

Heather A Lawson

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

962
citations

623734

14
h-index

642732

23
g-index

34
all docs

34
docs citations

34
times ranked

1395
citing authors

#	ARTICLE	IF	CITATIONS
1	Parent-of-origin effects propagate through networks to shape metabolic traits. <i>ELife</i> , 2022, 11, .	6.0	6
2	The Human Pangenome Project: a global resource to map genomic diversity. <i>Nature</i> , 2022, 604, 437-446.	27.8	192
3	Genetic, epigenetic, and environmental mechanisms govern allele-specific gene expression. <i>Genome Research</i> , 2022, 32, 1042-1057.	5.5	6
4	Epigenomic analysis reveals prevalent contribution of transposable elements to <i>cis</i> -regulatory elements, tissue-specific expression, and alternative promoters in zebrafish. <i>Genome Research</i> , 2022, 32, 1424-1436.	5.5	7
5	Pancreatic β -cell heterogeneity in health and diabetes: classes, sources, and subtypes. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2021, 320, E716-E731.	3.5	21
6	Brown Adipose Expansion and Remission of Glycemic Dysfunction in Obese SM/J Mice. <i>Cell Reports</i> , 2020, 33, 108237.	6.4	9
7	Spontaneous restoration of functional β -cell mass in obese SM/J mice. <i>Physiological Reports</i> , 2020, 8, e14573.	1.7	5
8	Genetic background and diet affect brown adipose gene coexpression networks associated with metabolic phenotypes. <i>Physiological Genomics</i> , 2020, 52, 223-233.	2.3	4
9	Dietary iron interacts with genetic background to influence glucose homeostasis. <i>Nutrition and Metabolism</i> , 2019, 16, 13.	3.0	7
10	The NIEHS TaRGET II Consortium and environmental epigenomics. <i>Nature Biotechnology</i> , 2018, 36, 225-227.	17.5	79
11	Epigenetics of metabolic syndrome. <i>Physiological Genomics</i> , 2018, 50, 947-955.	2.3	36
12	Ironing out the Details: Untangling Dietary Iron and Genetic Background in Diabetes. <i>Nutrients</i> , 2018, 10, 1437.	4.1	15
13	Animal Models of Metabolic Syndrome. , 2017, , 221-243.		8
14	Reduced efficiency of sarcolipin-dependent respiration in myocytes from humans with severe obesity. <i>Obesity</i> , 2015, 23, 1440-1449.	3.0	41
15	Using whole-genome sequences of the LG/J and SM/J inbred mouse strains to prioritize quantitative trait genes and nucleotides. <i>BMC Genomics</i> , 2015, 16, 415.	2.8	31
16	Genomic imprinting and parent-of-origin effects on complex traits. <i>Nature Reviews Genetics</i> , 2013, 14, 609-617.	16.3	219
17	Animal Models of Metabolic Syndrome. , 2013, , 243-264.		4
18	Imputation of Single-Nucleotide Polymorphisms in Inbred Mice Using Local Phylogeny. <i>Genetics</i> , 2012, 190, 449-458.	2.9	42

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19	Obesityâ€“insulin targeted genes in the 3p26-25 region in human studies and LG/J and SM/J mice. <i>Metabolism: Clinical and Experimental</i> , 2012, 61, 1129-1141.	3.4	9
20	Dietâ€“Dependent Genetic and Genomic Imprinting Effects on Obesity in Mice. <i>Obesity</i> , 2011, 19, 160-170.	3.0	49
21	Genetic factors and diet affect long-bone length in the F34 LG,SM advanced intercross. <i>Mammalian Genome</i> , 2011, 22, 178-196.	2.2	25
22	The importance of context to the genetic architecture of diabetes-related traits is revealed in a genome-wide scan of a LG/JÃ—SM/J murine model. <i>Mammalian Genome</i> , 2011, 22, 197-208.	2.2	38
23	Genetic Effects at Pleiotropic Loci Are Context-Dependent with Consequences for the Maintenance of Genetic Variation in Populations. <i>PLoS Genetics</i> , 2011, 7, e1002256.	3.5	47
24	Metabolic Syndrome Components in Murine Models. <i>Endocrine, Metabolic and Immune Disorders - Drug Targets</i> , 2010, 10, 25-40.	1.2	23
25	Genetic, epigenetic, and gene-by-diet interaction effects underlie variation in serum lipids in a LG/JÃ—SM/J murine model. <i>Journal of Lipid Research</i> , 2010, 51, 2976-2984.	4.2	32