

Zhe Han

List of Publications by Year in descending order

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Version: 2024-02-01

53
papers

2,856
citations

257450

24
h-index

182427

51
g-index

61
all docs

61
docs citations

61
times ranked

3664
citing authors

#	ARTICLE	IF	CITATIONS
1	Pharmacological or genetic inhibition of hypoxia signaling attenuates oncogenic RAS-induced cancer phenotypes. <i>DMM Disease Models and Mechanisms</i> , 2022, 15, .	2.4	6
2	Using <i>Drosophila</i> Nephrocytes to Understand the Formation and Maintenance of the Podocyte Slit Diaphragm. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, 837828.	3.7	3
3	Lpt, trr, and Hcf regulate histone mono- and dimethylation that are essential for <i>Drosophila</i> heart development. <i>Developmental Biology</i> , 2022, 490, 53-65.	2.0	4
4	Autophagy inhibition rescues structural and functional defects caused by the loss of mitochondrial chaperone <i>Hsc70-5</i> in <i>Drosophila</i> . <i>Autophagy</i> , 2021, 17, 3160-3174.	9.1	5
5	Exome Sequencing and Congenital Heart Disease in Sub-Saharan Africa. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003108.	3.6	16
6	Heterozygosity for a Pathogenic Variant in <i>SLC12A3</i> That Causes Autosomal Recessive Gitelman Syndrome Is Associated with Lower Serum Potassium. <i>Journal of the American Society of Nephrology: JASN</i> , 2021, 32, 756-765.	6.1	11
7	Inactivating histone deacetylase HDA promotes longevity by mobilizing trehalose metabolism. <i>Nature Communications</i> , 2021, 12, 1981.	12.8	29
8	Characterization of SARS-CoV-2 proteins reveals Orf6 pathogenicity, subcellular localization, host interactions and attenuation by Selinexor. <i>Cell and Bioscience</i> , 2021, 11, 58.	4.8	92
9	Functional analysis of SARS-CoV-2 proteins in <i>Drosophila</i> identifies Orf6-induced pathogenic effects with Selinexor as an effective treatment. <i>Cell and Bioscience</i> , 2021, 11, 59.	4.8	18
10	Slit diaphragm maintenance requires dynamic clathrin-mediated endocytosis facilitated by AP-2, Lap, Aux and Hsc70-4 in nephrocytes. <i>Cell and Bioscience</i> , 2021, 11, 83.	4.8	13
11	<i>Drosophila</i> , a powerful model to study virus-host interactions and pathogenicity in the fight against SARS-CoV-2. <i>Cell and Bioscience</i> , 2021, 11, 110.	4.8	12
12	Understanding Individual SARS-CoV-2 Proteins for Targeted Drug Development against COVID-19. <i>Molecular and Cellular Biology</i> , 2021, 41, e0018521.	2.3	21
13	Phosphorylation of slit diaphragm proteins NEPHRIN and NEPH1 upon binding of HGF promotes podocyte repair. <i>Journal of Biological Chemistry</i> , 2021, 297, 101079.	3.4	4
14	Novel frameshift variant in <i>MYL2</i> reveals molecular differences between dominant and recessive forms of hypertrophic cardiomyopathy. <i>PLoS Genetics</i> , 2020, 16, e1008639.	3.5	16
15	Single-cell RNA sequencing identifies novel cell types in <i>Drosophila</i> blood. <i>Journal of Genetics and Genomics</i> , 2020, 47, 175-186.	3.9	73
16	Zika virus non-structural protein NS4A restricts eye growth in <i>Drosophila</i> through regulation of JAK/STAT signaling. <i>DMM Disease Models and Mechanisms</i> , 2020, 13, .	2.4	22
17	Exocyst Genes Are Essential for Recycling Membrane Proteins and Maintaining Slit Diaphragm in <i>Drosophila</i> Nephrocytes. <i>Journal of the American Society of Nephrology: JASN</i> , 2020, 31, 1024-1034.	6.1	12
18	Master regulator genes and their impact on major diseases. <i>PeerJ</i> , 2020, 8, e9952.	2.0	19

