Chen-Hsin Yu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11583030/publications.pdf

Version: 2024-02-01

| | | 1163117 | 1474206 | |
|----------|----------------|--------------|----------------|--|
| 8 | 325 | 8 | 9 | |
| papers | citations | h-index | g-index | |
| | | | | |
| | | | | |
| 0 | 0 | 0 | 707 | |
| 9 | 9 | 9 | 787 | |
| all docs | docs citations | times ranked | citing authors | |
| | | | | |

| # | Article | IF | CITATIONS |
|---|--|-----|-----------|
| 1 | Population genomics and the evolution of virulence in the fungal pathogen <i>Cryptococcus neoformans</i> . Genome Research, 2017, 27, 1207-1219. | 5.5 | 134 |
| 2 | Consensus Genome-Wide Expression Quantitative Trait Loci and Their Relationship with Human Complex Trait Disease. OMICS A Journal of Integrative Biology, 2016, 20, 400-414. | 2.0 | 46 |
| 3 | Genomeâ€wide analysis of the regulation of Cu metabolism in <i>Cryptococcus neoformans</i> Molecular Microbiology, 2018, 108, 473-494. | 2.5 | 34 |
| 4 | Cryptococcus neoformans resists to drastic conditions by switching to viable but non-culturable cell phenotype. PLoS Pathogens, 2019, 15, e1007945. | 4.7 | 31 |
| 5 | Landscape of gene expression variation of natural isolates of Cryptococcus neoformans in response to biologically relevant stresses. Microbial Genomics, 2020, 6, . | 2.0 | 24 |
| 6 | Gene Expression of Diverse Cryptococcus Isolates during Infection of the Human Central Nervous System. MBio, 2021, 12, e0231321. | 4.1 | 23 |
| 7 | Insights from GWAS: emerging landscape of mechanisms underlying complex trait disease. BMC Genomics, 2015, 16, S4. | 2.8 | 16 |
| 8 | Matching phenotypes to whole genomes: Lessons learned from four iterations of the personal genome project community challenges. Human Mutation, 2017, 38, 1266-1276. | 2.5 | 14 |