

Kelsey E Grinde

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

281
citations

1307594

7
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

1110
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Generalizing polygenic risk scores from Europeans to Hispanics/Latinos. <i>Genetic Epidemiology</i> , 2019, 43, 50-62. | 1.3 | 89 |
| 2 | A fully adjusted two-stage procedure for rank-normalization in genetic association studies. <i>Genetic Epidemiology</i> , 2019, 43, 263-275. | 1.3 | 60 |
| 3 | Local Ancestry Inference in a Large US-Based Hispanic/Latino Study: Hispanic Community Health Study/Study of Latinos (HCHS/SOL). <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 1525-1534. | 1.8 | 51 |
| 4 | Genome-wide Significance Thresholds for Admixture Mapping Studies. <i>American Journal of Human Genetics</i> , 2019, 104, 454-465. | 6.2 | 25 |
| 5 | Coagulation factor VIII: Relationship to cardiovascular disease risk and whole genome sequence and epigenome-wide analysis in African Americans. <i>Journal of Thrombosis and Haemostasis</i> , 2020, 18, 1335-1347. | 3.8 | 17 |
| 6 | Whole genome sequence analyses of eGFR in 23,732 people representing multiple ancestries in the NHLBI trans-omics for precision medicine (TOPMed) consortium. <i>EBioMedicine</i> , 2021, 63, 103157. | 6.1 | 14 |
| 7 | A general approach for combining diverse rare variant association tests provides improved robustness across a wider range of genetic architectures. <i>European Journal of Human Genetics</i> , 2016, 24, 767-773. | 2.8 | 12 |
| 8 | A general method for combining different family-based rare-variant tests of association to improve power and robustness of a wide range of genetic architectures. <i>BMC Proceedings</i> , 2016, 10, 165-170. | 1.6 | 4 |
| 9 | Illustrating, Quantifying, and Correcting for Bias in Post-hoc Analysis of Gene-Based Rare Variant Tests of Association. <i>Frontiers in Genetics</i> , 2017, 8, 117. | 2.3 | 4 |
| 10 | Circular RNA Profiles in Viremia and ART Suppression Predict Competing circRNA-miRNA-mRNA Networks Exclusive to HIV-1 Viremic Patients. <i>Viruses</i> , 2022, 14, 683. | 3.3 | 3 |
| 11 | A multistep approach to single nucleotide polymorphism-set analysis: an evaluation of power and type I error of gene-based tests of association after pathway-based association tests. <i>BMC Proceedings</i> , 2016, 10, 349-355. | 1.6 | 2 |
| 12 | Mo1146 ANESTHESIA PROFESSIONAL-DELIVERED SEDATION IS ASSOCIATED WITH SIMILAR OUTCOMES COMPARED TO NURSE ADMINISTERED SEDATION IN PATIENTS WITH ACUTE UPPER GASTROINTESTINAL BLEEDING. <i>Gastrointestinal Endoscopy</i> , 2018, 87, AB418-AB419. | 1.0 | 0 |