## Tianyuan Lu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5107424/publications.pdf

Version: 2024-02-01

		1039406	996533
25	293	9	15
papers	citations	h-index	g-index
32	32	32	301
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	H3K27M in Gliomas Causes a One-Step Decrease in H3K27 Methylation and Reduced Spreading within the Constraints of H3K36 Methylation. Cell Reports, 2020, 33, 108390.	2.9	50
2	Individuals with common diseases but with a low polygenic risk score could be prioritized for rare variant screening. Genetics in Medicine, 2021, 23, 508-515.	1.1	39
3	Improved prediction of fracture risk leveraging a genome-wide polygenic risk score. Genome Medicine, 2021, 13, 16.	3.6	35
4	Reshuffling of the ancestral core-eudicot genome shaped chromatin topology and epigenetic modification in Panax. Nature Communications, 2022, 13, 1902.	5.8	30
5	Polygenic risk for coronary heart disease acts through atherosclerosis in type 2 diabetes. Cardiovascular Diabetology, 2020, 19, 12.	2.7	23
6	Whole-genome bisulfite sequencing in systemic sclerosis provides novel targets to understand disease pathogenesis. BMC Medical Genomics, 2019, 12, 144.	0.7	22
7	Investigating transcriptome-wide sex dimorphism by multi-level analysis of single-cell RNA sequencing data in ten mouse cell types. Biology of Sex Differences, 2020, 11, 61.	1.8	19
8	A Polygenic Risk Score to Predict Future Adult Short Stature Among Children. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 1918-1928.	1.8	19
9	Polygenic risk score as a possible tool for identifying familial monogenic causes of complex diseases. Genetics in Medicine, 2022, 24, 1545-1555.	1.1	12
10	Rapid Divergence Followed by Adaptation to Contrasting Ecological Niches of Two Closely Related Columbine SpeciesAquilegia japonicaandA. oxysepala. Genome Biology and Evolution, 2019, 11, 919-930.	1.1	7
11	Capturing additional genetic risk from family history for improved polygenic risk prediction. Communications Biology, 2022, 5, .	2.0	6
12	Simultaneous SNP selection and adjustment for population structure in high dimensional prediction models. PLoS Genetics, 2020, 16, e1008766.	1.5	5
13	Detecting cord blood cell type-specific epigenetic associations with gestational diabetes mellitus and early childhood growth. Clinical Epigenetics, 2021, 13, 131.	1.8	5
14	Block coordinate descent algorithm improves variable selection and estimation in errorâ€inâ€variables regression. Genetic Epidemiology, 2021, 45, 874-890.	0.6	5
15	Identifying Causes of Fracture Beyond Bone Mineral Density: Evidence From Human Genetics. Journal of Bone and Mineral Research, 2020, 37, 1592-1602.	3.1	5
16	Evolutionary Contribution of Duplicated Genes to Genome Evolution in the Ginseng Species Complex. Genome Biology and Evolution, 2021, 13, .	1.1	4
17	Indirect effect inference and application to GAW20 data. BMC Genetics, 2018, 19, 67.	2.7	3
18	Take a Step Down and Beware of H3K36me2: The H3K27m Mutation in Glioma Directs H3K27 Methylation. SSRN Electronic Journal, 0, , .	0.4	0

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#	Article	IF	CITATIONS
19	Title is missing!. , 2020, 16, e1008766.		O
20	Title is missing!. , 2020, 16, e1008766.		0
21	Title is missing!. , 2020, 16, e1008766.		0
22	Title is missing!. , 2020, 16, e1008766.		0
23	Title is missing!. , 2020, 16, e1008766.		0
24	Title is missing!. , 2020, 16, e1008766.		0
25	Genetic and Epigenetic Signatures Associated with the Divergence of Aquilegia Species. Genes, 2022, 13, 793.	1.0	0