

# Jeffrey C Murray

## List of Publications by Year in descending order

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136  
papers

16,645  
citations

61984

43  
h-index

18647

119  
g-index

143  
all docs

143  
docs citations

143  
times ranked

22151  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	27.8	3,823
2	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	21.4	1,818
3	Cleft lip and palate: understanding genetic and environmental influences. <i>Nature Reviews Genetics</i> , 2011, 12, 167-178.	16.3	1,435
4	Cloning and characterization of a novel bicoid-related homeobox transcription factor gene, RIEG, involved in Rieger syndrome. <i>Nature Genetics</i> , 1996, 14, 392-399.	21.4	852
5	Mutations in IRF6 cause Van der Woude and popliteal pterygium syndromes. <i>Nature Genetics</i> , 2002, 32, 285-289.	21.4	784
6	A genome-wide association study of cleft lip with and without cleft palate identifies risk variants near MAFB and ABCA4. <i>Nature Genetics</i> , 2010, 42, 525-529.	21.4	518
7	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. <i>Nature Genetics</i> , 2019, 51, 804-814.	21.4	402
8	Disruption of an AP-2 binding site in an IRF6 enhancer is associated with cleft lip. <i>Nature Genetics</i> , 2008, 40, 1341-1347.	21.4	382
9	A novel homeobox gene PITX3 is mutated in families with autosomal-dominant cataracts and ASMD. <i>Nature Genetics</i> , 1998, 19, 167-170.	21.4	371
10	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015, 11, e1005378.	3.5	331
11	Domain-specific mutations in TGFB1 result in Camurati-Engelmann disease. <i>Nature Genetics</i> , 2000, 26, 19-20.	21.4	239
12	Genetic Evidence for Causal Relationships Between Maternal Obesity-Related Traits and Birth Weight. <i>JAMA - Journal of the American Medical Association</i> , 2016, 315, 1129.	7.4	220
13	Integrated human genome-wide maps constructed using the CEPH reference panel. <i>Nature Genetics</i> , 1994, 6, 391-393.	21.4	216
14	Genetics of Nonsyndromic Orofacial Clefts. <i>Cleft Palate-Craniofacial Journal</i> , 2012, 49, 73-91.	0.9	212
15	Long term follow up study of survival associated with cleft lip and palate at birth. <i>BMJ: British Medical Journal</i> , 2004, 328, 1405.	2.3	205
16	Candidate genes for nonsyndromic cleft lip and palate and maternal cigarette smoking and alcohol consumption: Evaluation of genotype-environment interactions from a population-based case-control study of orofacial clefts. , 1999, 59, 39-50.		198
17	Dominant Mutations in GRHL3 Cause Van der Woude Syndrome and Disrupt Oral Periderm Development. <i>American Journal of Human Genetics</i> , 2014, 94, 23-32.	6.2	195
18	A cohort study of recurrence patterns among more than 54 000 relatives of oral cleft cases in Denmark: support for the multifactorial threshold model of inheritance. <i>Journal of Medical Genetics</i> , 2010, 47, 162-168.	3.2	188

