

# Mary L Marazita

## List of Publications by Year in descending order

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161  
papers

8,757  
citations

61857

43  
h-index

53109

85  
g-index

169  
all docs

169  
docs citations

169  
times ranked

7078  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cleft lip and palate: understanding genetic and environmental influences. <i>Nature Reviews Genetics</i> , 2011, 12, 167-178.	7.7	1,435
2	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.	9.4	518
3	Disruption of an AP-2 binding site in an IRF6 enhancer is associated with cleft lip. <i>Nature Genetics</i> , 2008, 40, 1341-1347.	9.4	382
4	Genetics of cleft lip and cleft palate. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2013, 163, 246-258.	0.7	336
5	Genome-wide meta-analyses of nonsyndromic cleft lip with or without cleft palate identify six new risk loci. <i>Nature Genetics</i> , 2012, 44, 968-971.	9.4	311
6	Genome-wide analyses of non-syndromic cleft lip with palate identify 14 novel loci and genetic heterogeneity. <i>Nature Communications</i> , 2017, 8, 14364.	5.8	207
7	Genome-wide mapping of global-to-local genetic effects on human facial shape. <i>Nature Genetics</i> , 2018, 50, 414-423.	9.4	205
8	Meta-Analysis of 13 Genome Scans Reveals Multiple Cleft Lip/Palate Genes with Novel Loci on 9q21 and 2q32-35. <i>American Journal of Human Genetics</i> , 2004, 75, 161-173.	2.6	200
9	Genome-wide analysis of dental caries and periodontitis combining clinical and self-reported data. <i>Nature Communications</i> , 2019, 10, 2773.	5.8	183
10	Evidence for Autosomal Dominant Inheritance and Race-specific Heterogeneity in Early Onset Periodontitis. <i>Journal of Periodontology</i> , 1994, 65, 623-630.	1.7	177
11	The Evolution of Human Genetic Studies of Cleft Lip and Cleft Palate. <i>Annual Review of Genomics and Human Genetics</i> , 2012, 13, 263-283.	2.5	174
12	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.	1.4	163
13	Genetic factors influencing risk to orofacial clefts: today's challenges and tomorrow's opportunities. <i>F1000Research</i> , 2016, 5, 2800.	0.8	155
14	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.	2.6	150
15	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.	2.6	146
16	Evidence for gene-environment interaction in a genome wide study of nonsyndromic cleft palate. <i>Genetic Epidemiology</i> , 2011, 35, n/a-n/a.	0.6	145
17	Genome-Wide Association Study Reveals Multiple Loci Influencing Normal Human Facial Morphology. <i>PLoS Genetics</i> , 2016, 12, e1006149.	1.5	140
18	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.	1.8	139

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19	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.	1.4	136
20	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.	0.4	113
21	Current concepts in the embryology and genetics of cleft lip and cleft palate. <i>Clinics in Plastic Surgery</i> , 2004, 31, 125-140.	0.7	112
22	Genome Scan for Loci Involved in Cleft Lip With or Without Cleft Palate, in Chinese Multiplex Families. <i>American Journal of Human Genetics</i> , 2002, 71, 349-364.	2.6	107
23	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
24	Insights into the genetic architecture of the human face. <i>Nature Genetics</i> , 2021, 53, 45-53.	9.4	94
25	Case/control family study of autonomic nervous system dysfunction in idiopathic congenital central hypoventilation syndrome. <i>American Journal of Medical Genetics Part A</i> , 2001, 100, 237-245.	2.4	83
26	Genetic analysis of cleft lip with or without cleft palate in Danish kindreds. <i>American Journal of Medical Genetics Part A</i> , 1984, 19, 9-18.	2.4	82
27	The 3D Facial Norms Database: Part 1. A Web-Based Craniofacial Anthropometric and Image Repository for the Clinical and Research Community. <i>Cleft Palate-Craniofacial Journal</i> , 2016, 53, 185-197.	0.5	80
28	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. <i>Nature Communications</i> , 2020, 11, 5562.	5.8	80
29	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.	1.2	79
30	Genetic correlation between smoking behaviors and schizophrenia. <i>Schizophrenia Research</i> , 2018, 194, 86-90.	1.1	71
31	Orbicularis oris muscle defects as an expanded phenotypic feature in nonsyndromic cleft lip with or without cleft palate. <i>American Journal of Medical Genetics, Part A</i> , 2007, 143A, 1143-1149.	0.7	69
32	Study protocol of the Center for Oral Health Research in Appalachia (COHRA) etiology study. <i>BMC Oral Health</i> , 2008, 8, 18.	0.8	69
33	Genome-wide association study of facial morphology reveals novel associations with FREM1 and PARK2. <i>PLoS ONE</i> , 2017, 12, e0176566.	1.1	68
34	Using the 3D Facial Norms Database to investigate craniofacial sexual dimorphism in healthy children, adolescents, and adults. <i>Biology of Sex Differences</i> , 2016, 7, 23.	1.8	65
35	The FaceBase Consortium: A comprehensive resource for craniofacial researchers. <i>Development (Cambridge)</i> , 2016, 143, 2677-88.	1.2	62
36	Genomic analyses in African populations identify novel risk loci for cleft palate. <i>Human Molecular Genetics</i> , 2019, 28, 1038-1051.	1.4	61

