Andrew P Morgan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8110989/publications.pdf

Version: 2024-02-01

25 papers 2,241 citations

393982 19 h-index 25 g-index

34 all docs

34 docs citations

times ranked

34

3474 citing authors

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Population structure and inbreeding in wild house mice (Mus musculus) at different geographic scales. Heredity, 2022, 129, 183-194. | 1.2 | 12 |
| 2 | The epidemiology of Plasmodium vivax among adults in the Democratic Republic of the Congo. Nature Communications, 2021, 12, 4169. | 5.8 | 18 |
| 3 | <i>Prdm9</i> Intersubspecific Interactions in Hybrid Male Sterility of House Mouse. Molecular Biology and Evolution, 2020, 37, 3423-3438. | 3.5 | 24 |
| 4 | The impact of antimalarial resistance on the genetic structure of Plasmodium falciparum in the DRC. Nature Communications, 2020, 11, 2107. | 5.8 | 57 |
| 5 | Falciparum malaria from coastal Tanzania and Zanzibar remains highly connected despite effective control efforts on the archipelago. Malaria Journal, 2020, 19, 47. | 0.8 | 30 |
| 6 | Instability of the Pseudoautosomal Boundary in House Mice. Genetics, 2019, 212, 469-487. | 1.2 | 15 |
| 7 | Structural Variation Shapes the Landscape of Recombination in Mouse. Genetics, 2017, 206, 603-619. | 1.2 | 51 |
| 8 | Male Infertility Is Responsible for Nearly Half of the Extinction Observed in the Mouse Collaborative Cross. Genetics, 2017, 206, 557-572. | 1.2 | 66 |
| 9 | Genomes of the Mouse Collaborative Cross. Genetics, 2017, 206, 537-556. | 1.2 | 189 |
| 10 | Wild Mouse Gut Microbiota Promotes Host Fitness and Improves Disease Resistance. Cell, 2017, 171, 1015-1028.e13. | 13.5 | 603 |
| 11 | Sequence and Structural Diversity of Mouse Y Chromosomes. Molecular Biology and Evolution, 2017, 34, 3186-3204. | 3.5 | 54 |
| 12 | Genetic architecture of atherosclerosis dissected by QTL analyses in three F2 intercrosses of apolipoprotein E-null mice on C57BL6/J, DBA/2J and 129S6/SvEvTac backgrounds. PLoS ONE, 2017, 12, e0182882. | 1.1 | 4 |
| 13 | Allelic Variation in the Toll-Like Receptor Adaptor Protein <i>Ticam2 < /i>Contributes to SARS-Coronavirus Pathogenesis in Mice. G3: Genes, Genomes, Genetics, 2017, 7, 1653-1663.</i> | 0.8 | 75 |
| 14 | argyle: An R Package for Analysis of Illumina Genotyping Arrays. G3: Genes, Genomes, Genetics, 2016, 6, 281-286. | 0.8 | 36 |
| 15 | The Mouse Universal Genotyping Array: From Substrains to Subspecies. G3: Genes, Genomes, Genetics, 2016, 6, 263-279. | 0.8 | 199 |
| 16 | Whole Genome Sequence of Two Wild-Derived <i>Mus musculus domesticus</i> Inbred Strains, LEWES/EiJ and ZALENDE/EiJ, with Different Diploid Numbers. G3: Genes, Genomes, Genetics, 2016, 6, 4211-4216. | 0.8 | 12 |
| 17 | Diversity Outbred Mice at 21: Maintaining Allelic Variation in the Face of Selection. G3: Genes, Genomes, Genetics, 2016, 6, 3893-3902. | 0.8 | 66 |
| 18 | The Evolutionary Fates of a Large Segmental Duplication in Mouse. Genetics, 2016, 204, 267-285. | 1.2 | 21 |

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | <i>R2d2</i> Drives Selfish Sweeps in the House Mouse. Molecular Biology and Evolution, 2016, 33, 1381-1395. | 3.5 | 55 |
| 20 | A Multi-Megabase Copy Number Gain Causes Maternal Transmission Ratio Distortion on Mouse Chromosome 2. PLoS Genetics, 2015, 11, e1004850. | 1.5 | 76 |
| 21 | Analyses of allele-specific gene expression in highly divergent mouse crosses identifies pervasive allelic imbalance. Nature Genetics, 2015, 47, 353-360. | 9.4 | 204 |
| 22 | Informatics resources for the Collaborative Cross and related mouse populations. Mammalian Genome, 2015, 26, 521-539. | 1.0 | 50 |
| 23 | The Antipsychotic Olanzapine Interacts with the Gut Microbiome to Cause Weight Gain in Mouse. PLoS ONE, 2014, 9, e115225. | 1.1 | 147 |
| 24 | High-Resolution Sex-Specific Linkage Maps of the Mouse Reveal Polarized Distribution of Crossovers in Male Germline. Genetics, 2014, 197, 91-106. | 1.2 | 85 |
| 25 | The Collaborative Cross as a Resource for Modeling Human Disease: CC011/Unc, a New Mouse Model for Spontaneous Colitis. Mammalian Genome, 2014, 25, 95-108. | 1.0 | 78 |