequencing analysis in very early onset inflammatory bowel disease: a case report. BMC Gastroenterology, 2015, 15, 160.	0.8	38
65	Computerized neurocognitive profile in young people with 22q11.2 deletion syndrome compared to youths with schizophrenia and Atâ€Risk for psychosis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 87-93.	1.1	37
66	Association of a NovelACTA1Mutation With a Dominant Progressive Scapuloperoneal Myopathy in an Extended Family. JAMA Neurology, 2015, 72, 689.	4.5	35
67	THBS2 Is a Candidate Modifier of Liver Disease Severity in Alagille Syndrome. Cellular and Molecular Gastroenterology and Hepatology, 2016, 2, 663-675.e2.	2.3	35
68	Genetic Predisposition to Familial Neuroblastoma: Identification of Two Novel Genomic Regions at 2p and 12p. Human Heredity, 2007, 63, 205-211.	0.4	34
69	ADHD genetics: 2007 update. Current Psychiatry Reports, 2007, 9, 434-439.	2.1	34
70	Linkage analysis conditional on HLA status in a large North American pedigree supports the presence of a multiple sclerosis susceptibility locus on chromosome 12p12. Human Molecular Genetics, 2002, 11, 295-300.	1.4	33
71	Candidate gene analysis in an on-going genome-wide association study of attention-deficit hyperactivity disorder: suggestive association signals in ADRA1A. Psychiatric Genetics, 2009, 19, 134-141.	0.6	33
72	The Unique Disease Course of Children with Very Early onset-Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2020, 26, 909-918.	0.9	32

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73	CARD15 Genotyping in Inflammatory Bowel Disease Patients by Multiplex Pyrosequencing. Clinical Chemistry, 2003, 49, 1675-1679.	1.5	30
74	Heterogeneity in the Magnitude of the Insulin Gene Effect on HLA Risk in Type 1 Diabetes. Diabetes, 2004, 53, 3286-3291.	0.3	30
75	Estimates of Genetic and Environmental Contribution to 43 Quantitative Traits Support Sharing of a Homogeneous Environment in an Isolated Population from South Tyrol, Italy. Human Heredity, 2008, 65, 175-182.	0.4	30
76	Expanding the spectrum of genes responsible for hereditary motor neuropathies. Journal of Neurology, Neurosurgery and Psychiatry, 2019, 90, 1171-1179.	0.9	30
77	Ophthalmologic Findings in Cornelia de Lange Syndrome. JAMA Ophthalmology, 2006, 124, 552.	2.6	29
78	A novel locus for syndromic chronic idiopathic intestinal pseudo-obstruction maps to chromosome 8q23â€“q24. European Journal of Human Genetics, 2007, 15, 889-897.	1.4	29
79	Exclusion of linkage to theCDL1 gene region on chromosome 3q26.3 in some familial cases of Cornelia de Lange syndrome. American Journal of Medical Genetics Part A, 2001, 101, 120-129.	2.4	27
80	Contribution of IBD5 Locus to Clinical Features of IBD Patients. American Journal of Gastroenterology, 2006, 101, 318-325.	0.2	27
81	Analysis of GWAS top hits in ADHD suggests association to two polymorphisms located in genes expressed in the cerebellum. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 1127-1133.	1.1	27
82	Genome-Wide Linkage Analysis to Identify Genetic Modifiers of <i>ALK</i> Mutation Penetrance in Familial Neuroblastoma. Human Heredity, 2011, 71, 135-139.	0.4	27
83	Novel COCH mutation in a family with autosomal dominant late onset sensorineural hearing impairment and tinnitus. American Journal of Otolaryngology - Head and Neck Medicine and Surgery, 2013, 34, 230-235.	0.6	27
84	Five families with arginine519-cysteine mutation in COL2A1: Evidence for three distinct founders. , 1998, 12, 172-176.		26
85	Exclusion of linkage between RET and neuronal intestinal dysplasia type B. , 1996, 62, 195-198.		23
86	Pathway analysis supports association of nonsyndromic cryptorchidism with genetic loci linked to cytoskeleton-dependent functions. Human Reproduction, 2015, 30, 2439-2451.	0.4	23
87	A nonsense mutation (R1158X) and a splicing mutation (3849 + 4A â†' G) in exon 19 of the cystic fibrosis transmembrane conductance regulator gene. Genomics, 1992, 12, 417-418.	1.3	22