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37	Novel genetic loci affecting facial shape variation in humans. <i>ELife</i> , 2019, 8, .	2.8	58
38	Sonic Hedgehog regulation of <i>Foxf2</i> promotes cranial neural crest mesenchyme proliferation and is disrupted in cleft lip morphogenesis. <i>Development (Cambridge)</i> , 2017, 144, 2082-2091.	1.2	55
39	Genome-Wide Association Study of Periodontal Health Measured by Probing Depth in Adults Ages 18-49 years. <i>G3: Genes, Genomes, Genetics</i> , 2014, 4, 307-314.	0.8	54
40	Genetic segregation analysis of autonomic nervous system dysfunction in families of probands with idiopathic congenital central hypoventilation syndrome. <i>American Journal of Medical Genetics Part A</i> , 2001, 100, 229-236.	2.4	49
41	Subclinical features in non-syndromic cleft lip with or without cleft palate (CL/P): review of the evidence that subepithelial orbicularis oris muscle defects are part of an expanded phenotype for CL/P. <i>Orthodontics and Craniofacial Research</i> , 2007, 10, 82-87.	1.2	49
42	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. <i>Nature Communications</i> , 2019, 10, 3927.	5.8	49
43	Nonsyndromic Cleft Lip with or without Cleft Palate in China: Assessment of Candidate Regions. <i>Cleft Palate-Craniofacial Journal</i> , 2002, 39, 149-156.	0.5	48
44	Identification of common non-coding variants at 1p22 that are functional for non-syndromic orofacial clefting. <i>Nature Communications</i> , 2017, 8, 14759.	5.8	48
45	Facial recognition from DNA using face-to-DNA classifiers. <i>Nature Communications</i> , 2019, 10, 2557.	5.8	46
46	Cleft lip with or without cleft palate: Reanalysis of a three-generation family study from England. <i>Genetic Epidemiology</i> , 1986, 3, 335-342.	0.6	45
47	Investigating the shared genetics of non-syndromic cleft lip/palate and facial morphology. <i>PLoS Genetics</i> , 2018, 14, e1007501.	1.5	44
48	Identifying Genetic Sources of Phenotypic Heterogeneity in Orofacial Clefts by Targeted Sequencing. <i>Birth Defects Research</i> , 2017, 109, 1030-1038.	0.8	41
49	Heritability of face shape in twins: a preliminary study using 3D stereophotogrammetry and geometric morphometrics. <i>Dentistry 3000</i> , 2013, 1, 7-11.	0.1	39
50	Genome-scan for loci involved in cleft lip with or without cleft palate in consanguineous families from Turkey. , 2004, 126A, 111-122.		37
51	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.	0.7	36
52	A systematic genetic analysis and visualization of phenotypic heterogeneity among orofacial cleft GWAS signals. <i>Genetic Epidemiology</i> , 2019, 43, 704-716.	0.6	36
53	Rapid Testing of SNPs and Gene-Environment Interactions in Case-Parent Trio Data Based on Exact Analytic Parameter Estimation. <i>Biometrics</i> , 2012, 68, 766-773.	0.8	34
54	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.	1.1	33

