

# Dorota Kaminska

## List of Publications by Year in descending order

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Version: 2024-02-01

36  
papers

2,075  
citations

361045

20  
h-index

344852

36  
g-index

37  
all docs

37  
docs citations

37  
times ranked

4095  
citing authors

#	ARTICLE	IF	CITATIONS
1	Transmembrane 6 superfamily member 2 gene variant disentangles nonalcoholic steatohepatitis from cardiovascular disease. <i>Hepatology</i> , 2015, 61, 506-514.	3.6	424
2	Statin use and non-alcoholic steatohepatitis in at risk individuals. <i>Journal of Hepatology</i> , 2015, 63, 705-712.	1.8	309
3	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.	2.0	183
4	Conjugated Bile Acids Associate with Altered Rates of Glucose and Lipid Oxidation after Roux-en-Y Gastric Bypass. <i>Obesity Surgery</i> , 2012, 22, 1473-1480.	1.1	135
5	Interleukin-1alpha has antiallodynic and antihyperalgesic activities in a rat neuropathic pain model. <i>Pain</i> , 2008, 138, 587-597.	2.0	88
6	Serum interleukin 1 receptor antagonist as an independent marker of non-alcoholic steatohepatitis in humans. <i>Journal of Hepatology</i> , 2012, 56, 663-670.	1.8	87
7	Morphine effects on striatal transcriptome in mice. <i>Genome Biology</i> , 2007, 8, R128.	13.9	74
8	Adipose Tissue TCF7L2 Splicing Is Regulated by Weight Loss and Associates With Glucose and Fatty Acid Metabolism. <i>Diabetes</i> , 2012, 61, 2807-2813.	0.3	67
9	Ketone body production is differentially altered in steatosis and nonalcoholic steatohepatitis in obese humans. <i>Liver International</i> , 2015, 35, 1853-1861.	1.9	62
10	Lipoprotein subclass metabolism in nonalcoholic steatohepatitis. <i>Journal of Lipid Research</i> , 2014, 55, 2676-2684.	2.0	59
11	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, 634-641.	1.9	50
12	Gene expression profiling in the striatum of inbred mouse strains with distinct opioid-related phenotypes. <i>BMC Genomics</i> , 2006, 7, 146.	1.2	48
13	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.	3.3	43
14	Desmosterol in human nonalcoholic steatohepatitis. <i>Hepatology</i> , 2013, 58, 976-982.	3.6	42
15	Parental metabolic syndrome epigenetically reprograms offspring hepatic lipid metabolism in mice. <i>Journal of Clinical Investigation</i> , 2020, 130, 2391-2407.	3.9	42
16	Adipose tissue INSR splicing in humans associates with fasting insulin level and is regulated by weight loss. <i>Diabetologia</i> , 2014, 57, 347-351.	2.9	41
17	Downregulation of <i>CPPED1</i> Expression Improves Glucose Metabolism In Vitro in Adipocytes. <i>Diabetes</i> , 2013, 62, 3747-3750.	0.3	32
18	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.	2.0	28

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19	Indole-3-Propionic Acid, a Gut-Derived Tryptophan Metabolite, Associates with Hepatic Fibrosis. <i>Nutrients</i> , 2021, 13, 3509.	1.7	25
20	Serum aromatic and branched-chain amino acids associated with NASH demonstrate divergent associations with serum lipids. <i>Liver International</i> , 2021, 41, 754-763.	1.9	23
21	Identification of TBX15 as an adipose master trans regulator of abdominal obesity genes. <i>Genome Medicine</i> , 2021, 13, 123.	3.6	23
22	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.	3.6	21
23	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.	2.6	20
24	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, 793-796.	1.5	19
25	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 (Elk)-dependent transcription; small interfering RNA-based strategy. <i>FEBS Journal</i> , 2008, 275, 3836-3849.	2.2	15
26	Dietary polyunsaturated fatty acids and the Pro12Ala polymorphisms of PPARC regulate serum lipids through divergent pathways: a randomized crossover clinical trial. <i>Genes and Nutrition</i> , 2015, 10, 43.	1.2	15
27	Î±-Synuclein expression in the brain and blood during abstinence from chronic alcohol drinking in mice. <i>Neuropharmacology</i> , 2008, 54, 1239-1246.	2.0	14
28	Regulation of alternative splicing in obesity and weight loss. <i>Adipocyte</i> , 2013, 2, 143-147.	1.3	13
29	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.	1.8	13
30	Total liver phosphatidylcholine content associates with non-alcoholic steatohepatitis and glycine N-methyltransferase expression. <i>Liver International</i> , 2019, 39, 1895-1905.	1.9	12
31	Regulation of alternative splicing in human obesity loci. <i>Obesity</i> , 2016, 24, 2033-2037.	1.5	11
32	Hyperinsulinemia Is Highly Associated With Markers of Hepatocytic Senescence in Two Independent Cohorts. <i>Diabetes</i> , 2022, 71, 1929-1936.	0.3	11
33	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.	1.0	10
34	Serum, liver and bile sitosterol and sitostanol in obese patients with and without NAFLD. <i>Bioscience Reports</i> , 2018, 38, .	1.1	6
35	Response to Brosch et al.. <i>Cell Metabolism</i> , 2012, 15, 267-269.	7.2	5
36	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.	2.0	4