Bin Wu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2585665/publications.pdf

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		1307594	1125743	
18	176	7	13	
papers	citations	h-index	g-index	
19	19	19	260	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Birth weight was associated positively with gluteofemoral fat mass and inversely with 2-h postglucose insulin concentrations, a marker of insulin resistance, in young normal-weight Japanese women. Diabetology International, 2022, 13, 375-380.	1.4	3
2	Weight Trajectory Since Birth, Current Body Composition, Dietary Intake, and Glucose Tolerance in Young Underweight Japanese Women. Women S Health Reports, 2022, 3, 215-221.	0.8	1
3	Higher circulating orosomucoid and lower early-phase insulin secretion in midlife Japanese with slower glucose disposal during oral glucose tolerance tests. Diabetology International, 2020, 11, 27-32.	1.4	3
4	Higher circulating orosomucoid, an acute-phase protein, and reduced glucose-induced insulin secretion in middle-aged Japanese people with prediabetes. BMJ Open Diabetes Research and Care, 2020, 8, e001392.	2.8	3
5	Muscle fatigue detection and treatment system driven by internet of things. BMC Medical Informatics and Decision Making, 2019, 19, 275.	3.0	7
6	An integrative approach to investigate the association among high-sensitive C-reactive protein, body fat mass distribution, and other cardiometabolic risk factors in young healthy women. Methods, 2018, 145, 60-66.	3.8	4
7	Different Associations of Trunk and Lower-Body Fat Mass Distribution with Cardiometabolic Risk Factors between Healthy Middle-Aged Men and Women. International Journal of Endocrinology, 2018, 2018, 1-10.	1.5	12
8	Involvement of MicroRNAs in Diabetes and Its Complications. Methods in Molecular Biology, 2017, 1617, 225-239.	0.9	18
9	MicroRNA Regulatory Networks as Biomarkers in Obesity: The Emerging Role. Methods in Molecular Biology, 2017, 1617, 241-260.	0.9	7
10	Co-expression analysis among microRNAs, long non-coding RNAs, and messenger RNAs to understand the pathogenesis and progression of diabetic kidney disease at the genetic level. Methods, 2017, 124, 46-56.	3.8	11
11	Combine biological experiments, statistical analysis, and semantic search to discover association among high-sensitive C-reactive protein, body fat mass distribution, and other cardiometabolic risk factors in young healthy women., 2017,,.		O
12	Innovative microRNA-lncRNA-mRNA co-expression analysis to understand the pathogenesis and progression of diabetic kidney disease. , 2016 , , .		O
13	The Non-Coding RNA Ontology (NCRO): a comprehensive resource for the unification of non-coding RNA biology. Journal of Biomedical Semantics, 2016, 7, 24.	1.6	10
14	The development of non-coding RNA ontology. International Journal of Data Mining and Bioinformatics, 2016, 15, 214.	0.1	9
15	OmniSearch: a semantic search system based on the Ontology for MIcroRNA Target (OMIT) for microRNA-target gene interaction data. Journal of Biomedical Semantics, 2016, 7, 25.	1.6	27
16	A domain ontology for the Non-Coding RNA field. , 2015, , .		0
17	Relationships of Systemic Oxidative Stress to Body Fat Distribution, Adipokines and Inflammatory Markers in Healthy Middle-aged Women. Endocrine Journal, 2009, 56, 773-782.	1.6	50
18	Effect of acupuncture on immunomodulation in patients with malignant tumors., 1996, 2, 266-269.		7