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55	Using the PhenX Toolkit to Add Standard Measures to a Study. <i>Current Protocols in Human Genetics</i> , 2015, 86, 1.21.1-1.21.17.	3.5	33
56	Oral Health in a Sample of Pregnant Women from Northern Appalachia (2011â€“2015). <i>International Journal of Dentistry</i> , 2015, 2015, 1-12.	0.5	32
57	Consortium-based genome-wide meta-analysis for childhood dental caries traits. <i>Human Molecular Genetics</i> , 2018, 27, 3113-3127.	1.4	32
58	Segregation analysis of attention deficit hyperactivity disorder. , 1999, 88, 71-78.		31
59	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.	0.7	31
60	Caries Experience Differs between Females and Males across Age Groups in Northern Appalachia. <i>International Journal of Dentistry</i> , 2015, 2015, 1-8.	0.5	30
61	Geneâ€“Gene Interaction Among <i>WNT</i> Genes for Oral Cleft in Trios. <i>Genetic Epidemiology</i> , 2015, 39, 385-394.	0.6	30
62	Effects of Specimen Collection Methodologies and Storage Conditions on the Short-Term Stability of Oral Microbiome Taxonomy. <i>Applied and Environmental Microbiology</i> , 2016, 82, 5519-5529.	1.4	30
63	Limb development genes underlie variation in human fingerprint patterns. <i>Cell</i> , 2022, 185, 95-112.e18.	13.5	30
64	Genome-wide association study of dental caries in the Hispanic Communities Health Study/Study of Latinos (HCHS/SOL). <i>Human Molecular Genetics</i> , 2016, 25, 807-816.	1.4	29
65	Multiethnic CWAS Reveals Polygenic Architecture of Earlobe Attachment. <i>American Journal of Human Genetics</i> , 2017, 101, 913-924.	2.6	29
66	FaceBase 3: analytical tools and FAIR resources for craniofacial and dental research. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	25
67	Evidence for SNP-SNP interaction identified through targeted sequencing of cleft case-parent trios. <i>Genetic Epidemiology</i> , 2017, 41, 244-250.	0.6	24
68	Identification of 16q21 as a modifier of nonsyndromic orofacial cleft phenotypes. <i>Genetic Epidemiology</i> , 2017, 41, 887-897.	0.6	24
69	Exploring the genomic basis of early childhood caries: a pilot study. <i>International Journal of Paediatric Dentistry</i> , 2018, 28, 217-225.	1.0	24
70	SNPs Associated With Testosterone Levels Influence Human Facial Morphology. <i>Frontiers in Genetics</i> , 2018, 9, 497.	1.1	23
71	Aquaporin 5 Interacts with Fluoride and Possibly Protects against Caries. <i>PLoS ONE</i> , 2015, 10, e0143068.	1.1	22
72	Search for genetic modifiers of IRF6 and genotypeâ€“phenotype correlations in Van der Woude and popliteal pterygium syndromes. <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2535-2544.	0.7	21

