

Jacqueline B Hetmanski

List of Publications by Year in descending order

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16
papers

1,257
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840776

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#	ARTICLE	IF	CITATIONS
1	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.	3.7	2
2	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
3	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
4	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
5	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
6	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
7	Identifying a Deletion Affecting Total Lung Capacity Among Subjects in the COPD Gene Study Cohort. <i>Genetic Epidemiology</i> , 2016, 40, 81-88.	1.3	5
8	Hemizygous Deletion on Chromosome 3p26.1 Is Associated with Heavy Smoking among African American Subjects in the COPD Gene Study. <i>PLoS ONE</i> , 2016, 11, e0164134.	2.5	4
9	Novel evidence of association with nonsyndromic cleft lip with or without cleft palate was shown for single nucleotide polymorphisms in <i>FOXF2</i> gene in an Asian population. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2015, 103, 857-862.	1.6	11
10	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
11	Gene-Gene Interaction Among <i>WNT</i> Genes for Oral Cleft in Trios. <i>Genetic Epidemiology</i> , 2015, 39, 385-394.	1.3	30
12	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
13	Joint Testing of Genotypic and Gene-Environment Interaction Identified Novel Association for BMP4 with Non-Syndromic CL/P in an Asian Population Using Data from an International Cleft Consortium. <i>PLoS ONE</i> , 2014, 9, e109038.	2.5	17
14	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
15	Risk loci for chronic obstructive pulmonary disease: a genome-wide association study and meta-analysis. <i>Lancet Respiratory Medicine</i> , 2014, 2, 214-225.	10.7	291
16	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