Craig A Glastonbury

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/259200/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Meta-analysis of genome-wide association studies for body fat distribution in 694Â649 individuals of European ancestry. Human Molecular Genetics, 2019, 28, 166-174.	2.9	752
2	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. Nature Genetics, 2016, 48, 189-194.	21.4	211
3	Regulatory variants at KLF14 influence type 2 diabetes risk via a female-specific effect on adipocyte size and body composition. Nature Genetics, 2018, 50, 572-580.	21.4	143
4	Smoking induces coordinated DNA methylation and gene expression changes in adipose tissue with consequences for metabolic health. Clinical Epigenetics, 2018, 10, 126.	4.1	110
5	Cell-Type Heterogeneity in Adipose Tissue Is Associated with Complex Traits and Reveals Disease-Relevant Cell-Specific eQTLs. American Journal of Human Genetics, 2019, 104, 1013-1024.	6.2	76
6	Count-ception: Counting by Fully Convolutional Redundant Counting. , 2017, , .		75
7	Integration of human adipocyte chromosomal interactions with adipose gene expression prioritizes obesity-related genes from GWAS. Nature Communications, 2018, 9, 1512.	12.8	75
8	Evaluating the cardiovascular safety of sclerostin inhibition using evidence from meta-analysis of clinical trials and human genetics. Science Translational Medicine, 2020, 12, .	12.4	68
9	Estrogen receptor $\hat{I}\pm$ controls metabolism in white and brown adipocytes by regulating <i>Polg1</i> and mitochondrial remodeling. Science Translational Medicine, 2020, 12, .	12.4	64
10	GWAS Identifies Risk Locus for Erectile Dysfunction and Implicates Hypothalamic Neurobiology and Diabetes in Etiology. American Journal of Human Genetics, 2019, 104, 157-163.	6.2	55
11	Metabolomic profiling to dissect the role of visceral fat in cardiometabolic health. Obesity, 2016, 24, 1380-1388.	3.0	41
12	Adiposity-Dependent Regulatory Effects on Multi-tissue Transcriptomes. American Journal of Human Genetics, 2016, 99, 567-579.	6.2	26
13	Machine Learning based histology phenotyping to investigate the epidemiologic and genetic basis of adipocyte morphology and cardiometabolic traits. PLoS Computational Biology, 2020, 16, e1008044.	3.2	16
14	Further evidence supporting a potential role for ADH1B in obesity. Scientific Reports, 2021, 11, 1932.	3.3	11
15	Fasting and time of day independently modulate circadian rhythm relevant gene expression in adipose and skin tissue. BMC Genomics, 2018, 19, 659.	2.8	9
16	The genetic underpinnings of body fat distribution. Expert Review of Endocrinology and Metabolism, 2017, 12, 417-427.	2.4	3
17	Response to comment on "Evaluating the cardiovascular safety of sclerostin inhibition using evidence from meta-analysis of clinical trials and human genetics― Science Translational Medicine, 2021, 13, eabf4530.	12.4	1
18	Abstract 50: Procollagen C-endopeptidase Enhancer protein 2 (PCPE2) Deficiency Profoundly Affects Adipose Distribution in Mice and Humans and Links HDL Metabolism to Adipocyte Biology. Arteriosclerosis, Thrombosis, and Vascular Biology, 2016, 36, .	2.4	0

#	Article	IF	CITATIONS
19	Title is missing!. , 2020, 16, e1008044.		0
20	Title is missing!. , 2020, 16, e1008044.		0
21	Title is missing!. , 2020, 16, e1008044.		0
22	Title is missing!. , 2020, 16, e1008044.		0
23	Title is missing!. , 2020, 16, e1008044.		0
24	Title is missing!. , 2020, 16, e1008044.		0