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73	Cleft lip/palate and educational attainment: cause, consequence or correlation? A Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2020, 49, 1282-1293.	0.9	21
74	Depression and Rural Environment Are Associated With Poor Oral Health Among Pregnant Women in Northern Appalachia. <i>Behavior Modification</i> , 2016, 40, 325-340.	1.1	20
75	Toward a genetic understanding of dental fear: evidence of heritability. <i>Community Dentistry and Oral Epidemiology</i> , 2017, 45, 66-73.	0.9	20
76	A Preliminary Genome-Wide Association Study of Pain-Related Fear: Implications for Orofacial Pain. <i>Pain Research and Management</i> , 2017, 2017, 1-12.	0.7	20
77	Six NSCL/P Loci Show Associations With Normal-Range Craniofacial Variation. <i>Frontiers in Genetics</i> , 2018, 9, 502.	1.1	20
78	The effects of family, dentition, and dental caries on the salivary microbiome. <i>Annals of Epidemiology</i> , 2016, 26, 348-354.	0.9	19
79	Periodontal Status and Quality of Life: Impact of Fear of Pain and Dental Fear. <i>Pain Research and Management</i> , 2017, 2017, 1-9.	0.7	19
80	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.	1.8	19
81	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.	0.7	18
82	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.	1.5	18
83	Exploring the effect of dentition, dental decay and familiarity on oral health using metabolomics. <i>Infection, Genetics and Evolution</i> , 2014, 22, 201-207.	1.0	17
84	Predictors of dental care utilization in north-central Appalachia in the USA. <i>Community Dentistry and Oral Epidemiology</i> , 2019, 47, 283-290.	0.9	17
85	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.	1.8	16
86	Using the PhenX Toolkit to Select Standard Measurement Protocols for Your Research Study. <i>Current Protocols</i> , 2021, 1, e149.	1.3	16
87	Genome-Wide Association Study (GWAS) of dental caries in diverse populations. <i>BMC Oral Health</i> , 2021, 21, 377.	0.8	16
88	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.	1.1	16
89	Expanding the cleft phenotype: the dental characteristics of unaffected parents of Australian children with non-syndromic cleft lip and palate. <i>International Journal of Paediatric Dentistry</i> , 2014, 24, 286-292.	1.0	15
90	Genome-wide interaction studies identify sex-specific risk alleles for nonsyndromic orofacial clefts. <i>Genetic Epidemiology</i> , 2018, 42, 664-672.	0.6	15

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91	Oral Health Disparities in Appalachia. <i>Journal of the American Dental Association</i> , 2008, 139, 598-604.	0.7	14
92	Effects of Smoking and Genotype on the PSR Index of Periodontal Disease in Adults Aged 18â€“49. <i>International Journal of Environmental Research and Public Health</i> , 2012, 9, 2839-2850.	1.2	14
93	Transmission of dental fear from parent to adolescent in an Appalachian sample in the USA. <i>International Journal of Paediatric Dentistry</i> , 2019, 29, 720-727.	1.0	14
94	Genetic association and differential expression of PITX2 with acute appendicitis. <i>Human Genetics</i> , 2019, 138, 37-47.	1.8	14
95	Low levels of salivary metals, oral microbiome composition and dental decay. <i>Scientific Reports</i> , 2020, 10, 14640.	1.6	14
96	<i>FAT4</i> identified as a potential modifier of orofacial cleft laterality. <i>Genetic Epidemiology</i> , 2021, 45, 721-735.	0.6	14
97	Non-random distribution of deleterious mutations in the DNA and protein-binding domains of <i>IRF6</i> are associated with Van Der Woude syndrome. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e1355.	0.6	13
98	Genome scans of facial features in East Africans and cross-population comparisons reveal novel associations. <i>PLoS Genetics</i> , 2021, 17, e1009695.	1.5	13
99	Genetic Etiologies of Facial Clefting. , 0, , 147-161.		12
100	Genetic Modifiers of Patent Ductus Arteriosus in Term Infants. <i>Journal of Pediatrics</i> , 2016, 176, 57-61.e1.	0.9	12
101	Genetic variants in pachyonychia congenita-associated keratins increase susceptibility to tooth decay. <i>PLoS Genetics</i> , 2018, 14, e1007168.	1.5	12
102	Exploring palatal and dental shape variation with 3D shape analysis and geometric deep learning. <i>Orthodontics and Craniofacial Research</i> , 2021, 24, 134-143.	1.2	12
103	Variants on chromosome 4q21 near PKD2 and SIBLINGs are associated with dental caries. <i>Journal of Human Genetics</i> , 2017, 62, 491-496.	1.1	11
104	Mapping genetic variants for cranial vault shape in humans. <i>PLoS ONE</i> , 2018, 13, e0196148.	1.1	11
105	Whole exome association of rare deletions in multiplex oral cleft families. <i>Genetic Epidemiology</i> , 2017, 41, 61-69.	0.6	10
106	Vitamin D metabolic loci and preeclampsia risk in multi-ethnic pregnant women. <i>Physiological Reports</i> , 2018, 6, e13468.	0.7	10
107	Vitamin D metabolic loci and vitamin D status in Black and White pregnant women. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2018, 220, 61-68.	0.5	10
108	Effects of Male Facial Masculinity on Perceived Attractiveness. <i>Adaptive Human Behavior and Physiology</i> , 2021, 7, 73-88.	0.6	10

