

Kikuko Hotta

List of Publications by Citations

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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

4,282
citations

12
h-index

16
g-index

16
ext. papers

4,561
ext. citations

4.9
avg, IF

4.29
L-index

#	Paper	IF	Citations
15	Hypoadiponectinemia in obesity and type 2 diabetes: close association with insulin resistance and hyperinsulinemia. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2001 , 86, 1930-5	5.6	2703
14	Adipocyte-derived plasma protein, adiponectin, suppresses lipid accumulation and class A scavenger receptor expression in human monocyte-derived macrophages. <i>Circulation</i> , 2001 , 103, 1057-63	16.7	1066
13	Genome-wide scan revealed that polymorphisms in the PNPLA3, SAMM50, and PARVB genes are associated with development and progression of nonalcoholic fatty liver disease in Japan. <i>Human Genetics</i> , 2013 , 132, 783-92	6.3	138
12	Association of the rs738409 polymorphism in PNPLA3 with liver damage and the development of nonalcoholic fatty liver disease. <i>BMC Medical Genetics</i> , 2010 , 11, 172	2.1	110
11	Targeted-bisulfite sequence analysis of the methylation of CpG islands in genes encoding PNPLA3, SAMM50, and PARVB of patients with non-alcoholic fatty liver disease. <i>Journal of Hepatology</i> , 2015 , 63, 494-502	13.4	53
10	Association of polymorphisms in GCKR and TRIB1 with nonalcoholic fatty liver disease and metabolic syndrome traits. <i>Endocrine Journal</i> , 2014 , 61, 683-9	2.9	41
9	Characteristics of non-obese non-alcoholic fatty liver disease: Effect of genetic and environmental factors. <i>Hepatology Research</i> , 2016 , 46, 1011-8	5.1	39
8	The expression of SPARC in adipose tissue and its increased plasma concentration in patients with coronary artery disease. <i>Obesity</i> , 2001 , 9, 388-93		37
7	Identification of the genomic region under epigenetic regulation during non-alcoholic fatty liver disease progression. <i>Hepatology Research</i> , 2018 , 48, E320-E334	5.1	23
6	Gene expression profiling of non-alcoholic steatohepatitis using gene set enrichment analysis. <i>Hepatology Research</i> , 2008 , 38, 1204-12	5.1	22
5	Identification of differentially methylated region (DMR) networks associated with progression of nonalcoholic fatty liver disease. <i>Scientific Reports</i> , 2018 , 8, 13567	4.9	21
4	Targeted next-generation sequencing and fine linkage disequilibrium mapping reveals association of PNPLA3 and PARVB with the severity of nonalcoholic fatty liver disease. <i>Journal of Human Genetics</i> , 2014 , 59, 241-6	4.3	14
3	Identification of core gene networks and hub genes associated with progression of non-alcoholic fatty liver disease by RNA sequencing. <i>Hepatology Research</i> , 2017 , 47, 1445-1458	5.1	10
2	A novel compound heterozygous mutation in identified in a Japanese patient. <i>Human Genome Variation</i> , 2019 , 6, 14	1.8	5
1	Cell Biology of Visceral Fat 1999 , 48, 963-970,1194		