

Zhuo Chen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4082335/publications.pdf>

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9
papers

961
citations

1040056

9
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

1612
citing authors

#	ARTICLE	IF	CITATIONS
1	miR-379 deletion ameliorates features of diabetic kidney disease by enhancing adaptive mitophagy via FIS1. <i>Communications Biology</i> , 2021, 4, 30.	4.4	20
2	LncRNA DRAIR is downregulated in diabetic monocytes and modulates inflammatory phenotype via epigenetic mechanisms. <i>JCI Insight</i> , 2021, 6, .	5.0	15
3	DNA methylation mediates development of HbA1c-associated complications in type 1 diabetes. <i>Nature Metabolism</i> , 2020, 2, 744-762.	11.9	53
4	Novel Long Noncoding RNA, Macrophage Inflammation-Suppressing Transcript (<i>MIST</i>), Regulates Macrophage Activation During Obesity. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 914-928.	2.4	32
5	Regulation of angiotensin II actions by enhancers and super-enhancers in vascular smooth muscle cells. <i>Nature Communications</i> , 2017, 8, 1467.	12.8	89
6	Epigenomic profiling reveals an association between persistence of DNA methylation and metabolic memory in the DCCT/EDIC type 1 diabetes cohort. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E3002-11.	7.1	179
7	An endoplasmic reticulum stress-regulated lncRNA hosting a microRNA megacluster induces early features of diabetic nephropathy. <i>Nature Communications</i> , 2016, 7, 12864.	12.8	206
8	Evaluating the Role of Epigenetic Histone Modifications in the Metabolic Memory of Type 1 Diabetes. <i>Diabetes</i> , 2014, 63, 1748-1762.	0.6	208
9	Regulation of Inflammatory Phenotype in Macrophages by a Diabetes-Induced Long Noncoding RNA. <i>Diabetes</i> , 2014, 63, 4249-4261.	0.6	155