

Tzumin Lee

List of Publications by Year in descending order

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Version: 2024-02-01

49
papers

7,930
citations

147566

31
h-index

197535

49
g-index

67
all docs

67
docs citations

67
times ranked

6235
citing authors

#	ARTICLE	IF	CITATIONS
1	Mosaic Analysis with a Repressible Cell Marker for Studies of Gene Function in Neuronal Morphogenesis. <i>Neuron</i> , 1999, 22, 451-461.	3.8	2,368
2	Optimized CRISPR/Cas tools for efficient germline and somatic genome engineering in <i>Drosophila</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2967-76.	3.3	947
3	Mosaic analysis with a repressible cell marker (MARCM) for <i>Drosophila</i> neural development. <i>Trends in Neurosciences</i> , 2001, 24, 251-254.	4.2	845
4	Genetic mosaic with dual binary transcriptional systems in <i>Drosophila</i> . <i>Nature Neuroscience</i> , 2006, 9, 703-709.	7.1	478
5	Organization and Postembryonic Development of Glial Cells in the Adult Central Brain of <i>Drosophila</i> . <i>Journal of Neuroscience</i> , 2008, 28, 13742-13753.	1.7	280
6	Sparse, decorrelated odor coding in the mushroom body enhances learned odor discrimination. <i>Nature Neuroscience</i> , 2014, 17, 559-568.	7.1	268
7	Cell-Autonomous Requirement of the USP/EcR-B Ecdysone Receptor for Mushroom Body Neuronal Remodeling in <i>Drosophila</i> . <i>Neuron</i> , 2000, 28, 807-818.	3.8	255
8	TGF- β Signaling Activates Steroid Hormone Receptor Expression during Neuronal Remodeling in the <i>Drosophila</i> Brain. <i>Cell</i> , 2003, 112, 303-315.	13.5	215
9	Gradients of the <i>Drosophila</i> Chinmo BTB-Zinc Finger Protein Govern Neuronal Temporal Identity. <i>Cell</i> , 2006, 127, 409-422.	13.5	213
10	Clonal analysis of <i>Drosophila</i> antennal lobe neurons: diverse neuronal architectures in the lateral neuroblast lineage. <i>Development (Cambridge)</i> , 2008, 135, 2883-2893.	1.2	182
11	Clonal Development and Organization of the Adult <i>Drosophila</i> Central Brain. <i>Current Biology</i> , 2013, 23, 633-643.	1.8	161
12	Twin-spot MARCM to reveal the developmental origin and identity of neurons. <i>Nature Neuroscience</i> , 2009, 12, 947-953.	7.1	149
13	A Complete Developmental Sequence of a <i>Drosophila</i> Neuronal Lineage as Revealed by Twin-Spot MARCM. <i>PLoS Biology</i> , 2010, 8, e1000461.	2.6	140
14	Opposing intrinsic temporal gradients guide neural stem cell production of varied neuronal fates. <i>Science</i> , 2015, 350, 317-320.	6.0	130
15	Stem Cell-Intrinsic, Seven-up-Triggered Temporal Factor Gradients Diversify Intermediate Neural Progenitors. <i>Current Biology</i> , 2017, 27, 1303-1313.	1.8	81
16	Neurotransmitter identity is acquired in a lineage-restricted manner in the <i>Drosophila</i> CNS. <i>ELife</i> , 2019, 8, .	2.8	78
17	Lineage Analysis of <i>Drosophila</i> Lateral Antennal Lobe Neurons Reveals Notch-Dependent Binary Temporal Fate Decisions. <i>PLoS Biology</i> , 2012, 10, e1001425.	2.6	67
18	Diverse neuronal lineages make stereotyped contributions to the <i>Drosophila</i> locomotor control center, the central complex. <i>Journal of Comparative Neurology</i> , 2013, 521, 2645-2662.	0.9	67

