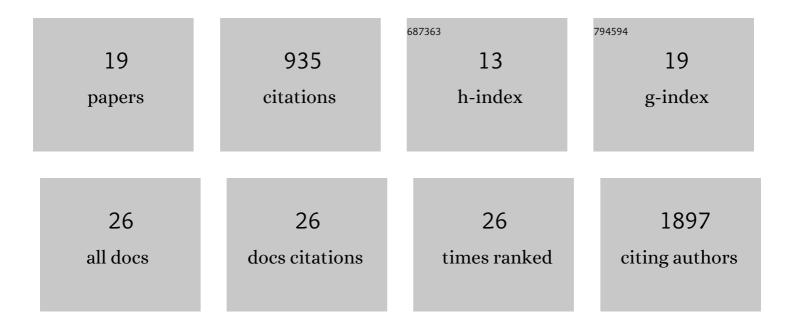
Lawrence H Uricchio

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2162252/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Revisiting the malaria hypothesis: accounting for polygenicity and pleiotropy. Trends in Parasitology, 2022, 38, 290-301. | 3.3 | 5 |
| 2 | Understanding the emergence of contingent and deterministic exclusion in multispecies communities. Ecology Letters, 2021, 24, 2155-2168. | 6.4 | 8 |
| 3 | How will mosquitoes adapt to climate warming?. ELife, 2021, 10, . | 6.0 | 46 |
| 4 | Evolutionary perspectives on polygenic selection, missing heritability, and GWAS. Human Genetics, 2020, 139, 5-21. | 3.8 | 37 |
| 5 | Priority Effects and Nonhierarchical Competition Shape Species Composition in a Complex Grassland Community. American Naturalist, 2019, 193, 213-226. | 2.1 | 31 |
| 6 | Ultrarare variants drive substantial cis heritability of human gene expression. Nature Genetics, 2019, 51, 1349-1355. | 21.4 | 98 |
| 7 | An evolutionary compass for detecting signals of polygenic selection and mutational bias. Evolution Letters, 2019, 3, 69-79. | 3.3 | 29 |
| 8 | Exploiting selection at linked sites to infer the rate and strength of adaptation. Nature Ecology and Evolution, 2019, 3, 977-984. | 7.8 | 43 |
| 9 | An admixture mapping meta-analysis implicates genetic variation at 18q21 with asthma susceptibility in Latinos. Journal of Allergy and Clinical Immunology, 2019, 143, 957-969. | 2.9 | 33 |
| 10 | Singleton Variants Dominate the Genetic Architecture of Human Gene Expression. SSRN Electronic Journal, 2018, , . | 0.4 | 4 |
| 11 | Selection and explosive growth alter genetic architecture and hamper the detection of causal rare variants. Genome Research, 2016, 26, 863-873. | 5.5 | 63 |
| 12 | An analytical upper bound on the number of loci required for all splits of a species tree to appear in a set of gene trees. BMC Bioinformatics, 2016, 17, 417. | 2.6 | 1 |
| 13 | Population Genetic Simulations of Complex Phenotypes with Implications for Rare Variant Association Tests. Genetic Epidemiology, 2015, 39, 35-44. | 1.3 | 16 |
| 14 | Robust Forward Simulations of Recurrent Hitchhiking. Genetics, 2014, 197, 221-236. | 2.9 | 24 |
| 15 | Population Genetics of Rare Variants and Complex Diseases. Human Heredity, 2012, 74, 118-128. | 0.8 | 53 |
| 16 | Accurate Imputation of Rare and Common Variants in a Founder Population From a Small Number of Sequenced Individuals. Genetic Epidemiology, 2012, 36, 312-319. | 1.3 | 19 |
| 17 | Exome sequencing reveals a novel mutation for autosomal recessive non-syndromic mental retardation in the TECR gene on chromosome 19p13. Human Molecular Genetics, 2011, 20, 1285-1289. | 2.9 | 94 |
| 18 | Inflammasome-Mediated Production of IL-1β Is Required for Neutrophil Recruitment against <i>Staphylococcus aureus</i> In Vivo. Journal of Immunology, 2007, 179, 6933-6942. | 0.8 | 294 |

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Coarse-grained entropy decrease and phase-space focusing in Hamiltonian dynamics. Physical Review A, 2005, 72, . | 2.5 | 2 |