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19	Mutations in the human forkhead transcription factor FOXE3 associated with anterior segment ocular dysgenesis and cataracts. <i>Human Molecular Genetics</i> , 2001, 10, 231-236.	2.9	184
20	A multi-ethnic genome-wide association study identifies novel loci for non-syndromic cleft lip with or without cleft palate on 2p24.2, 17q23 and 19q13. <i>Human Molecular Genetics</i> , 2016, 25, ddw104.	2.9	163
21	Genome-wide association study of offspring birth weight in 86%577 women identifies five novel loci and highlights maternal genetic effects that are independent of fetal genetics. <i>Human Molecular Genetics</i> , 2018, 27, 742-756.	2.9	156
22	Identification of Functional Variants for Cleft Lip with or without Cleft Palate in or near PAX7, FGFR2, and NOG by Targeted Sequencing of GWAS Loci. <i>American Journal of Human Genetics</i> , 2015, 96, 397-411.	6.2	150
23	A Genome-wide Association Study of Nonsyndromic Cleft Palate Identifies an Etiologic Missense Variant in GRHL3. <i>American Journal of Human Genetics</i> , 2016, 98, 744-754.	6.2	146
24	Evidence for gene-environment interaction in a genome wide study of nonsyndromic cleft palate. <i>Genetic Epidemiology</i> , 2011, 35, n/a-n/a.	1.3	145
25	Genome-Wide Association Study Reveals Multiple Loci Influencing Normal Human Facial Morphology. <i>PLoS Genetics</i> , 2016, 12, e1006149.	3.5	140
26	Genome-wide meta-analyses of nonsyndromic orofacial clefts identify novel associations between FOXE1 and all orofacial clefts, and TP63 and cleft lip with or without cleft palate. <i>Human Genetics</i> , 2017, 136, 275-286.	3.8	139
27	FOXE1 association with both isolated cleft lip with or without cleft palate, and isolated cleft palate. <i>Human Molecular Genetics</i> , 2009, 18, 4879-4896.	2.9	136
28	Maternal Contributions to Preterm Delivery. <i>American Journal of Epidemiology</i> , 2009, 170, 1358-1364.	3.4	133
29	The Pitx2 protein in mouse development. <i>Developmental Dynamics</i> , 2000, 218, 195-200.	1.8	126
30	Genome Scan, Fine-Mapping, and Candidate Gene Analysis of Non-Syndromic Cleft Lip with or without Cleft Palate Reveals Phenotype-Specific Differences in Linkage and Association Results. <i>Human Heredity</i> , 2009, 68, 151-170.	0.8	113
31	Cancer Risk in Persons with Oral Clefts—A Population-based Study of 8,093 Cases. <i>American Journal of Epidemiology</i> , 2005, 161, 1047-1055.	3.4	111
32	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
33	Mutations in the Epithelial Cadherin-p120-Catenin Complex Cause Mendelian Non-Syndromic Cleft Lip with or without Cleft Palate. <i>American Journal of Human Genetics</i> , 2018, 102, 1143-1157.	6.2	94
34	Integrated Genomic Analyses in Bronchopulmonary Dysplasia. <i>Journal of Pediatrics</i> , 2015, 166, 531-537.e13.	1.8	93
35	Cleft palate: players, pathways, and pursuits. <i>Journal of Clinical Investigation</i> , 2004, 113, 1676-1678.	8.2	93
36	Analysis of select folate pathway genes, PAX3, and human T in a midwestern neural tube defect population. <i>Teratology</i> , 1999, 59, 331-341.	1.6	88

