Lili Hao

List of Publications by Year
in descending order

11 papers	119 citations	1306789 7 h-index	1281420 11 g-index
11 all docs	11 docs citations	11 times ranked	199 citing authors

#	Article	IF	Citations
1	Multiple gene variations contributed to congenital heart disease via GATA family transcriptional regulation. Journal of Translational Medicine, 2017, 15, 69.	1.8	22
2	The roles of SMYD4 in epigenetic regulation of cardiac development in zebrafish. PLoS Genetics, 2018, 14, e1007578.	1.5	17
3	Early onset developmental delay and epilepsy in pediatric patients with WDR45 variants. European Journal of Medical Genetics, 2019, 62, 149-160.	0.7	16
4	Hypermethylationâ€mediated downâ€regulation of lncRNA TBX5â€AS1:2 in Tetralogy of Fallot inhibits cell proliferation by reducingTBX5expression. Journal of Cellular and Molecular Medicine, 2020, 24, 6472-6484.	1.6	16
5	Hypomethylation and decreased expression of BRG1 in the myocardium of patients with congenital heart disease. Birth Defects Research, 2017, 109, 1183-1195.	0.8	14
6	Exome sequencing identifies a mutation in TMC1 as a novel cause of autosomal recessive nonsyndromic hearing loss. Journal of Translational Medicine, 2016, 14, 29.	1.8	10
7	Two novel ANK1 lossâ€ofâ€function mutations in Chinese families with hereditary spherocytosis. Journal of Cellular and Molecular Medicine, 2019, 23, 4454-4463.	1.6	10
8	Long non-coding RNA SAP30-2:1 is downregulated in congenital heart disease and regulates cell proliferation by targeting HAND2. Frontiers of Medicine, 2021, 15, 91-100.	1.5	7
9	Expression Profiles of Exosomal MicroRNAs Derived from Cerebrospinal Fluid in Patients with Congenital Hydrocephalus Determined by MicroRNA Sequencing. Disease Markers, 2022, 2022, 1-16.	0.6	3
10	WDR62 variants contribute to congenital heart disease by inhibiting cardiomyocyte proliferation. Clinical and Translational Medicine, 2022, 12, .	1.7	3
11	The Novel Pathogenic Mutation c.849dupT in BRCA2 Contributes to the Nonsense-Mediated mRNA Decay of BRCA2 in Familial Breast Cancer. Journal of Breast Cancer, 2018, 21, 330.	0.8	1