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109	The Intersection of the Genetic Architectures of Orofacial Clefts and Normal Facial Variation. <i>Frontiers in Genetics</i> , 2021, 12, 626403.	1.1	10
110	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.	0.5	9
111	A Multivariate Approach to Determine the Dimensionality of Human Facial Asymmetry. <i>Symmetry</i> , 2020, 12, 348.	1.1	9
112	The PAX1 locus at 20p11 is a potential genetic modifier for bilateral cleft lip. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100025.	1.0	9
113	Exploration of genetic factors determining cleft side in a pair of monozygotic twins with mirror-image cleft lip and palate using whole-genome sequencing and comparison of craniofacial morphology. <i>Archives of Oral Biology</i> , 2018, 96, 33-38.	0.8	8
114	GWAS reveals loci associated with velopharyngeal dysfunction. <i>Scientific Reports</i> , 2018, 8, 8470.	1.6	8
115	Novel caries loci in children and adults implicated by genome-wide analysis of families. <i>BMC Oral Health</i> , 2018, 18, 98.	0.8	8
116	Integrative approaches generate insights into the architecture of non-syndromic cleft lip and cleft palate. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100038.	1.0	8
117	Testing the face shape hypothesis in twins discordant for nonsyndromic orofacial clefting. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2886-2892.	0.7	7
118	Genetics of Orofacial Cleft Birth Defects. <i>Current Genetic Medicine Reports</i> , 2015, 3, 118-126.	1.9	6
119	Diagnosing subtle palatal anomalies: Validation of video-analysis and assessment protocol for diagnosing occult submucous cleft palate. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2017, 100, 242-246.	0.4	6
120	Inferring disease risk genes from sequencing data in multiplex pedigrees through sharing of rare variants. <i>Genetic Epidemiology</i> , 2019, 43, 37-49.	0.6	6
121	Individuals with nonsyndromic orofacial clefts have increased asymmetry of fingerprint patterns. <i>PLoS ONE</i> , 2020, 15, e0230534.	1.1	6
122	Fluctuating Asymmetry and Sexual Dimorphism in Human Facial Morphology: A Multi-Variate Study. <i>Symmetry</i> , 2021, 13, 304.	1.1	6
123	PRICKLE1-FOCAD Interaction Revealed by Genome-Wide vQTL Analysis of Human Facial Traits. <i>Frontiers in Genetics</i> , 2021, 12, 674642.	1.1	6
124	Identifying genetic risk loci for diabetic complications and showing evidence for heterogeneity of type 1 diabetes based on complications risk. <i>PLoS ONE</i> , 2018, 13, e0192696.	1.1	6
125	Symptoms of Attention-Deficit Hyperactivity Disorder, Nonsyndromic Orofacial Cleft Children, and Dopamine Polymorphisms. <i>Biological Research for Nursing</i> , 2015, 17, 257-262.	1.0	5
126	Whole-genome sequencing in a pair of monozygotic twins with discordant cleft lip and palate subtypes. <i>Oral Diseases</i> , 2018, 24, 1303-1309.	1.5	5



