

Qiuming Yao

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2382461/publications.pdf>

Version: 2024-02-01

16
papers

828
citations

1162889

8
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1058333

14
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17
all docs

17
docs citations

17
times ranked

1390
citing authors

#	ARTICLE	IF	CITATIONS
1	Deletion of murine <i>Rho</i> h leads to de-repression of <i>Bcl-6</i> via decreased KAISO levels and accelerates a malignancy phenotype in a murine model of lymphoma. <i>Small GTPases</i> , 2022, 13, 267-281.	0.7	3
2	Epigenetic Alterations in Keratinocyte Carcinoma. <i>Journal of Investigative Dermatology</i> , 2021, 141, 1207-1218.	0.3	9
3	Motif-Raptor: a cell type-specific and transcription factor centric approach for post-GWAS prioritization of causal regulators. <i>Bioinformatics</i> , 2021, 37, 2103-2111.	1.8	5
4	ZNF410 represses fetal globin by singular control of CHD4. <i>Nature Genetics</i> , 2021, 53, 719-728.	9.4	35
5	Dissecting ELANE neutropenia pathogenicity by human HSC gene editing. <i>Cell Stem Cell</i> , 2021, 28, 833-845.e5.	5.2	23
6	Augmenting and directing long-range CRISPR-mediated activation in human cells. <i>Nature Methods</i> , 2021, 18, 1075-1081.	9.0	17
7	Therapeutic base editing of human hematopoietic stem cells. <i>Nature Medicine</i> , 2020, 26, 535-541.	15.2	196
8	ZNF410 Represses Fetal Globin By Devoted Control of CHD4/NuRD. <i>Blood</i> , 2020, 136, 1-1.	0.6	0
9	Rational targeting of a NuRD subcomplex guided by comprehensive in situ mutagenesis. <i>Nature Genetics</i> , 2019, 51, 1149-1159.	9.4	83
10	Editing aberrant splice sites efficiently restores $\hat{\beta}^2$ -globin expression in $\hat{\beta}^2$ -thalassemia. <i>Blood</i> , 2019, 133, 2255-2262.	0.6	57
11	Highly efficient therapeutic gene editing of human hematopoietic stem cells. <i>Nature Medicine</i> , 2019, 25, 776-783.	15.2	344
12	Gene Editing ELANE in Human Hematopoietic Stem and Progenitor Cells Reveals Disease Mechanisms and Therapeutic Strategies for Severe Congenital Neutropenia. <i>Blood</i> , 2019, 134, 3-3.	0.6	8
13	CRISPRO: identification of functional protein coding sequences based on genome editing dense mutagenesis. <i>Genome Biology</i> , 2018, 19, 169.	3.8	34
14	Highly Efficient Therapeutic Gene Editing of BCL11A enhancer in Human Hematopoietic Stem Cells from $\hat{\Delta}^{\gamma}$ -Hemoglobinopathy Patients for Fetal Hemoglobin Induction. <i>Blood</i> , 2018, 132, 3482-3482.	0.6	2
15	Comprehensive Integrated Genomic Perturbations Reveal Molecular Mechanisms of Red Blood Cell Trait Associations. <i>Blood</i> , 2018, 132, 532-532.	0.6	0
16	Multi-resolution based hybrid spatiotemporal compression of encrypted videos. , 2009, , .		9