

Shengwen Huang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7369379/publications.pdf>

Version: 2024-02-01

9
papers

54
citations

2258059

3
h-index

1720034

7
g-index

14
all docs

14
docs citations

14
times ranked

69
citing authors

#	ARTICLE	IF	CITATIONS
1	CRISPR/Cas9-based multiplex genome editing of BCL11A and HBG efficiently induces fetal hemoglobin expression. <i>European Journal of Pharmacology</i> , 2022, 918, 174788.	3.5	10
2	Comparison of transcriptome profiles of nucleated red blood cells in cord blood between preterm and full-term neonates. <i>Hematology</i> , 2022, 27, 263-273.	1.5	2
3	Two novel mutations in PADI6 and TLE6 genes cause female infertility due to arrest in embryonic development. <i>Journal of Assisted Reproduction and Genetics</i> , 2021, 38, 1551-1559.	2.5	17
4	Identification of two CUL7 variants in two Chinese families with 3 α -M syndrome by whole ϵ xome sequencing. <i>Journal of Clinical Laboratory Analysis</i> , 2020, 34, e23265.	2.1	3
5	Association of polymorphisms in the HBG1 ϵ HBD intergenic region with HbF levels. <i>Journal of Clinical Laboratory Analysis</i> , 2020, 34, e23243.	2.1	2
6	Long noncoding RNA PCED1B-AS1 promotes erythroid differentiation coordinating with GATA1 and chromatin remodeling. <i>Blood Science</i> , 2019, 1, 161-167.	0.9	3
7	Molecular newborn screening of four genetic diseases in Guizhou Province of South China. <i>Gene</i> , 2016, 591, 119-122.	2.2	12
8	Association between apolipoprotein E gene polymorphism and the dose for warfarin maintenance. <i>Journal of Central South University (Medical Sciences)</i> , 2011, 36, 212-6.	0.1	3
9	Whole ϵ xome sequencing identified five novel <i>de novo</i> variants in patients with unexplained intellectual disability. <i>Journal of Clinical Laboratory Analysis</i> , 0, , .	2.1	1