

Dorota Kaminska

List of Publications by Year in Descending Order

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Version: 2024-04-20

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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.

35
papers

1,535
citations

17
h-index

37
g-index

37
ext. papers

1,829
ext. citations

7.5
avg, IF

3.76
L-index

#	Paper	IF	Citations
35	Identification of 90 NAFLD GWAS loci and establishment of NAFLD PRS and causal role of NAFLD in coronary artery disease.. <i>Human Genetics and Genomics Advances</i> , 2022 , 3, 100056	0.8	0
34	Indole-3-Propionic Acid, a Gut-Derived Tryptophan Metabolite, Associates with Hepatic Fibrosis. <i>Nutrients</i> , 2021 , 13,	6.7	6
33	Integrative analysis of liver-specific non-coding regulatory SNPs associated with the risk of coronary artery disease. <i>American Journal of Human Genetics</i> , 2021 , 108, 411-430	11	4
32	Serum aromatic and branched-chain amino acids associated with NASH demonstrate divergent associations with serum lipids. <i>Liver International</i> , 2021 , 41, 754-763	7.9	8
31	Protein Phosphatase 1 Regulatory Subunit 3B Genotype at rs4240624 Has a Major Effect on Gallbladder Bile Composition. <i>Hepatology Communications</i> , 2021 , 5, 244-257	6	1
30	Differential Mitochondrial Gene Expression in Adipose Tissue Following Weight Loss Induced by Diet or Bariatric Surgery. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, 1312-1324	5.6	1
29	Identification of TBX15 as an adipose master trans regulator of abdominal obesity genes. <i>Genome Medicine</i> , 2021 , 13, 123	14.4	2
28	Parental metabolic syndrome epigenetically reprograms offspring hepatic lipid metabolism in mice. <i>Journal of Clinical Investigation</i> , 2020 , 130, 2391-2407	15.9	14
27	Total liver phosphatidylcholine content associates with non-alcoholic steatohepatitis and glycine N-methyltransferase expression. <i>Liver International</i> , 2019 , 39, 1895-1905	7.9	3
26	Novel Lipid Long Intervening Noncoding RNA, Oligodendrocyte Maturation-Associated Long Intergenic Noncoding RNA, Regulates the Liver Steatosis Gene Stearoyl-Coenzyme A Desaturase As an Enhancer RNA. <i>Hepatology Communications</i> , 2019 , 3, 1356-1372	6	17
25	Serum, liver and bile sitosterol and sitostanol in obese patients with and without NAFLD. <i>Bioscience Reports</i> , 2018 , 38,	4.1	4
24	Identification and characterization of a FOXA2-regulated transcriptional enhancer at a type 2 diabetes intronic locus that controls GCKR expression in liver cells. <i>Genome Medicine</i> , 2017 , 9, 63	14.4	16
23	Regulation of alternative splicing in human obesity loci. <i>Obesity</i> , 2016 , 24, 2033-7	8	9
22	Transmembrane 6 superfamily member 2 gene variant disentangles nonalcoholic steatohepatitis from cardiovascular disease. <i>Hepatology</i> , 2015 , 61, 506-14	11.2	311
21	The 148 M allele of the PNPLA3 is associated with plasma irisin levels in a population sample of Caucasian children: The PANIC Study. <i>Metabolism: Clinical and Experimental</i> , 2015 , 64, 793-6	12.7	14
20	Ketone body production is differentially altered in steatosis and non-alcoholic steatohepatitis in obese humans. <i>Liver International</i> , 2015 , 35, 1853-61	7.9	48
19	Dietary polyunsaturated fatty acids and the Pro12Ala polymorphisms of PPARG regulate serum lipids through divergent pathways: a randomized crossover clinical trial. <i>Genes and Nutrition</i> , 2015 , 10, 43	4.3	11

18	Statin use and non-alcoholic steatohepatitis in at risk individuals. <i>Journal of Hepatology</i> , 2015 , 63, 705-12	3.4	227
17	Specific collagen XVIII isoforms promote adipose tissue accrual via mechanisms determining adipocyte number and affect fat deposition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, E3043-52	11.5	31
16	High-fat diet increases tau expression in the brain of T2DM and AD mice independently of peripheral metabolic status. <i>Journal of Nutritional Biochemistry</i> , 2014 , 25, 634-41	6.3	40
15	Lipoprotein subclass metabolism in nonalcoholic steatohepatitis. <i>Journal of Lipid Research</i> , 2014 , 55, 2676-84	6.3	51
14	Adipose tissue INSR splicing in humans associates with fasting insulin level and is regulated by weight loss. <i>Diabetologia</i> , 2014 , 57, 347-51	10.3	30
13	Desmosterol in human nonalcoholic steatohepatitis. <i>Hepatology</i> , 2013 , 58, 976-82	11.2	34
12	Downregulation of CPPED1 expression improves glucose metabolism in vitro in adipocytes. <i>Diabetes</i> , 2013 , 62, 3747-50	0.9	18
11	Regulation of alternative splicing in obesity and weight loss. <i>Adipocyte</i> , 2013 , 2, 143-7	3.2	12
10	Adipose tissue TCF7L2 splicing is regulated by weight loss and associates with glucose and fatty acid metabolism. <i>Diabetes</i> , 2012 , 61, 2807-13	0.9	57
9	Serum interleukin 1 receptor antagonist as an independent marker of non-alcoholic steatohepatitis in humans. <i>Journal of Hepatology</i> , 2012 , 56, 663-70	13.4	69
8	Response to Brosch et al. <i>Cell Metabolism</i> , 2012 , 15, 267-269	24.6	4
7	Conjugated bile acids associate with altered rates of glucose and lipid oxidation after Roux-en-Y gastric bypass. <i>Obesity Surgery</i> , 2012 , 22, 1473-80	3.7	127
6	Local peripheral opioid effects and expression of opioid genes in the spinal cord and dorsal root ganglia in neuropathic and inflammatory pain. <i>Pain</i> , 2009 , 141, 283-291	8	150
5	alpha-Synuclein expression in the brain and blood during abstinence from chronic alcohol drinking in mice. <i>Neuropharmacology</i> , 2008 , 54, 1239-46	5.5	14
4	Interleukin-1 alpha has antiallodynic and antihyperalgesic activities in a rat neuropathic pain model. <i>Pain</i> , 2008 , 138, 587-597	8	83
3	Regulation of ERK1/2 phosphorylation by acute and chronic morphine - implications for the role of cAMP-responsive element binding factor (CREB)-dependent and Ets-like protein-1 (Elk-1)-dependent transcription; small interfering RNA-based strategy. <i>FEBS Journal</i> , 2008 , 275, 3836-49	5.7	15
2	Morphine effects on striatal transcriptome in mice. <i>Genome Biology</i> , 2007 , 8, R128	18.3	60
1	Gene expression profiling in the striatum of inbred mouse strains with distinct opioid-related phenotypes. <i>BMC Genomics</i> , 2006 , 7, 146	4.5	43

