

# Miao Tian

## List of Publications by Year in descending order

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18  
papers

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docs citations

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| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Identification and utilization of a mutated 60S ribosomal subunit coding gene as an effective and cost-efficient selection marker for <i>Tetrahymena</i> genetic manipulation. <i>International Journal of Biological Macromolecules</i> , 2022, 204, 1-8. | 7.5  | 1         |
| 2  | Arrested crossover precursor structures form stable homologous bonds in a <i>Tetrahymena</i> meiotic mutant. <i>PLoS ONE</i> , 2022, 17, e0263691.   | 2.5  | 2         |
| 3  | Zfp1, a Cys2His2 zinc finger protein is required for meiosis initiation in <i>Tetrahymena thermophila</i> . <i>Cell Cycle</i> , 2022, , 1-12.  | 2.6  | 1         |
| 4  | Spatial constraints on chromosomes are instrumental to meiotic pairing. <i>Journal of Cell Science</i> , 2020, 133, .  | 2.0  | 12        |
| 5  | Non-coding RNA Transcription in <i>Tetrahymena</i> Meiotic Nuclei Requires Dedicated Mediator Complex-Associated Proteins. <i>Current Biology</i> , 2019, 29, 2359-2370.e5.  | 3.9  | 9         |
| 6  | A specialized condensin complex participates in somatic nuclear maturation in <i>Tetrahymena thermophila</i> . <i>Molecular Biology of the Cell</i> , 2019, 30, 1326-1338.   | 2.1  | 8         |
| 7  | An MCM family protein promotes interhomolog recombination by preventing precocious intersister repair of meiotic DSBs. <i>PLoS Genetics</i> , 2019, 15, e1008514.  | 3.5  | 6         |
| 8  | A DP-like transcription factor protein interacts with E2f1 to regulate meiosis in <i>Tetrahymena thermophila</i> . <i>Cell Cycle</i> , 2018, 17, 634-642.  | 2.6  | 31        |
| 9  | A chromatin-associated protein required for inducing and limiting meiotic DNA double-strand break formation. <i>Nucleic Acids Research</i> , 2018, 46, 11822-11834.  | 14.5 | 17        |
| 10 | E2f1 is a meiosis-specific transcription factor in the protist <i>Tetrahymena thermophila</i> . <i>Cell Cycle</i> , 2017, 16, 123-135.   | 2.6  | 9         |
| 11 | Nonsense-mediated mRNA decay in <i>Tetrahymena</i> is EJC independent and requires a protozoa-specific nuclease. <i>Nucleic Acids Research</i> , 2017, 45, 6848-6863.  | 14.5 | 22        |
| 12 | Cyc17, a meiosis-specific cyclin, is essential for anaphase initiation and chromosome segregation in <i>Tetrahymena thermophila</i> . <i>Cell Cycle</i> , 2016, 15, 1855-1864.   | 2.6  | 17        |
| 13 | Cdk3, a conjugation-specific cyclin-dependent kinase, is essential for the initiation of meiosis in <i>Tetrahymena thermophila</i> . <i>Cell Cycle</i> , 2016, 15, 2506-2514.  | 2.6  | 17        |
| 14 | Genome of the facultative scuticociliatosis pathogen <i>Pseudocohnilembus persalinus</i> provides insight into its virulence through horizontal gene transfer. <i>Scientific Reports</i> , 2015, 5, 15470.   | 3.3  | 46        |
| 15 | Phylogenomic analyses reveal subclass Scuticociliatia as the sister group of subclass Hymenostomatia within class Oligohymenophorea. <i>Molecular Phylogenetics and Evolution</i> , 2015, 90, 104-111.   | 2.7  | 37        |
| 16 | Phosphoproteomic Analysis of Protein Phosphorylation Networks in <i>Tetrahymena thermophila</i> , a Model Single-celled Organism. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 503-519.  | 3.8  | 21        |
| 17 | <i>Tetrahymena</i> Functional Genomics Database (TetraFGD): an integrated resource for <i>Tetrahymena</i> functional genomics. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat008.                                     | 3.0  | 51        |
| 18 | Transcriptome Analysis of the Model Protozoan, <i>Tetrahymena thermophila</i> , Using Deep RNA Sequencing. <i>PLoS ONE</i> , 2012, 7, e30630.  | 2.5  | 111       |