

Mary E Haas

List of Publications by Citations

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Version: 2024-04-27

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

15
papers

557
citations

10
h-index

16
g-index

16
ext. papers

773
ext. citations

13.6
avg, IF

3.25
L-index

#	Paper	IF	Citations
15	Hepatic insulin signaling is required for obesity-dependent expression of SREBP-1c mRNA but not for feeding-dependent expression. <i>Cell Metabolism</i> , 2012 , 15, 873-84	24.6	141
14	The regulation of ApoB metabolism by insulin. <i>Trends in Endocrinology and Metabolism</i> , 2013 , 24, 391-7	8.8	76
13	Role of Insulin in the Regulation of Proprotein Convertase Subtilisin/Kexin Type 9. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015 , 35, 1589-96	9.4	63
12	Genetic Association of Albuminuria with Cardiometabolic Disease and Blood Pressure. <i>American Journal of Human Genetics</i> , 2018 , 103, 461-473	11	62
11	The Role of Proprotein Convertase Subtilisin/Kexin Type 9 in Nephrotic Syndrome-Associated Hypercholesterolemia. <i>Circulation</i> , 2016 , 134, 61-72	16.7	56
10	Phenotypic Refinement of Heart Failure in a National Biobank Facilitates Genetic Discovery. <i>Circulation</i> , 2018 ,	16.7	51
9	A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease. <i>PLoS Genetics</i> , 2020 , 16, e1008629	6	49
8	Insulin Dissociates the Effects of Liver X Receptor on Lipogenesis, Endoplasmic Reticulum Stress, and Inflammation. <i>Journal of Biological Chemistry</i> , 2016 , 291, 1115-22	5.4	14
7	Effect of Leptin Replacement on PCSK9 in ob/ob Mice and Female Lipodystrophic Patients. <i>Endocrinology</i> , 2016 , 157, 1421-9	4.8	14
6	Association of Habitual Alcohol Intake With Risk of Cardiovascular Disease.. <i>JAMA Network Open</i> , 2022 , 5, e223849	10.4	10
5	Machine learning enables new insights into clinical significance of and genetic contributions to liver fat accumulation		5
4	A trans-ancestry genome-wide association study of unexplained chronic ALT elevation as a proxy for nonalcoholic fatty liver disease with histological and radiological validation		5
3	Electronic health record-based genome-wide meta-analysis provides insights on the genetic architecture of non-alcoholic fatty liver disease. <i>Cell Reports Medicine</i> , 2021 , 2, 100437	18	4
2	Machine learning enables new insights into genetic contributions to liver fat accumulation.. <i>Cell Genomics</i> , 2021 , 1,		3
1	A multiancestry genome-wide association study of unexplained chronic ALT elevation as a proxy for nonalcoholic fatty liver disease with histological and radiological validation. <i>Nature Genetics</i> ,	36.3	2