# Wei-Jen Chang 

## List of Publications by Year

 in descending orderSource: https:/|exaly.com/author-pdf/6254974/publications.pdf
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| 1 | The Oxytricha trifallax Macronuclear Genome: A Complex Eukaryotic Cenome with 16,000 Tiny Chromosomes. PLoS Biology, 2013, 11, e1001473. | 5.6 | 198 |
| :---: | :---: | :---: | :---: |
| 2 | Development of zebrafish epidermis. Birth Defects Research Part C: Embryo Today Reviews, 2011, 93, 205-214. | 3.6 | 79 |
| 3 | The evolutionary history of histone H3 suggests a deep eukaryotic root of chromatin modifying mechanisms. BMC Evolutionary Biology, 2010, 10, 259. | 3.2 | 75 |
| 4 | The transcription factor, glial cell missing 2 , is involved in differentiation and functional regulation of H<sup>+</sup>-ATPase-rich cells in zebrafish (<i>Danio rerio</i>). American Journal of Physiology Regulatory Integrative and Comparative Physiology, 2009, 296, R1192-R1201. | 1.8 | 56 |
| 5 | RNA-dependent control of gene amplification. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 22134-22139. | 7.1 | 50 |
| 6 | The evolutionary origin of a complex scrambled gene. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 15149-15154. | 7.1 | 47 |
| 7 | Compensatory regulation of $\mathrm{Na}+$ absorption by $\mathrm{Na}+/ \mathrm{H}+$ exchanger and $\mathrm{Na}+-\mathrm{Cl}-$ cotransporter in zebrafish (Danio rerio). Frontiers in Zoology, 2013, 10, 46. | 2.0 | 43 |

8 The Pathway to Detangle a Scrambled Gene. PLoS ONE, 2008, 3, e2330. 2.5
Macronuclear genome structure of the ciliate Nyctotherus ovalis: Single-gene chromosomes and tiny
introns. BMC Genomics, 2008, 9,587 .
Copy number variations of 11 macronuclear chromosomes and their gene expression in Oxytricha
trifallax. Gene, 2012, 505, 75-80.
11 A Micronuclear Locus Containing Three Protein-Coding Genes Remains Linked During Macronuclear Development in the Spirotrichous Ciliate Holosticha. Protist, 2004, 155, 245-255. ..... 1.5

14 Molecular genetic diversity and characterization of conjugation genes in the fish parasite Ichthyophthirius multifiliis. Molecular Phylogenetics and Evolution, 2015, 86, 1-7.
2.7

SIGAR: Inferring Features of Genome Architecture and DNA Rearrangements by Split-Read Mapping.

