

# 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	Genomic imprinting and parent-of-origin effects on complex traits. <i>Nature Reviews Genetics</i> , 2013, 14, 609-617.	16.3	219
2	The Human Pangenome Project: a global resource to map genomic diversity. <i>Nature</i> , 2022, 604, 437-446.	27.8	192
3	The NIEHS TaRGET II Consortium and environmental epigenomics. <i>Nature Biotechnology</i> , 2018, 36, 225-227.	17.5	79
4	Diet-Dependent Genetic and Genomic Imprinting Effects on Obesity in Mice. <i>Obesity</i> , 2011, 19, 160-170.	3.0	49
5	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
6	Imputation of Single-Nucleotide Polymorphisms in Inbred Mice Using Local Phylogeny. <i>Genetics</i> , 2012, 190, 449-458.	2.9	42
7	Reduced efficiency of sarcolipin-dependent respiration in myocytes from humans with severe obesity. <i>Obesity</i> , 2015, 23, 1440-1449.	3.0	41
8	The importance of context to the genetic architecture of diabetes-related traits is revealed in a genome-wide scan of a LG/J-ASM/J murine model. <i>Mammalian Genome</i> , 2011, 22, 197-208.	2.2	38
9	Epigenetics of metabolic syndrome. <i>Physiological Genomics</i> , 2018, 50, 947-955.	2.3	36
10	Genetic, epigenetic, and gene-by-diet interaction effects underlie variation in serum lipids in a LG/J-ASM/J murine model. <i>Journal of Lipid Research</i> , 2010, 51, 2976-2984.	4.2	32
11	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
12	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
13	Metabolic Syndrome Components in Murine Models. <i>Endocrine, Metabolic and Immune Disorders - Drug Targets</i> , 2010, 10, 25-40.	1.2	23
14	Pancreatic $\beta$ -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
15	Ironing out the Details: Untangling Dietary Iron and Genetic Background in Diabetes. <i>Nutrients</i> , 2018, 10, 1437.	4.1	15
16	Obesity-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
17	Brown Adipose Expansion and Remission of Glycemic Dysfunction in Obese SM/J Mice. <i>Cell Reports</i> , 2020, 33, 108237.	6.4	9
18	Animal Models of Metabolic Syndrome. , 2017, , 221-243.		8

#	ARTICLE	IF	CITATIONS
19	Dietary iron interacts with genetic background to influence glucose homeostasis. Nutrition and Metabolism, 2019, 16, 13.	3.0	7
20	Epigenomic analysis reveals prevalent contribution of transposable elements to <i>cis</i> -regulatory elements, tissue-specific expression, and alternative promoters in zebrafish. Genome Research, 2022, 32, 1424-1436.	5.5	7
21	Parent-of-origin effects propagate through networks to shape metabolic traits. ELife, 2022, 11, .	6.0	6
22	Genetic, epigenetic, and environmental mechanisms govern allele-specific gene expression. Genome Research, 2022, 32, 1042-1057.	5.5	6
23	Spontaneous restoration of functional $\beta$ -cell mass in obese SM/J mice. Physiological Reports, 2020, 8, e14573.	1.7	5
24	Animal Models of Metabolic Syndrome. , 2013, , 243-264.		4
25	Genetic background and diet affect brown adipose gene coexpression networks associated with metabolic phenotypes. Physiological Genomics, 2020, 52, 223-233.	2.3	4