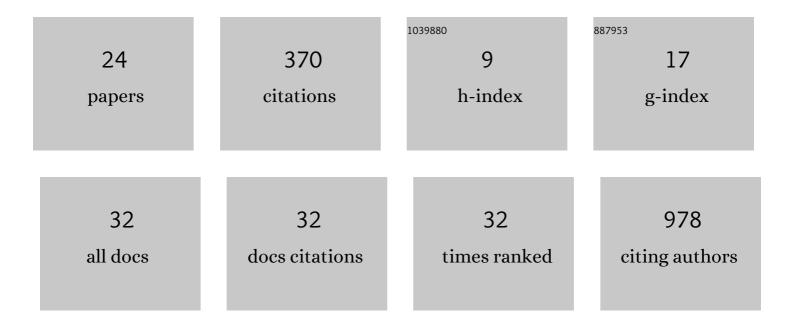
Kevin L Keys

List of Publications by Year in descending order

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KEVIN L KEVS

| # | Article | lF | CITATIONS |
|----|---|------|-----------|
| 1 | Whole-Genome Sequencing of Pharmacogenetic Drug Response in Racially Diverse Children with Asthma. American Journal of Respiratory and Critical Care Medicine, 2018, 197, 1552-1564. | 2.5 | 102 |
| 2 | On the cross-population generalizability of gene expression prediction models. PLoS Genetics, 2020, 16, e1008927. | 1.5 | 41 |
| 3 | Sequencing and imputation in GWAS: Costâ€effective strategies to increase power and genomic coverage across diverse populations. Genetic Epidemiology, 2020, 44, 537-549. | 0.6 | 30 |
| 4 | OpenMendel: a cooperative programming project for statistical genetics. Human Genetics, 2020, 139, 61-71. | 1.8 | 29 |
| 5 | Ambient air pollution, asthma drug response, and telomere length in African American youth. Journal of Allergy and Clinical Immunology, 2019, 144, 839-845.e10. | 1.5 | 24 |
| 6 | Genetic Determinants of Telomere Length in African American Youth. Scientific Reports, 2018, 8, 13265. | 1.6 | 20 |
| 7 | Genomeâ€wide association study reveals a novel locus for asthma with severe exacerbations in diverse populations. Pediatric Allergy and Immunology, 2021, 32, 106-115. | 1.1 | 17 |
| 8 | Ten Simple Rules for Getting Help from Online Scientific Communities. PLoS Computational Biology, 2011, 7, e1002202. | 1.5 | 12 |
| 9 | BioSimulator.jl: Stochastic simulation in Julia. Computer Methods and Programs in Biomedicine, 2018, 167, 23-35. | 2.6 | 12 |
| 10 | Epigenome-wide association study of lung function in Latino children and youth with asthma. Clinical Epigenetics, 2022, 14, 9. | 1.8 | 12 |
| 11 | Lung Function in African American Children with Asthma Is Associated with Novel Regulatory Variants of the KIT Ligand <i>KITLG/SCF</i> and Gene-By-Air-Pollution Interaction. Genetics, 2020, 215, 869-886. | 1.2 | 11 |
| 12 | Whole-Genome Sequencing Identifies Novel Functional Loci Associated with Lung Function in Puerto Rican Youth. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 962-972. | 2.5 | 11 |
| 13 | Iterative hard thresholding in genome-wide association studies: Generalized linear models, prior weights, and double sparsity. GigaScience, 2020, 9, . | 3.3 | 10 |
| 14 | A genome-wide study of DNA methylation in white blood cells and asthma in Latino children and youth. Epigenetics, 2021, 16, 577-585. | 1.3 | 10 |
| 15 | Iterative hard thresholding for model selection in genomeâ€wide association studies. Genetic Epidemiology, 2017, 41, 756-768. | 0.6 | 6 |
| 16 | Integrative genomic analysis in African American children with asthma finds three novel loci associated with lung function. Genetic Epidemiology, 2021, 45, 190-208. | 0.6 | 4 |
| 17 | Influence of pathway topology and functional class on the molecular evolution of human metabolic genes. PLoS ONE, 2018, 13, e0208782. | 1.1 | 3 |
| 18 | Proximal Distance Algorithms: Theory and Practice. Journal of Machine Learning Research, 2019, 20, . | 62.4 | 0 |

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| # | Article | IF | CITATIONS |
|----|--|----|-----------|
| 19 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | Ο |
| 20 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | 0 |
| 21 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | 0 |
| 22 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | 0 |
| 23 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | О |
| 24 | On the cross-population generalizability of gene expression prediction models. , 2020, 16, e1008927. | | 0 |