

Amanda A Seyerle

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

377
citations

1307594

7
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

1450
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. <i>American Journal of Human Genetics</i> , 2016, 98, 165-184.	6.2	266
2	GWAS of the electrocardiographic QT interval in Hispanics/Latinos generalizes previously identified loci and identifies population-specific signals. <i>Scientific Reports</i> , 2017, 7, 17075.	3.3	23
3	Evidence of Heterogeneity by Race/Ethnicity in Genetic Determinants of QT Interval. <i>Epidemiology</i> , 2014, 25, 790-798.	2.7	22
4	Fine mapping of QT interval regions in global populations refines previously identified QT interval loci and identifies signals unique to African and Hispanic descent populations. <i>Heart Rhythm</i> , 2017, 14, 572-580.	0.7	19
5	Genome-wide association study of PR interval in Hispanics/Latinos identifies novel locus at <i>ID2</i> . <i>Heart</i> , 2018, 104, 904-911.	2.9	12
6	A genome-wide interaction analysis of tricyclic/tetracyclic antidepressants and RR and QT intervals: a pharmacogenomics study from the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) consortium. <i>Journal of Medical Genetics</i> , 2017, 54, 313-323.	3.2	9
7	GWAS of QRS duration identifies new loci specific to Hispanic/Latino populations. <i>PLoS ONE</i> , 2019, 14, e0217796.	2.5	8
8	Genetic epidemiology: the potential benefits and challenges of using genetic information to improve human health. <i>North Carolina Medical Journal</i> , 2013, 74, 505-8.	0.2	8
9	Multi-Ethnic Genome-Wide Association Study of Decomposed Cardioelectric Phenotypes Illustrates Strategies to Identify and Characterize Evidence of Shared Genetic Effects for Complex Traits. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002680.	3.6	4
10	Comparison of 2 models for gene-environment interactions: an example of simulated gene-medication interactions on systolic blood pressure in family-based data. <i>BMC Proceedings</i> , 2016, 10, 371-377.	1.6	3
11	BinomiRare: A robust test for association of a rare genetic variant with a binary outcome for mixed models and any case-control proportion. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100040.	1.7	2