## Akio Toh-E

## List of Publications by Year in descending order

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933447 888059 18 691 10 17 citations h-index g-index papers 19 19 19 707 citing authors docs citations times ranked all docs

| #  | Article                                                                                                                                                                                                                           | IF   | CITATIONS |
|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 1  | Genetic system underlying responses of Cryptococcus neoformans to cadmium. Current Genetics, 2022, 68, 125-141.                                                                                                                   | 1.7  | 2         |
| 2  | Identification and characterization of a sulfite reductase gene and new insights regarding the sulfur-containing amino acid metabolism in the basidiomycetous yeast Cryptococcus neoformans. Current Genetics, 2021, 67, 115-128. | 1.7  | 5         |
| 3  | Vph1 is associated with the copper homeostasis of <i>Cryptococcus neoformans</i> serotype D. Journal of General and Applied Microbiology, 2021, 67, 195-206.                                                                      | 0.7  | 1         |
| 4  | Step-wise elimination of $\hat{l}$ ±-mitochondrial nucleoids and mitochondrial structure as a basis for the strict uniparental inheritance in Cryptococcus neoformans. Scientific Reports, 2020, 10, 2468.                        | 3.3  | 9         |
| 5  | Novel biosynthetic pathway for sulfur amino acids in Cryptococcus neoformans. Current Genetics, 2018, 64, 681-696.                                                                                                                | 1.7  | 13        |
| 6  | Creation, characterization and utilization of Cryptococcus neoformans mutants sensitive to micafungin. Current Genetics, 2017, 63, 1093-1104.                                                                                     | 1.7  | 12        |
| 7  | Putative orotate transporter of Cryptococcus neoformans, Oat1, is a member of the NCS1/PRT transporter super family and its loss causes attenuation of virulence. Current Genetics, 2017, 63, 697-707.                            | 1.7  | 0         |
| 8  | Local Anesthetics and Antipsychotic Phenothiazines Interact Nonspecifically with Membranes and Inhibit Hexose Transporters in Yeast. Genetics, 2016, 202, 997-1012.                                                               | 2.9  | 14        |
| 9  | Identification of genes involved in the phosphate metabolism in Cryptococcus neoformans. Fungal Genetics and Biology, 2015, 80, 19-30.                                                                                            | 2.1  | 26        |
| 10 | Positional cloning in Cryptococcus neoformans and its application for identification and cloning of the gene encoding methylenetetrahydrofolate reductase. Fungal Genetics and Biology, 2015, 76, 70-77.                          | 2.1  | 6         |
| 11 | Quantitative live-cell imaging reveals spatio-temporal dynamics and cytoplasmic assembly of the 26S proteasome. Nature Communications, 2014, 5, 3396.                                                                             | 12.8 | 111       |
| 12 | Functional characterization of PMT2, encoding a protein-O-mannosyltransferase, in the human pathogen Cryptococcus neoformans. Fungal Genetics and Biology, 2014, 69, 13-22.                                                       | 2.1  | 16        |
| 13 | Structural analysis of compounds with actions similar to local anesthetics and antipsychotic phenothiazines in yeast. Yeast, 2011, 28, 391-404.                                                                                   | 1.7  | 12        |
| 14 | Non-homologous end joining pathway of the human pathogen Cryptococcus neoformans influences homologous integration efficiency but not virulence. Mycoscience, 2010, 51, 272-280.                                                  | 0.8  | 10        |
| 15 | Transcriptional Repression by the Pho4 Transcription Factor Controls the Timing of <i>SNZ1</i> Expression. Eukaryotic Cell, 2008, 7, 949-957.                                                                                     | 3.4  | 14        |
| 16 | Forchlorfenuron, a phenylurea cytokinin, disturbs septin organization in Saccharomyces cerevisiae.<br>Genes and Genetic Systems, 2004, 79, 199-206.                                                                               | 0.7  | 39        |
| 17 | Phosphorylation of Sic1, a Cyclin-dependent Kinase (Cdk) Inhibitor, by Cdk Including Pho85 Kinase Is<br>Required for Its Prompt Degradation. Molecular Biology of the Cell, 1998, 9, 2393-2405.                                   | 2.1  | 96        |
| 18 | Isolation and Characterization of Acid Phosphatase Mutants in <i>Saccharomyces cerevisiae</i> Journal of Bacteriology, 1973, 113, 727-738.                                                                                        | 2.2  | 305       |