88	Compound heterozygous mutations in NEK8 in siblings with endâ€stage renal disease with hepatic and cardiac anomalies. American Journal of Medical Genetics, Part A, 2016, 170, 750-753.	0.7	22
89	Genetic loci linked to Type 1 Diabetes and Multiple Sclerosis families in Sardinia. BMC Medical Genetics, 2008, 9, 3.	2.1	21
90	Overt cleft palate phenotype and TBX1 genotype correlations in veloâ€cardioâ€facial/DiGeorge/22q11.2 deletion syndrome patients. American Journal of Medical Genetics, Part A, 2012, 158A, 2781-2787.	0.7	20

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91	Haplotype distribution and molecular defects at the phenylalanine hydroxylase locus in Italy. Human Genetics, 1990, 86, 69-72.	1.8	19
92	Novel ZBTB24 Mutation Associated with Immunodeficiency, Centromere Instability, and Facial Anomalies Type-2 Syndrome Identified in a Patient with Very Early Onset Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2017, 23, 2252-2255.	0.9	19
93	Combined segregation and linkage analysis of inflammatory bowel disease in the IBD1 region using severity to characterise Crohn's disease and ulcerative colitis. European Journal of Human Genetics, 2000, 8, 846-852.	1.4	18
94	Genetic Refinement and Physical Mapping of the CMT4B Gene on Chromosome 11q22. Genomics, 2000, 63, 271-278.	1.3	18
95	Neural crest-derived tumor neuroblastoma and melanoma share 1p13.2 as susceptibility locus that shows a long-range interaction with the SLC16A1 gene. Carcinogenesis, 2020, 41, 284-295.	1.3	18
96	Frequency of consanguineous marriages among parents and grandparents of Down patients. Human Genetics, 1985, 70, 256-8.	1.8	17
97	A refined physical and transcriptional map of the SPG9 locus on 10q23.3â€“q24.2. European Journal of Human Genetics, 2000, 8, 777-782.	1.4	17
98	Linkage Analysis in Families with Recurrent Neuroblastoma. Annals of the New York Academy of Sciences, 2002, 963, 74-84.	1.8	17
99	Phenotype Specific Association of the TGFBR3 Locus with Nonsyndromic Cryptorchidism. Journal of Urology, 2015, 193, 1637-1645.	0.2	17
100	Exome Sequencing in Individuals with Isolated Biliary Atresia. Scientific Reports, 2020, 10, 2709.	1.6	17
101	Novel X-linked mental retardation syndrome with short stature maps to Xq24. American Journal of Medical Genetics Part A, 2001, 103, 1-8.	2.4	14
102	Exploring Shared Susceptibility between Two Neural Crest Cells Originating Conditions: Neuroblastoma and Congenital Heart Disease. Genes, 2019, 10, 663.	1.0	14
103	A candidate gene approach to identify modifiers of the palatal phenotype in 22q11.2 deletion syndrome patients. International Journal of Pediatric Otorhinolaryngology, 2013, 77, 123-127.	0.4	13
104	Second family with hearing impairment linked to 19q13 and refined DFNA4 localisation. European Journal of Human Genetics, 2002, 10, 95-99.	1.4	12
105	Statistical tools for linkage analysis and genetic association studies. Expert Review of Molecular Diagnostics, 2005, 5, 781-796.	1.5	11
106	Genetic differences in cystic fibrosis patients with and without pancreatic insufficiency. Human Genetics, 1990, 84, 435-8.	1.8	10
107	Variants of ST8SIA1 Are Associated with Risk of Developing Multiple Sclerosis. PLoS ONE, 2008, 3, e2653.	1.1	10
108	Subphenotype meta-analysis of testicular cancer genome-wide association study data suggests a role for RBFOX family genes in cryptorchidism susceptibility. Human Reproduction, 2018, 33, 967-977.	0.4	10

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109	Frequency of the Î”F508 mutation in a sample of 175 Italian cystic fibrosis patients. Human Genetics, 1990, 85, 400-402.	1.8	9
110	Analysis of linkage disequilibrium between different cystic fibrosis mutations and three intragenic microsatellites in the Italian population. Human Mutation, 1995, 5, 23-27.	1.1	9
111	Exome sequencing reveals compound heterozygous mutations in ATP8B1 in a JAG1/NOTCH2 mutationâ€negative patient with clinically diagnosed Alagille syndrome. American Journal of Medical Genetics, Part A, 2015, 167, 891-893.	0.7	9