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37	Genome-Wide Association Studies in Dogs and Humans Identify ADAMTS20 as a Risk Variant for Cleft Lip and Palate. <i>PLoS Genetics</i> , 2015, 11, e1005059.	3.5	82
38	Whole Exome Sequencing of Distant Relatives in Multiplex Families Implicates Rare Variants in Candidate Genes for Oral Clefts. <i>Genetics</i> , 2014, 197, 1039-1044.	2.9	79
39	Pitx2 Regulates Procollagen Lysyl Hydroxylase (Plod) Gene Expression. <i>Journal of Cell Biology</i> , 2001, 152, 545-552.	5.2	78
40	Microdeletions at chromosome bands 1q32-q41 as a cause of Van der Woude syndrome. , 1999, 84, 145-150.		65
41	Clinical and environmental influences on metabolic biomarkers collected for newborn screening. <i>Clinical Biochemistry</i> , 2013, 46, 133-138.	1.9	62
42	Mapping a New Spontaneous Preterm Birth Susceptibility Gene, IGF1R, Using Linkage, Haplotype Sharing, and Association Analysis. <i>PLoS Genetics</i> , 2011, 7, e1001293.	3.5	61
43	Genomic analyses in African populations identify novel risk loci for cleft palate. <i>Human Molecular Genetics</i> , 2019, 28, 1038-1051.	2.9	61
44	Multimiomics Characterization of Preterm Birth in Low- and Middle-Income Countries. <i>JAMA Network Open</i> , 2020, 3, e2029655.	5.9	53
45	Antioxidant response genes sequence variants and BPD susceptibility in VLBW infants. <i>Pediatric Research</i> , 2015, 77, 477-483.	2.3	52
46	A LINE-1 Insertion in DLX6 Is Responsible for Cleft Palate and Mandibular Abnormalities in a Canine Model of Pierre Robin Sequence. <i>PLoS Genetics</i> , 2014, 10, e1004257.	3.5	49
47	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. <i>Nature Communications</i> , 2019, 10, 3927.	12.8	49
48	Genome-wide Enrichment of De Novo Coding Mutations in Orofacial Cleft Trios. <i>American Journal of Human Genetics</i> , 2020, 107, 124-136.	6.2	48
49	Gene-Centric Analysis of Preeclampsia Identifies Maternal Association at <i>PLEKHG1</i> . <i>Hypertension</i> , 2018, 72, 408-416.	2.7	46
50	A solution pathway for preterm birth: accelerating a priority research agenda. <i>The Lancet Global Health</i> , 2013, 1, e328-e330.	6.3	44
51	Identifying Genetic Sources of Phenotypic Heterogeneity in Orofacial Clefts by Targeted Sequencing. <i>Birth Defects Research</i> , 2017, 109, 1030-1038.	1.5	41
52	A new human homeobox gene OGI2X is a member of the most conserved homeobox gene family and is expressed during heart development in mouse. <i>Human Molecular Genetics</i> , 1998, 7, 415-422.	2.9	40
53	A phase 2 study of valproic acid and radiation, followed by maintenance valproic acid and bevacizumab in children with newly diagnosed diffuse intrinsic pontine glioma or high-grade glioma. <i>Pediatric Blood and Cancer</i> , 2020, 67, e28283.	1.5	40
54	Dissecting maternal and fetal genetic effects underlying the associations between maternal phenotypes, birth outcomes, and adult phenotypes: A mendelian-randomization and haplotype-based genetic score analysis in 10,734 mother-infant pairs. <i>PLoS Medicine</i> , 2020, 17, e1003305.	8.4	37

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55	Association studies of low-frequency coding variants in nonsyndromic cleft lip with or without cleft palate. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 1531-1538.	1.2	36
56	A systematic genetic analysis and visualization of phenotypic heterogeneity among orofacial cleft GWAS signals. <i>Genetic Epidemiology</i> , 2019, 43, 704-716.	1.3	36
57	Whole exome sequencing reveals HSPA1L as a genetic risk factor for spontaneous preterm birth. <i>PLoS Genetics</i> , 2018, 14, e1007394.	3.5	35
58	Evidence of Gene-Environment Interaction for Two Genes on Chromosome 4 and Environmental Tobacco Smoke in Controlling the Risk of Nonsyndromic Cleft Palate. <i>PLoS ONE</i> , 2014, 9, e88088.	2.5	33
59	Genome-wide association study of sepsis in extremely premature infants. <i>Archives of Disease in Childhood: Fetal and Neonatal Edition</i> , 2017, 102, F439-F445.	2.8	32
60	Novel IRF6 mutations in Japanese patients with Van der Woude Syndrome: two missense mutations (R45Q and P396S) and a 17-kb deletion. <i>Journal of Human Genetics</i> , 2003, 48, 622-628.	2.3	31
61	Replication of 13q31.1 association in nonsyndromic cleft lip with cleft palate in Europeans. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1054-1060.	1.2	31
62	Recurrence risk for offspring of twins discordant for oral cleft: A population-based cohort study of the Danish 1936-2004 cleft twin cohort. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 2468-2474.	1.2	28
63	Mutations in GDF11 and the extracellular antagonist, Follistatin, as a likely cause of Mendelian forms of orofacial clefting in humans. <i>Human Mutation</i> , 2019, 40, 1813-1825.	2.5	26
64	Evidence for SNP-SNP interaction identified through targeted sequencing of cleft case-parent trios. <i>Genetic Epidemiology</i> , 2017, 41, 244-250.	1.3	24
65	Identification of 16q21 as a modifier of nonsyndromic orofacial cleft phenotypes. <i>Genetic Epidemiology</i> , 2017, 41, 887-897.	1.3	24
66	Polymorphisms of CYP51A1 from Cholesterol Synthesis: Associations with Birth Weight and Maternal Lipid Levels and Impact on CYP51 Protein Structure. <i>PLoS ONE</i> , 2013, 8, e82554.	2.5	24
67	The effect of follow-up on limiting non-participation bias in genetic epidemiologic investigations. <i>European Journal of Epidemiology</i> , 1998, 14, 129-138.	5.7	23
68	Assignment of gene responsible for progressive pseudorheumatoid dysplasia to chromosome 6 and examination of COL10A1 as candidate gene. <i>European Journal of Human Genetics</i> , 1998, 6, 251-256.	2.8	22
69	Maternal and neonatal epidemiological features in clinical subtypes of preterm birth. <i>Journal of Maternal-Fetal and Neonatal Medicine</i> , 2016, 29, 3153-3161.	1.5	22
70	Polymorphisms in urea cycle enzyme genes are associated with persistent pulmonary hypertension of the newborn. <i>Pediatric Research</i> , 2018, 83, 142-147.	2.3	22
71	Search for genetic modifiers of IRF6 and genotype-phenotype correlations in Van der Woude and popliteal pterygium syndromes. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2535-2544.	1.2	21
72	Exclusion of the branchio-oto-renal syndrome locus (EYA1) from patients with branchio-oculo-facial syndrome. , 2000, 91, 387-390.		19

