

# Xiaoqing Zhang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3404020/publications.pdf>

Version: 2024-02-01

19  
papers

243  
citations

1307594

7  
h-index

996975

15  
g-index

21  
all docs

21  
docs citations

21  
times ranked

587  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical features, laboratory findings and persistence of virus in 10 children with coronavirus disease 2019 (COVID-19). <i>Biomedical Journal</i> , 2021, 44, 94-100.	3.1	8
2	Identification of the genetic basis of sporadic polydactyly in China by targeted sequencing. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 3482-3490.	4.1	0
3	FGD5-AS1 Is a Hub lncRNA ceRNA in Hearts With Tetralogy of Fallot Which Regulates Congenital Heart Disease Genes Transcriptionally and Epigenetically. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 630634.	3.7	13
4	Long Noncoding RNA lnc-TSSK2-8 Activates Canonical Wnt/ $\beta$ 2-Catenin Signaling Through Small Heat Shock Proteins HSPA6 and CRYAB. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 660576.	3.7	7
5	SORBS2 is a genetic factor contributing to cardiac malformation of 4q deletion syndrome patients. <i>ELife</i> , 2021, 10, .	6.0	9
6	Generation of a homozygous CRISPR/Cas9-mediated knockout human iPSC line for PTCH1 gene. <i>Stem Cell Research</i> , 2021, 56, 102517.	0.7	0
7	Copy number variation analysis in Chinese children with complete atrioventricular canal and single ventricle. <i>BMC Medical Genomics</i> , 2021, 14, 243.	1.5	0
8	A Novel Nonsense GLI3 Variant Is Associated With Polydactyly and Syndactyly in a Family by Blocking the Sonic Hedgehog Signaling Pathway. <i>Frontiers in Genetics</i> , 2020, 11, 542004.	2.3	5
9	Rare Copy Number Variations Might Not be Involved in the Molecular Pathogenesis of PA $\alpha$ IVS in an Unselected Chinese Cohort. <i>Pediatric Cardiology</i> , 2019, 40, 762-767.	1.3	3
10	Intracellular osteopontin negatively regulates toll-like receptor 4-mediated inflammatory response via regulating GSK3 $\beta$ and 4EBP1 phosphorylation. <i>Cytokine</i> , 2018, 108, 89-95.	3.2	13
11	A novel dNTP-limited PCR and HRM assay to detect Williams-Beuren syndrome. <i>Clinica Chimica Acta</i> , 2018, 481, 171-176.	1.1	1
12	A loss-of-function mutation p.T52S in RIPPLY3 is a potential predisposing genetic risk factor for Chinese Han conotruncal heart defect patients without the 22q11.2 deletion/duplication. <i>Journal of Translational Medicine</i> , 2018, 16, 260.	4.4	3
13	Accurate diagnosis of spinal muscular atrophy and 22q11.2 deletion syndrome using limited deoxynucleotide triphosphates and high-resolution melting. <i>BMC Genomics</i> , 2018, 19, 485.	2.8	7
14	A Novel Multiplex HRM Assay to Detect Clopidogrel Resistance. <i>Scientific Reports</i> , 2017, 7, 16021.	3.3	6
15	A Novel Missense Mutation of GATA4 in a Chinese Family with Congenital Heart Disease. <i>PLoS ONE</i> , 2016, 11, e0158904.	2.5	19
16	Three novel missense mutations in the filamin B gene are associated with isolated congenital talipes equinovarus. <i>Human Genetics</i> , 2016, 135, 1181-1189.	3.8	22
17	A novel missense mutation of <i>TNNI2</i> in a Chinese family cause distal arthrogyriposis type 1. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 135-141.	1.2	8
18	A modified multiplex ligation-dependent probe amplification method for the detection of 22q11.2 copy number variations in patients with congenital heart disease. <i>BMC Genomics</i> , 2015, 16, 364.	2.8	44

#	ARTICLE	IF	CITATIONS
19	Chromosome microarray testing for patients with congenital heart defects reveals novel disease causing loci and high diagnostic yield. BMC Genomics, 2014, 15, 1127.	2.8	74