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19	Cell Class-Lineage Analysis Reveals Sexually Dimorphic Lineage Compositions in the <i>Drosophila</i> Brain. <i>Current Biology</i> , 2016, 26, 2583-2593.	1.8	67
20	Lineage-specific effects of Notch/Numb signaling in post-embryonic development of the <i>Drosophila</i> brain. <i>Development (Cambridge)</i> , 2010, 137, 43-51.	1.2	62
21	Imp and Syp RNA-binding proteins govern decommissioning of <i>Drosophila</i> neural stem cells. <i>Development (Cambridge)</i> , 2017, 144, 3454-3464.	1.2	62
22	Making <i>Drosophila</i> lineage-restricted drivers via patterned recombination in neuroblasts. <i>Nature Neuroscience</i> , 2014, 17, 631-637.	7.1	57
23	Dissection of the <i>Drosophila</i> neuropeptide F circuit using a high-throughput two-choice assay. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E8091-E8099.	3.3	55
24	<i>Drosophila</i> intermediate neural progenitors produce lineage-dependent related series of diverse neurons. <i>Development (Cambridge)</i> , 2014, 141, 253-258.	1.2	52
25	Transcriptomes of lineage-specific <i>Drosophila</i> neuroblasts profiled via genetic targeting and robotic sorting. <i>Development (Cambridge)</i> , 2015, 143, 411-21.	1.2	49
26	Temporal control of <i>Drosophila</i> central nervous system development. <i>Current Opinion in Neurobiology</i> , 2019, 56, 24-32.	2.0	47
27	Generating neuronal diversity in the <i>Drosophila</i> central nervous system. <i>Developmental Dynamics</i> , 2012, 241, 57-68.	0.8	45
28	Hierarchical Deployment of Factors Regulating Temporal Fate in a Diverse Neuronal Lineage of the <i>Drosophila</i> Central Brain. <i>Neuron</i> , 2012, 73, 677-684.	3.8	44
29	The bHLH Repressor Deadpan Regulates the Self-renewal and Specification of <i>Drosophila</i> Larval Neural Stem Cells Independently of Notch. <i>PLoS ONE</i> , 2012, 7, e46724.	1.1	44
30	Nuclear Receptor Unfulfilled Regulates Axonal Guidance and Cell Identity of <i>Drosophila</i> Mushroom Body Neurons. <i>PLoS ONE</i> , 2009, 4, e8392.	1.1	43
31	Extremes of Lineage Plasticity in the <i>Drosophila</i> Brain. <i>Current Biology</i> , 2013, 23, 1908-1913.	1.8	43
32	Birth time/order-dependent neuron type specification. <i>Current Opinion in Neurobiology</i> , 2010, 20, 14-21.	2.0	32
33	Unlimited Genetic Switches for Cell-Type-Specific Manipulation. <i>Neuron</i> , 2019, 104, 227-238.e7.	3.8	29
34	Conservation and divergence of related neuronal lineages in the <i>Drosophila</i> central brain. <i>ELife</i> , 2020, 9, .	2.8	29
35	An Enhanced Gene Targeting Toolkit for <i>Drosophila</i> : Golic+. <i>Genetics</i> , 2015, 199, 683-694.	1.2	28
36	Mamo decodes hierarchical temporal gradients into terminal neuronal fate. <i>ELife</i> , 2019, 8, .	2.8	23

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37	Wiring the Drosophila Brain with Individually Tailored Neural Lineages. <i>Current Biology</i> , 2017, 27, R77-R82.	1.8	21
38	Lineage-guided Notch-dependent gliogenesis by <i>Drosophila</i> multi-potent progenitors. <i>Development (Cambridge)</i> , 2018, 145, .	1.2	21
39	High-throughput dense reconstruction of cell lineages. <i>Open Biology</i> , 2019, 9, 190229.	1.5	21
40	Generating mosaics for lineage analysis in flies. <i>Wiley Interdisciplinary Reviews: Developmental Biology</i> , 2014, 3, 69-81.	5.9	20
41	A programmable sequence of reporters for lineage analysis. <i>Nature Neuroscience</i> , 2020, 23, 1618-1628.	7.1	18
42	Neuronal upregulation of Prospero protein is driven by alternative mRNA polyadenylation and Syncrip-mediated mRNA stabilisation. <i>Biology Open</i> , 2020, 9, .	0.6	14
43	New genetic tools for cell lineage analysis in <i>Drosophila</i> . <i>Nature Methods</i> , 2009, 6, 566-568.	9.0	11
44	The art of lineage tracing: From worm to human. <i>Progress in Neurobiology</i> , 2021, 199, 101966.	2.8	9
45	Enhanced Golic+: Highly effective CRISPR gene targeting and transgene HACKing in <i>Drosophila</i> . <i>Development (Cambridge)</i> , 2020, 147, .	1.2	6
46	YAP1 nuclear efflux and transcriptional reprogramming follow membrane diminution upon VSV-G-induced cell fusion. <i>Nature Communications</i> , 2021, 12, 4502.	5.8	5
47	Hormone-controlled changes in the differentiation state of post-mitotic neurons. <i>Current Biology</i> , 2022, , .	1.8	4
48	Diverse neuronal lineages make stereotyped contributions to the <i>Drosophila</i> locomotor control center, the central complex. <i>Journal of Comparative Neurology</i> , 2013, 521, Spc1-Spc1.	0.9	3
49	CAMIO: a transgenic CRISPR pipeline to create diverse targeted genome deletions in <i>Drosophila</i> . <i>Nucleic Acids Research</i> , 2020, 48, 4344-4356.	6.5	3