Aaron P Turkewitz

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#	Paper	IF	Citations
48	Macronuclear genome sequence of the ciliate Tetrahymena thermophila, a model eukaryote. <i>PLoS Biology</i> , 2006 , 4, e286	9.7	544
47	Functional genomics: the coming of age for Tetrahymena thermophila. <i>Trends in Genetics</i> , 2002 , 18, 35-	- 40 .5	94
46	Elucidation of clathrin-mediated endocytosis in tetrahymena reveals an evolutionarily convergent recruitment of dynamin. <i>PLoS Genetics</i> , 2005 , 1, e52	6	82
45	Evolutionary cell biology: two origins, one objective. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 16990-4	11.5	75
44	Comprehensive analysis reveals dynamic and evolutionary plasticity of Rab GTPases and membrane traffic in Tetrahymena thermophila. <i>PLoS Genetics</i> , 2010 , 6, e1001155	6	63
43	Whole-cell biosensors for detection of heavy metal ions in environmental samples based on metallothionein promoters from Tetrahymena thermophila. <i>Microbial Biotechnology</i> , 2011 , 4, 513-22	6.3	61
42	Out with a bang! Tetrahymena as a model system to study secretory granule biogenesis. <i>Traffic</i> , 2004 , 5, 63-8	5.7	55
41	Genetic tool development in marine protists: emerging model organisms for experimental cell biology. <i>Nature Methods</i> , 2020 , 17, 481-494	21.6	39
40	Functional GFP-metallothionein fusion protein from Tetrahymena thermophila: a potential whole-cell biosensor for monitoring heavy metal pollution and a cell model to study metallothionein overproduction effects. <i>BioMetals</i> , 2014 , 27, 195-205	3.4	38
39	Proteolytic processing and Ca2+-binding activity of dense-core vesicle polypeptides in Tetrahymena. <i>Molecular Biology of the Cell</i> , 1998 , 9, 497-511	3.5	37
38	Lysosomal sorting receptors are essential for secretory granule biogenesis in Tetrahymena. <i>Journal of Cell Biology</i> , 2013 , 203, 537-50	7.3	36
37	In vivo analysis of the major exocytosis-sensitive phosphoprotein in Tetrahymena. <i>Journal of Cell Biology</i> , 1997 , 139, 1197-207	7.3	29
36	Large-scale purification of murine I-Ak and I-Ek antigens and characterization of the purified proteins. <i>Molecular Immunology</i> , 1983 , 20, 1139-47	4.3	28
35	Analysis of expressed sequence tags (ESTs) in the ciliated protozoan Tetrahymena thermophila. <i>Journal of Eukaryotic Microbiology</i> , 2002 , 49, 99-107	3.6	27
34	Genomic and proteomic evidence for a second family of dense core granule cargo proteins in Tetrahymena thermophila. <i>Journal of Eukaryotic Microbiology</i> , 2005 , 52, 291-7	3.6	27
33	N-methyldeoxyadenosine directs nucleosome positioning in Tetrahymena DNA. <i>Genome Biology</i> , 2018 , 19, 200	18.3	26
32	Purification of murine MHC antigens by monoclonal antibody affinity chromatography. <i>Methods in Enzymology</i> , 1983 , 92, 86-109	1.7	25