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127	Co-occurrence of yeast, streptococci, dental decay, and gingivitis in the post-partum period: results of a longitudinal study. <i>Journal of Oral Microbiology</i> , 2020, 12, 1746494.	1.2	5
128	Prevalence of Torus Palatinus and association with dental arch shape in a multi-ethnic cohort. <i>HOMO- Journal of Comparative Human Biology</i> , 2020, 71, 273-280.	0.3	5
129	Gene-gene interaction of single nucleotide polymorphisms in 16p13.3 may contribute to the risk of non-syndromic cleft lip with or without cleft palate in Chinese case-parent trios. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 1489-1494.	0.7	4
130	Soft tissue nasal asymmetry as an indicator of orofacial cleft predisposition. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1296-1303.	0.7	4
131	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.	0.6	4
132	Primary teeth microhardness and lead (Pb) levels. <i>Heliyon</i> , 2019, 5, e01551.	1.4	4
133	Predictors of use of dental care by children in north-central Appalachia in the USA. <i>PLoS ONE</i> , 2021, 16, e0250488.	1.1	4
134	Mother's Perceived Social Support and Children's Dental Caries in Northern Appalachia. <i>Pediatric Dentistry (discontinued)</i> , 2019, 41, 200-205.	0.4	4
135	Genome-wide association study of multiethnic nonsyndromic orofacial cleft families identifies novel loci specific to family and phenotypic subtypes. <i>Genetic Epidemiology</i> , 2022, , .	0.6	4
136	Racism in oral healthcare settings: Implications for dental care-related fear/anxiety and utilization among Black/African American women in Appalachia. <i>Journal of Public Health Dentistry</i> , 2022, 82, 28-35.	0.5	4
137	Is the Fagerstr�m test for nicotine dependence invariant across secular trends in smoking? A question for cross-birth cohort analysis of nicotine dependence. <i>Drug and Alcohol Dependence</i> , 2018, 185, 127-132.	1.6	3
138	Impact of low-frequency coding variants on human facial shape. <i>Scientific Reports</i> , 2021, 11, 748.	1.6	3
139	Replication of GWAS significant loci in a sub-Saharan African Cohort with early childhood caries: a pilot study. <i>BMC Oral Health</i> , 2021, 21, 274.	0.8	3
140	Variant analyses of candidate genes in orofacial clefts in multi-ethnic populations. <i>Oral Diseases</i> , 2022, 28, 1921-1935.	1.5	3
141	Pittsburgh Registry of Infant Multiplets (PRIM). , 0, .		3
142	Genome-wide Interaction Study Implicates VGLL2 and Alcohol Exposure and PRL and Smoking in Orofacial Cleft Risk. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 621261.	1.8	3
143	Pittsburgh Registry of Infant Multiplets (PRIM): An Update. <i>Twin Research and Human Genetics</i> , 2006, 9, 1006-1008.	0.3	2
144	Detection of de novo copy number deletions from targeted sequencing of trios. <i>Bioinformatics</i> , 2019, 35, 571-578.	1.8	2

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145	Speech Phenotyping in Unaffected Family Members of Individuals With Nonsyndromic Cleft Lip With or Without Palate. <i>Cleft Palate-Craniofacial Journal</i> , 2019, 56, 867-876.	0.5	2
146	Detecting Gene-Environment Interaction for Maternal Exposures Using Case-Parent Trios Ascertained Through a Case With Non-Syndromic Orofacial Cleft. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 621018.	1.8	2
147	Exploring Mothersâ€™ Perspectives About Why Grandparents in Appalachia Give Their Grandchildren Cariogenic Foods and Beverages: A Qualitative Study. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2022, , .	0.4	2
148	The Influence of Sex and Ancestry on Three-Dimensional Palate Shape. <i>Journal of Craniofacial Surgery</i> , 2021, 32, 2883-2887.	0.3	1
149	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.	0.5	1
150	Direct Sequencing of Candidate Genes for Nonsyndromic Cleft Lip and Palate. <i>PLoS Genetics</i> , 2005, preprint, e64.	1.5	1
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152	Linkage analysis and multi-point mapping of 11p markers. <i>Genetic Epidemiology</i> , 1986, 3, 159-164.	0.6	0
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156	Variant Analyses of Candidate Genes in Orofacial Clefts in Multiâ€¢Ethnic Populations. <i>FASEB Journal</i> , 2021, 35, .	0.2	0
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