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73	Whole genome sequencing of orofacial cleft trios from the Gabriella Miller Kids First Pediatric Research Consortium identifies a new locus on chromosome 21. <i>Human Genetics</i> , 2020, 139, 215-226.	3.8	19
74	Association of low-frequency genetic variants in regulatory regions with nonsyndromic orofacial clefts. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 467-474.	1.2	18
75	SPECC1L regulates palate development downstream of IRF6. <i>Human Molecular Genetics</i> , 2020, 29, 845-858.	2.9	18
76	Pleiotropy method reveals genetic overlap between orofacial clefts at multiple novel loci from GWAS of multi-ethnic trios. <i>PLoS Genetics</i> , 2021, 17, e1009584.	3.5	18
77	Common Genetic Variants in FOXP2 Are Not Associated with Individual Differences in Language Development. <i>PLoS ONE</i> , 2016, 11, e0152576.	2.5	18
78	Genomic structure, sequence, and mapping of human FGF8 with no evidence for its role in craniosynostosis/limb defect syndromes. <i>Development</i> , 1997, 72, 354-362.		16
79	Temporal lobe pleomorphic xanthoastrocytoma and acquired BRAF mutation in an adolescent with the constitutional 22q11.2 deletion syndrome. <i>Journal of Neuro-Oncology</i> , 2011, 102, 509-514.	2.9	16
80	Deletions and loss-of-function variants in TP63 associated with orofacial clefting. <i>European Journal of Human Genetics</i> , 2019, 27, 1101-1112.	2.8	16
81	Genetic variants associated with patent ductus arteriosus in extremely preterm infants. <i>Journal of Perinatology</i> , 2019, 39, 401-408.	2.0	16
82	Genome-Wide Association Study of Non-syndromic Orofacial Clefts in a Multiethnic Sample of Families and Controls Identifies Novel Regions. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 621482.	3.7	16
83	Genome-wide analysis of parent-of-origin interaction effects with environmental exposure (PoOxE): An application to European and Asian cleft palate trios. <i>PLoS ONE</i> , 2017, 12, e0184358.	2.5	16
84	Genome-wide interaction studies identify sex-specific risk alleles for nonsyndromic orofacial clefts. <i>Genetic Epidemiology</i> , 2018, 42, 664-672.	1.3	15
85	FAT4 identified as a potential modifier of orofacial cleft laterality. <i>Genetic Epidemiology</i> , 2021, 45, 721-735.	1.3	14
86	Relationship of Genetic Variants With Procedural Pain, Anxiety, and Distress in Children. <i>Biological Research for Nursing</i> , 2017, 19, 339-349.	1.9	13
87	Advancing human health in the decade ahead: pregnancy as a key window for discovery. <i>American Journal of Obstetrics and Gynecology</i> , 2020, 223, 312-321.	1.3	13
88	Non-random distribution of deleterious mutations in the DNA and protein-binding domains of IRF6 are associated with Van Der Woude syndrome. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e1355.	1.2	13
89	Association of maternal prenatal selenium concentration and preterm birth: a multicountry meta-analysis. <i>BMJ Global Health</i> , 2021, 6, e005856.	4.7	13
90	Determining the prevalence of cytomegalovirus infection in a cohort of preterm infants. <i>Journal of Neonatal-Perinatal Medicine</i> , 2015, 8, 137-141.	0.8	12