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19	Forward genetic screen in human podocytes identifies diphthamide biosynthesis genes as regulators of adhesion. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 317, F1593-F1604.	2.7	4
20	Mutations in NUP160 Are Implicated in Steroid-Resistant Nephrotic Syndrome. <i>Journal of the American Society of Nephrology: JASN</i> , 2019, 30, 840-853.	6.1	21
21	APOL1 risk allele RNA contributes to renal toxicity by activating protein kinase R. <i>Communications Biology</i> , 2018, 1, 188.	4.4	59
22	Molecular mechanisms of heart failure: insights from <i>Drosophila</i> . <i>Heart Failure Reviews</i> , 2017, 22, 91-98.	3.9	18
23	Comprehensive functional analysis of Rab GTPases in <i>Drosophila</i> nephrocytes. <i>Cell and Tissue Research</i> , 2017, 368, 615-627.	2.9	40
24	The E3 ubiquitin ligase Nedd4/Nedd4L is directly regulated by microRNA 1. <i>Development (Cambridge)</i> , 2017, 144, 866-875.	2.5	18
25	A Personalized Model of COQ2 Nephropathy Rescued by the Wild-Type COQ2 Allele or Dietary Coenzyme Q10 Supplementation. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 2607-2617.	6.1	15
26	A <i>Drosophila</i> model system to assess the function of human monogenic podocyte mutations that cause nephrotic syndrome. <i>Human Molecular Genetics</i> , 2017, 26, 768-780.	2.9	26
27	Transmembrane TNF- α Facilitates HIV-1 Infection of Podocytes Cultured from Children with HIV-Associated Nephropathy. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 862-875.	6.1	22
28	APOL1-G1 in Nephrocytes Induces Hypertrophy and Accelerates Cell Death. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 1106-1116.	6.1	66
29	Epigenetic mechanisms underlying maternal diabetes-associated risk of congenital heart disease. <i>JCI Insight</i> , 2017, 2, .	5.0	59
30	Validating Candidate Congenital Heart Disease Genes in <i>Drosophila</i> . <i>Bio-protocol</i> , 2017, 7, .	0.4	10
31	High throughput in vivo functional validation of candidate congenital heart disease genes in <i>Drosophila</i> . <i>ELife</i> , 2017, 6, .	6.0	41
32	The E3 ubiquitin ligase Nedd4/Nedd4L is directly regulated by microRNA 1. <i>Journal of Cell Science</i> , 2017, 130, e1.2-e1.2.	2.0	0
33	Wnt4 is required for ostia development in the <i>Drosophila</i> heart. <i>Developmental Biology</i> , 2016, 413, 188-198.	2.0	13
34	Gia/Mth5 is an aorta specific GPCR required for <i>Drosophila</i> heart tube morphology and normal pericardial cell positioning. <i>Developmental Biology</i> , 2016, 414, 100-107.	2.0	10
35	A transgenic resource for conditional competitive inhibition of conserved <i>Drosophila</i> microRNAs. <i>Nature Communications</i> , 2015, 6, 7279.	12.8	63
36	KANK deficiency leads to podocyte dysfunction and nephrotic syndrome. <i>Journal of Clinical Investigation</i> , 2015, 125, 2375-2384.	8.2	159

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37	Cubilin and Amnionless Mediate Protein Reabsorption in <i>Drosophila</i> Nephrocytes. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 209-216.	6.1	98
38	An In Vivo Functional Analysis System for Renal Gene Discovery in <i>Drosophila</i> Pericardial Nephrocytes. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 191-197.	6.1	92
39	ADCK4 mutations promote steroid-resistant nephrotic syndrome through CoQ10 biosynthesis disruption. <i>Journal of Clinical Investigation</i> , 2013, 123, 5179-5189.	8.2	275
40	ARHGDI1 mutations cause nephrotic syndrome via defective RHO GTPase signaling. <i>Journal of Clinical Investigation</i> , 2013, 123, 3243-3253.	8.2	196
41	miR-92b regulates Mef2 levels through a negative-feedback circuit during <i>Drosophila</i> muscle development. <i>Development (Cambridge)</i> , 2012, 139, 3543-3552.	2.5	49
42	Spatial specificity of mesodermal even-skipped expression relies on multiple repressor sites. <i>Developmental Biology</i> , 2008, 313, 876-886.	2.0	9
43	Palisade is required in the <i>Drosophila</i> ovary for assembly and function of the protective vitelline membrane. <i>Developmental Biology</i> , 2008, 319, 359-369.	2.0	17
44	Heterotrimeric G Proteins Regulate a Noncanonical Function of Septate Junction Proteins to Maintain Cardiac Integrity in <i>Drosophila</i> . <i>Developmental Cell</i> , 2008, 15, 704-713.	7.0	50
45	The Him Gene Reveals a Balance of Inputs Controlling Muscle Differentiation in <i>Drosophila</i> . <i>Current Biology</i> , 2007, 17, 1409-1413.	3.9	33
46	Hand, an evolutionarily conserved bHLH transcription factor required for <i>Drosophila</i> cardiogenesis and hematopoiesis. <i>Development (Cambridge)</i> , 2006, 133, 1175-1182.	2.5	104
47	The Mevalonate Pathway Controls Heart Formation in <i>Drosophila</i> by Isoprenylation of GÅ1. <i>Science</i> , 2006, 313, 1301-1303.	12.6	83
48	Hand is a direct target of Tinman and GATA factors during <i>Drosophila</i> cardiogenesis and hematopoiesis. <i>Development (Cambridge)</i> , 2005, 132, 3525-3536.	2.5	131
49	Embryonic even-skipped-dependent muscle and heart cell fates are required for normal adult activity, heart function, and lifespan. <i>Circulation Research</i> , 2005, 97, 1108-1114.	4.5	37
50	MicroRNA1 influences cardiac differentiation in <i>Drosophila</i> and regulates Notch signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 18986-18991.	7.1	411
51	A myocardin-related transcription factor regulates activity of serum response factor in <i>Drosophila</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 12567-12572.	7.1	68
52	Myogenic cell fates are antagonized by Notch only in asymmetric lineages of the <i>Drosophila</i> heart, with or without cell division. <i>Development (Cambridge)</i> , 2003, 130, 3039-3051.	2.5	89
53	Transcriptional integration of competence modulated by mutual repression generates cell-type specificity within the cardiogenic mesoderm. <i>Developmental Biology</i> , 2002, 252, 225-240.	2.0	57