

Shu Ran

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5367160/publications.pdf>

Version: 2024-02-01

13
papers

135
citations

1478505

6
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

371
citing authors

#	ARTICLE	IF	CITATIONS
1	Total body bone mineral density and severe COVID-19: A Mendelian randomization analysis in five age strata. <i>Bone</i> , 2022, 155, 116281.	2.9	6
2	Three pleiotropic loci associated with bone mineral density and lean body mass. <i>Molecular Genetics and Genomics</i> , 2021, 296, 55-65.	2.1	4
3	The Association Between Coronavirus Disease 2019 Infection and Blood Constituents: A Mendelian Randomization Analysis. <i>Journal of Infectious Diseases</i> , 2021, 224, 922-924.	4.0	1
4	Four pleiotropic loci associated with fat mass and lean mass. <i>International Journal of Obesity</i> , 2020, 44, 2113-2123.	3.4	2
5	Whole-exome sequencing and genome-wide association studies identify novel sarcopenia risk genes in Han Chinese. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1267.	1.2	6
6	Association of 3p27.1 Variants with Whole Body Lean Mass Identified by a Genome-wide Association Study. <i>Scientific Reports</i> , 2020, 10, 4293.	3.3	2
7	Replication of FTO Gene associated with lean mass in a Meta-Analysis of Genome-Wide Association Studies. <i>Scientific Reports</i> , 2020, 10, 5057.	3.3	12
8	Joint Association Analysis Identified 18 New Loci for Bone Mineral Density. <i>Journal of Bone and Mineral Research</i> , 2019, 34, 1086-1094.	2.8	27
9	Identification of a 1p21 independent functional variant for abdominal obesity. <i>International Journal of Obesity</i> , 2019, 43, 2480-2490.	3.4	5
10	Genome-Wide Association Study Identified Copy Number Variants Important for Appendicular Lean Mass. <i>PLoS ONE</i> , 2014, 9, e89776.	2.5	12
11	Bivariate Genome-Wide Association Analyses Identified Genes with Pleiotropic Effects for Femoral Neck Bone Geometry and Age at Menarche. <i>PLoS ONE</i> , 2013, 8, e60362.	2.5	18
12	Genome-wide association study of copy number variation identified gremlin1 as a candidate gene for lean body mass. <i>Journal of Human Genetics</i> , 2012, 57, 33-37.	2.3	30
13	Bivariate genome-wide association study suggests that the DARC gene influences lean body mass and age at menarche. <i>Science China Life Sciences</i> , 2012, 55, 516-520.	4.9	10