## Xingjie Hao

List of Publications by Year in descending order

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XINCUE HAO

#	Article	IF	CITATIONS
1	Association of Public Health Interventions With the Epidemiology of the COVID-19 Outbreak in Wuhan, China. JAMA - Journal of the American Medical Association, 2020, 323, 1915.	7.4	1,333
2	Reconstruction of the full transmission dynamics of COVID-19 in Wuhan. Nature, 2020, 584, 420-424.	27.8	371
3	New Insights on Water Buffalo Genomic Diversity and Post-Domestication Migration Routes From Medium Density SNP Chip Data. Frontiers in Genetics, 2018, 9, 53.	2.3	79
4	Association of <i>MAP4K4 gene</i> single nucleotide polymorphism with mastitis and milk traits in Chinese Holstein cattle. Journal of Dairy Research, 2017, 84, 76-79.	1.4	38
5	Identifying and exploiting trait-relevant tissues with multiple functional annotations in genome-wide association studies. PLoS Genetics, 2018, 14, e1007186.	3.5	30
6	Trans-ethnic genome-wide association study of severe COVID-19. Communications Biology, 2021, 4, 1034.	4.4	29
7	Genome-wide association study identifies candidate genes for piglet splay leg syndrome in different populations. BMC Genetics, 2017, 18, 64.	2.7	22
8	Modelling the initial epidemic trends of COVID-19 in Italy, Spain, Germany, and France. PLoS ONE, 2020, 15, e0241743.	2.5	19
9	Rab11a is required for porcine reproductive and respiratory syndrome virus induced autophagy to promote viral replication. Biochemical and Biophysical Research Communications, 2017, 492, 236-242.	2.1	17
10	Incubation period of coronavirus disease 2019: new implications for intervention and control. International Journal of Environmental Health Research, 2021, , 1-9.	2.7	13
11	Longitudinal relationships of polycyclic aromatic hydrocarbons exposure and genetic susceptibility with blood lipid profiles. Environment International, 2022, 164, 107259.	10.0	13
12	Exploring the Relationship Between Psychiatric Traits and the Risk of Mouth Ulcers Using Bi-Directional Mendelian Randomization. Frontiers in Genetics, 2020, 11, 608630.	2.3	10
13	Causal Graph Among Serum Lipids and Glycemic Traits: A Mendelian Randomization Study. Diabetes, 2022, 71, 1818-1826.	0.6	10
14	Liver and Kidney Function Biomarkers, Blood Cell Traits and Risk of Severe COVID-19: A Mendelian Randomization Study. Frontiers in Genetics, 2021, 12, 647303.	2.3	9
15	Polymorphisms of HOMER1 gene are associated with piglet splay leg syndrome and one significant SNP can affect its intronic promoter activity in vitro. BMC Genetics, 2018, 19, 110.	2.7	7
16	Deconvolution of Bulk Gene Expression Profiles with Single-Cell Transcriptomics to Develop a Cell Type Composition-Based Prognostic Model for Acute Myeloid Leukemia. Frontiers in Cell and Developmental Biology, 2021, 9, 762260.	3.7	5
17	Differences in small noncoding RNAs profile between bull X and Y sperm. PeerJ, 2020, 8, e9822.	2.0	4
18	Integrative analysis of scRNA-seq and GWAS data pinpoints periportal hepatocytes as the relevant liver cell types for blood lipids. Human Molecular Genetics, 2020, 29, 3145-3153.	2.9	2