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91	Integrative genetic, genomic and transcriptomic analysis of heat shock protein and nuclear hormone receptor gene associations with spontaneous preterm birth. <i>Scientific Reports</i> , 2021, 11, 17115.	3.3	12
92	CDH1 Mutation Distribution and Type Suggests Genetic Differences between the Etiology of Orofacial Clefting and Gastric Cancer. <i>Genes</i> , 2020, 11, 391.	2.4	11
93	Cloning and chromosomal localization of two novel human genes encoding LIM-domain binding factors CLIM1 and CLIM2/LDB1/NLI. <i>Mammalian Genome</i> , 1998, 9, 921-924.	2.2	10
94	Whole exome association of rare deletions in multiplex oral cleft families. <i>Genetic Epidemiology</i> , 2017, 41, 61-69.	1.3	10
95	Using an aquatic model, <i>Xenopus laevis</i> , to uncover the role of chromodomain 1 in craniofacial disorders. <i>Genesis</i> , 2021, 59, e23394.	1.6	10
96	Identification of single nucleotide polymorphisms in hematopoietic cell transplant patients affecting early recognition of, and response to, endotoxin. <i>Innate Immunity</i> , 2014, 20, 697-711.	2.4	9
97	Novel <i>GREM1</i> Variations in Sub-Saharan African Patients With Cleft Lip and/or Cleft Palate. <i>Cleft Palate-Craniofacial Journal</i> , 2018, 55, 736-742.	0.9	9
98	The PAX1 locus at 20p11 is a potential genetic modifier for bilateral cleft lip. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100025.	1.7	9
99	Genomic structure of the human retinoic acid receptor-alpha1 gene. <i>Mammalian Genome</i> , 1999, 10, 528-529.	2.2	8
100	Time for T. <i>Nature Genetics</i> , 2001, 29, 107-109.	21.4	8
101	Polymorphisms in NR5A2, gene encoding liver receptor homolog-1 are associated with preterm birth. <i>Pediatric Research</i> , 2016, 79, 776-780.	2.3	8
102	Association of candidate gene polymorphisms with clinical subtypes of preterm birth in a Latin American population. <i>Pediatric Research</i> , 2017, 82, 554-559.	2.3	8
103	DNA methylation of a novel PAK4 locus influences ototoxicity susceptibility following cisplatin and radiation therapy for pediatric embryonal tumors. <i>Neuro-Oncology</i> , 2017, 19, 1372-1379.	1.2	7
104	Cost-effectiveness of a gestational age metabolic algorithm for preterm and small-for-gestational-age classification. <i>American Journal of Obstetrics &amp; Gynecology MFM</i> , 2021, 3, 100279.	2.6	7
105	Characterization of the calcitonin/CGRP gene in Williams syndrome. <i>American Journal of Medical Genetics Part A</i> , 1991, 39, 28-33.	2.4	6
106	Inferring disease risk genes from sequencing data in multiplex pedigrees through sharing of rare variants. <i>Genetic Epidemiology</i> , 2019, 43, 37-49.	1.3	6
107	Genetic variation in CYB5R3 is associated with methemoglobin levels in preterm infants receiving nitric oxide therapy. <i>Pediatric Research</i> , 2015, 77, 472-476.	2.3	5
108	Assessing the impact of nicotine dependence genes on the risk of facial clefts: An example of the use of national registry and biobank data. <i>Norsk Epidemiologi</i> , 2012, 21, 241-250.	0.3	5