112	Polygenic inheritance of cryptorchidism susceptibility in the LE/orl rat. Molecular Human Reproduction, 2016, 22, 18-34.	1.3	9
113	Impaired Redox and Protein Homeostasis as Risk Factors and Therapeutic Targets in Toxin-Induced Biliary Atresia. Gastroenterology, 2020, 159, 1068-1084.e2.	0.6	9
114	Colonoids From Patients With Pediatric Inflammatory Bowel Disease Exhibit Decreased Growth Associated With Inflammation Severity and Durable Upregulation of Antigen Presentation Genes. Inflammatory Bowel Diseases, 2021, 27, 256-267.	0.9	7
115	Variants in STXBP3 are Associated with Very Early Onset Inflammatory Bowel Disease, Bilateral Sensorineural Hearing Loss and Immune Dysregulation. Journal of Crohn's and Colitis, 2021, 15, 1908-1919.	0.6	7
116	Molecular Diagnostic Outcomes from 700 Cases. Journal of Molecular Diagnostics, 2022, 24, 274-286.	1.2	7
117	Association of a Polymorphic Variant of the Adiponectin Gene with Insulin Resistance in African Americans. Clinical and Translational Science, 2008, 1, 194-199.	1.5	6
118	IQ and hemizygosity for the Val¹⁵⁸Met functional polymorphism of COMT in 22q11DS. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 1112-1115.	1.1	6
119	Comparative analysis of different approaches for dealing with candidate regions in the context of a genome-wide association study. BMC Proceedings, 2009, 3, S93.	1.8	5
120	Next-Generation Linkage Analysis. Human Heredity, 2011, 72, 227-227.	0.4	5
121	Introduction. Human Heredity, 2008, 66, 65-66.	0.4	4
122	Integration of Omics Data in Genetic Epidemiology. Human Heredity, 2015, 79, 109-110.	0.4	4
123	Genomic copy number variation association study in Caucasian patients with nonsyndromic cryptorchidism. BMC Urology, 2016, 16, 62.	0.6	4
124	An Exploratory Association Analysis of the Insulin Gene Region With Diabetes Mellitus in Two Dog Breeds. Journal of Heredity, 2019, 110, 793-800.	1.0	4
125	Immune Dysregulation in Human ITCH Deficiency Successfully Treated with Hematopoietic Cell Transplantation. Journal of Allergy and Clinical Immunology: in Practice, 2021, 9, 2885-2893.e3.	2.0	4
126	Prenatal diagnosis and linkage disequilibrium with cystic fibrosis for markers surrounding D7S8. Human Genetics, 1990, 85, 275-8.	1.8	3

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127	Comparison of Sib Pairâ€Based Approaches for Identifying Quantitative Trait Loci Underlying Asthma in the Busselton Families. Genetic Epidemiology, 2001, 21, S198-203.	0.6	3
128	N-myc ONCOGENE AMPLIFICATION AND CATECHOLAMINE METABOLISM IN PATIENTS WITH NEUROBLASTOMA. Lancet, The, 1987, 330, 795.	6.3	2
129	Linkage study of early-onset obesity to leptin receptor gene in Italian children. Nutrition Research, 2000, 20, 1059-1063.	1.3	2
130	Genetic Mapping of Quantitative Trait Loci for Disease-Related Phenotypes. Methods in Molecular Biology, 2012, 871, 281-311.	0.4	2
131	Commentary on Mutations in Interleukin-10 Receptor and Clinical Phenotypes in Patients with Very Early-onset Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2017, 23, 591-592.	0.9	2
132	Genetic analysis in African American children supports ancestry specific neuroblastoma susceptibility. Cancer Epidemiology Biomarkers and Prevention, 2022, , cebp.EPI-21-0782-A.2021.	1.1	1
133	Preliminary results on the frequency of the Î”F508 mutation in cystic fibrosis patients from the USSR. Human Genetics, 1990, 85, 423-424.	1.8	0
134	Analysis of Human Genetic Data: A Celebratory Issue of Human Heredity Honoring Dr. JÃ¼rg Ott. Human Heredity, 2011, 71, 85-85.	0.4	0
135	Attention-deficit hyperactivity disorder. , 0, , 168-182.		0
136	P-197â€ƒIdentification of a Homozygous Mutation in the ZBTB24 Gene in a Patient with Very Early Onset Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2016, 22, S68.	0.9	0
137	O-003â€ƒUnderstanding the Relevance of Whole Exome Sequencing Identified Variants in Patients with Very Early-Onset-IBD. Inflammatory Bowel Diseases, 2016, 22, S1-S2.	0.9	0