Qingying Meng

List of Publications by Year in descending order

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22 papers	1,389 citations	471509 17 h-index	713466 21 g-index
23 all docs	23 docs citations	23 times ranked	3557 citing authors

#	Article	IF	Citations
1	Multiâ€Tissue Multiâ€Omics Nutrigenomics Indicates Contextâ€Specific Effects of Docosahexaenoic Acid on Rat Brain. Molecular Nutrition and Food Research, 2020, 64, e2000788.	3.3	2
2	Prenatal Bisphenol A Exposure in Mice Induces Multitissue Multiomics Disruptions Linking to Cardiometabolic Disorders. Endocrinology, 2019, 160, 409-429.	2.8	35
3	Maternal High-Protein and Low-Protein Diets Perturb Hypothalamus and Liver Transcriptome and Metabolic Homeostasis in Adult Mouse Offspring. Frontiers in Genetics, 2018, 9, 642.	2.3	6
4	Biglycan gene connects metabolic dysfunction with brain disorder. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2018, 1864, 3679-3687.	3.8	18
5	Traumatic Brain Injury Induces Genome-Wide Transcriptomic, Methylomic, and Network Perturbations in Brain and Blood Predicting Neurological Disorders. EBioMedicine, 2017, 16, 184-194.	6.1	88
6	Systems Nutrigenomics Reveals Brain Gene Networks Linking Metabolic and Brain Disorders. EBioMedicine, 2016, 7, 157-166.	6.1	59
7	Network-Based Identification and Prioritization of Key Regulators of Coronary Artery Disease Loci. Arteriosclerosis, Thrombosis, and Vascular Biology, 2016, 36, 928-941.	2.4	66
8	Abstract 58: Network-based Identification and Prioritization of Key Regulators of Coronary Artery Disease Loci. Arteriosclerosis, Thrombosis, and Vascular Biology, 2016, 36, .	2.4	0
9	Dissecting the Roles of MicroRNAs in Coronary Heart Disease via Integrative Genomic Analyses. Arteriosclerosis, Thrombosis, and Vascular Biology, 2015, 35, 1011-1021.	2.4	53
10	A systems genetics study of swine illustrates mechanisms underlying human phenotypic traits. BMC Genomics, 2015, 16, 88.	2.8	28
11	Integrative network analysis reveals molecular mechanisms of blood pressure regulation. Molecular Systems Biology, 2015, 11, 799.	7.2	102
12	Integrative Genomics Reveals Novel Molecular Pathways and Gene Networks for Coronary Artery Disease. PLoS Genetics, 2014, 10, e1004502.	3.5	192
13	Shared Molecular Pathways and Gene Networks for Cardiovascular Disease and Type 2 Diabetes Mellitus in Women Across Diverse Ethnicities. Circulation: Cardiovascular Genetics, 2014, 7, 911-919.	5.1	48
14	Systems Biology Approaches and Applications in Obesity, Diabetes, and Cardiovascular Diseases. Current Cardiovascular Risk Reports, 2013, 7, 73-83.	2.0	49
15	A Systems Biology Framework Identifies Molecular Underpinnings of Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 1427-1434.	2.4	157
16	Genetic Regulation of Life Span, Metabolism, and Body Weight in Pohn, a New Wild-Derived Mouse Strain. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2013, 68, 27-35.	3.6	15
17	Genetic coregulation of age of female sexual maturation and lifespan through circulating IGF1 among inbred mouse strains. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 8224-8229.	7.1	98
18	Systems analysis of eleven rodent disease models reveals an inflammatome signature and key drivers. Molecular Systems Biology, 2012, 8, 594.	7.2	134

#	Article	IF	Citations
19	Identification of genetic determinants of IGFâ€1 levels and longevity among mouse inbred strains. Aging Cell, 2010, 9, 823-836.	6.7	32
20	Effects of epigallocatechin-3-gallate on mitochondrial integrity and antioxidative enzyme activity in the aging process of human fibroblast. Free Radical Biology and Medicine, 2008, 44, 1032-1041.	2.9	79
21	Regulating the Age-Related Oxidative Damage, Mitochondrial Integrity, and Antioxidative Enzyme Activity in Fischer 344 Rats by Supplementation of the Antioxidant Epigallocatechin-3-Gallate. Rejuvenation Research, 2008, 11, 649-660.	1.8	48
22	Age-related changes in mitochondrial function and antioxidative enzyme activity in fischer 344 rats. Mechanisms of Ageing and Development, 2007, 128, 286-292.	4.6	80