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109	Re-evaluating surgery and re-irradiation for locally recurrent pediatric ependymoma – a multi-institutional study. <i>Neuro-Oncology Advances</i> , 2021, 3, vdab158.	0.7	5
110	Identification of paternal uniparental disomy on chromosome 22 and a <i>de novo</i> deletion on chromosome 18 in individuals with orofacial clefts. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2018, 6, 924-932.	1.2	4
111	Genome-wide association study of multiethnic nonsyndromic orofacial cleft families identifies novel loci specific to family and phenotypic subtypes. <i>Genetic Epidemiology</i> , 2022, , .	1.3	4
112	Exclusion of <i>Ifa</i> and <i>Ifb</i> as the <i>Lps</i> gene and mapping of three markers near the <i>Lps</i> locus. <i>Mammalian Genome</i> , 1997, 8, 785-786.	2.2	3
113	Genetic predictors of severe intraventricular hemorrhage in extremely low-birthweight infants. <i>Journal of Perinatology</i> , 2021, 41, 286-294.	2.0	3
114	Co-occurrence of orofacial clefts and clubfoot phenotypes in a sub-Saharan African cohort: Whole-exome sequencing implicates multiple syndromes and genes. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1655.	1.2	3
115	Variant analyses of candidate genes in orofacial clefts in multiethnic populations. <i>Oral Diseases</i> , 2022, 28, 1921-1935.	3.0	3
116	Analysis of select folate pathway genes, <i>PAX3</i> , and human <i>T</i> in a midwestern neural tube defect population. <i>Teratology</i> , 1999, 59, 331-341.	1.6	3
117	Matroshka and ectopic polymorphisms: Two new classes of DNA sequence variation identified at the Van der Woude syndrome locus on 1q32-q41. <i>Human Mutation</i> , 2001, 18, 422-434.	2.5	2
118	Detection of <i>de novo</i> copy number deletions from targeted sequencing of trios. <i>Bioinformatics</i> , 2019, 35, 571-578.	4.1	2
119	Targeted newborn metabolomics: prediction of gestational age from cord blood. <i>Journal of Perinatology</i> , 2022, 42, 181-186.	2.0	2
120	Genome-Wide Scan for Parent-of-Origin Effects in a sub-Saharan African Cohort With Nonsyndromic Cleft Lip and/or Cleft Palate (CL/P). <i>Cleft Palate-Craniofacial Journal</i> , 2021, , 105566562110363.	0.9	1
121	Genetic Variants and the Cortisol Response in Children: An Exploratory Study. <i>Biological Research for Nursing</i> , 2019, 21, 157-165.	1.9	1
122	Direct Sequencing of Candidate Genes for Nonsyndromic Cleft Lip and Palate. <i>PLoS Genetics</i> , 2005, preprint, e64.	3.5	1
123	The chromosome 4 workshop report. <i>Genomics</i> , 1992, 12, 857-858.	2.9	0
124	Clinical Pathology Case: Hyperimmunoglobulin E Syndrome (Job's Syndrome). <i>Fetal and Pediatric Pathology</i> , 2001, 20, 223-226.	0.3	0
125	2013 Presidential Address: Just Another President's Speech (but It's All about You). <i>American Journal of Human Genetics</i> , 2014, 94, 319-323.	6.2	0
126	Front Cover, Volume 40, Issue 10. <i>Human Mutation</i> , 2019, 40, i.	2.5	0

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127	Reply to "Diversity is essential for good science and Reproductive science is no different: A response to the recent formulation of the Burroughs Wellcome Fund Pregnancy Think-Tank" American Journal of Obstetrics and Gynecology, 2020, 223, 951-952.	1.3	0
128	Interactions between PDA-associated polymorphisms and genetic ancestry alter ductus arteriosus gene expression. Pediatric Research, 2021, , .	2.3	0
129	Identifying Genetic Sources of Phenotypic Heterogeneity in Orofacial Clefts by Targeted Sequencing. Birth Defects Research, 2017, , .	1.5	0
130	CLINICAL PATHOLOGY CASE: HYPERIMMUNOGLOBULIN E SYNDROME (JOB'S SYNDROME). Fetal and Pediatric Pathology, 2001, 20, 223-226.	0.3	0
131	Title is missing!. , 2020, 17, e1003305.		0
132	Title is missing!. , 2020, 17, e1003305.		0
133	Title is missing!. , 2020, 17, e1003305.		0
134	Title is missing!. , 2020, 17, e1003305.		0
135	Title is missing!. , 2020, 17, e1003305.		0
136	Title is missing!. , 2020, 17, e1003305.		0