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31	Genetic, genomic, and functional analysis of the granule lattice proteins in Tetrahymena secretory granules. <i>Molecular Biology of the Cell</i> , 2005 , 16, 4046-60	3.5	24	
30	Analysis of a mutant exhibiting conditional sorting to dense core secretory granules in Tetrahymena thermophila. <i>Genetics</i> , 2001 , 159, 1605-16	4	24	
29	Core formation and the acquisition of fusion competence are linked during secretory granule maturation in Tetrahymena. <i>Traffic</i> , 2005 , 6, 303-23	5.7	23	
28	A role for convergent evolution in the secretory life of cells. <i>Trends in Cell Biology</i> , 2007 , 17, 157-64	18.3	21	
27	Immunocytochemical analysis of secretion mutants of Tetrahymena using a mucocyst-specific monoclonal antibody. <i>Genesis</i> , 1992 , 13, 151-9		20	
26	An evolutionary balance: conservation vs innovation in ciliate membrane trafficking. <i>Traffic</i> , 2017 , 18, 18-28	5.7	19	
25	A dynamin-related protein required for nuclear remodeling in Tetrahymena. <i>Current Biology</i> , 2008 , 18, 1227-33	6.3	19	
24	New class of cargo protein in Tetrahymena thermophila dense core secretory granules. <i>Eukaryotic Cell</i> , 2002 , 1, 583-93		19	
23	Resolving the homology-function relationship through comparative genomics of membrane-trafficking machinery and parasite cell biology. <i>Molecular and Biochemical Parasitology</i> , 2016 , 209, 88-103	1.9	17	
22	The cytochrome b5 dependent C-5(6) sterol desaturase DES5A from the endoplasmic reticulum of Tetrahymena thermophila complements ergosterol biosynthesis mutants in Saccharomyces cerevisiae. <i>Steroids</i> , 2012 , 77, 1313-20	2.8	15	
21	Conservation and innovation in Tetrahymena membrane traffic: proteins, lipids, and compartments. <i>Methods in Cell Biology</i> , 2012 , 109, 141-75	1.8	15	
20	Remodeling the Specificity of an Endosomal CORVET Tether Underlies Formation of Regulated Secretory Vesicles in the Ciliate Tetrahymena thermophila. <i>Current Biology</i> , 2018 , 28, 697-710.e13	6.3	14	
19	Independent transport and sorting of functionally distinct protein families in Tetrahymena thermophila dense core secretory granules. <i>Eukaryotic Cell</i> , 2009 , 8, 1575-83		14	
18	Regulated protein secretion in Tetrahymena thermophila. <i>Methods in Cell Biology</i> , 2000 , 62, 347-62	1.8	14	
17	An endosomal syntaxin and the AP-3 complex are required for formation and maturation of candidate lysosome-related secretory organelles (mucocysts) in. <i>Molecular Biology of the Cell</i> , 2017 , 28, 1551-1564	3.5	13	
16	An aspartyl cathepsin, CTH3, is essential for proprotein processing during secretory granule maturation in Tetrahymena thermophila. <i>Molecular Biology of the Cell</i> , 2014 , 25, 2444-60	3.5	13	
15	Proprotein processing within secretory dense core granules of Tetrahymena thermophila. <i>Journal of Biological Chemistry</i> , 2003 , 278, 4087-95	5.4	13	
14	Extreme metal adapted, knockout and knockdown strains reveal a coordinated gene expression among different Tetrahymena thermophila metallothionein isoforms. <i>PLoS ONE</i> , 2017 , 12, e0189076	3.7	13	

13	An Alveolata secretory machinery adapted to parasite host cell invasion. <i>Nature Microbiology</i> , 2021 , 6, 425-434	26.6	13
12	Tetrahymena thermophila: a divergent perspective on membrane traffic. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , 2014 , 322, 500-16	1.8	12
11	Secretion of Polypeptide Crystals from Tetrahymena thermophila Secretory Organelles (Mucocysts) Depends on Processing by a Cysteine Cathepsin, Cth4p. <i>Eukaryotic Cell</i> , 2015 , 14, 817-33		10
10	Genome plasticity in response to stress in Tetrahymena thermophila: selective and reversible chromosome amplification and paralogous expansion of metallothionein genes. <i>Environmental Microbiology</i> , 2018 , 20, 2410-2421	5.2	9
9	A Rab-based view of membrane traffic in the ciliate Tetrahymena thermophila. <i>Small GTPases</i> , 2011 , 2, 222-226	2.7	8
8	Whole Genome Sequencing Identifies a Novel Factor Required for Secretory Granule Maturation in Tetrahymena thermophila. <i>G3: Genes, Genomes, Genetics</i> , 2016 , 6, 2505-16	3.2	8
7	Diversification of CORVET tethers facilitates transport complexity in. <i>Journal of Cell Science</i> , 2020 , 133,	5.3	5
6	The evolution of germ-soma nuclear differentiation in eukaryotic unicells. <i>Current Biology</i> , 2020 , 30, RS	50 8 -∕R51	105
5	ESCargo: a regulatable fluorescent secretory cargo for diverse model organisms. <i>Molecular Biology of the Cell</i> , 2020 , 31, 2892-2903	3.5	4
4	Biogenesis of Dense-Core Secretory Granules 2009 , 183-209		4
3	The Co-regulation Data Harvester: automating gene annotation starting from a transcriptome database. <i>SoftwareX</i> , 2017 , 6, 165-171	2.7	1
2	Stalking the wild Tetrahymena. <i>Molecular Ecology</i> , 2013 , 22, 912-4	5.7	1
1	A novel membrane complex is required for docking and regulated exocytosis of lysosome-related organelles in Tetrahymena thermophila <i>PLoS Genetics</i> , 2022 , 18, e1010194	6	0