Maria Kulecka

List of Publications by Year in descending order

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516561 610775 45 745 16 24 citations h-index g-index papers 46 46 46 1501 all docs docs citations times ranked citing authors

#	Article	IF	Citations
1	Limited prolonged effects of rifaximin treatment on irritable bowel syndrome-related differences in the fecal microbiome and metabolome. Gut Microbes, 2016, 7, 397-413.	4.3	68
2	Prolonged transfer of feces from the lean mice modulates gut microbiota in obese mice. Nutrition and Metabolism, 2016, 13, 57.	1.3	55
3	IDH1/2 Mutations Predict Shorter Survival in Chondrosarcoma. Journal of Cancer, 2018, 9, 998-1005.	1.2	50
4	The composition and richness of the gut microbiota differentiate the top Polish endurance athletes from sedentary controls. Gut Microbes, 2020, 11, 1374-1384.	4.3	48
5	Wholeâ€exome sequencing identifies novel pathogenic variants across the ⟨i>ATP7B ⟨li>gene and some modifiers of Wilson's disease phenotype. Liver International, 2019, 39, 177-186.	1.9	38
6	New recurrent BRCA1/2 mutations in Polish patients with familial breast/ovarian cancer detected by next generation sequencing. BMC Medical Genomics, 2015, 8, 19.	0.7	34
7	Genetic architecture differences between pediatric and adult-onset inflammatory bowel diseases in the Polish population. Scientific Reports, 2016, 6, 39831.	1.6	33
8	Effect of Saccharomyces boulardii and Mode of Delivery on the Early Development of the Gut Microbial Community in Preterm Infants. PLoS ONE, 2016, 11, e0150306.	1.1	29
9	Candidate diagnostic miRNAs that can detect cancer in prostate biopsy. Prostate, 2018, 78, 178-185.	1.2	24
10	Genome-wide co-localization of active EGFR and downstream ERK pathway kinases mirrors mitogen-inducible RNA polymerase 2 genomic occupancy. Nucleic Acids Research, 2016, 44, gkw763.	6.5	23
11	A preliminary evaluation of next-generation sequencing as a screening tool for targeted genotyping of erythrocyte and platelet antigens in blood donors. Blood Transfusion, 2018, 16, 285-292.	0.3	23
12	RNA sequencing reveals widespread transcriptome changes in a renal carcinoma cell line. Oncotarget, 2018, 9, 8597-8613.	0.8	22
13	Exome scale map of genetic alterations promoting metastasis in colorectal cancer. BMC Genetics, 2018, 19, 85.	2.7	22
14	Redefining the Practical Utility of Blood Transcriptome Biomarkers in Inflammatory Bowel Diseases. Journal of Crohn's and Colitis, 2019, 13, 626-633.	0.6	22
15	Lack of miR-378 attenuates muscular dystrophy in mdx mice. JCI Insight, 2020, 5, .	2.3	22
16	A heterozygous mutation in GOT1 is associated with familial macro-aspartate aminotransferase. Journal of Hepatology, 2017, 67, 1026-1030.	1.8	18
17	Common functional alterations identified in blood transcriptome of autoimmune cholestatic liver and inflammatory bowel diseases. Scientific Reports, 2019, 9, 7190.	1.6	18
18	Ectopic overexpression of MCPIP1 impairs adipogenesis by modulating microRNAs. Biochimica Et Biophysica Acta - Molecular Cell Research, 2018, 1865, 186-195.	1.9	17

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19	Mitochondrial amidoxime-reducing component 2 (MARC2) has a significant role in N-reductive activity and energy metabolism. Journal of Biological Chemistry, 2019, 294, 17593-17602.	1.6	15
20	Keratinocyte-specific ablation of Mcpip1 impairs skin integrity and promotes local and systemic inflammation. Journal of Molecular Medicine, 2019, 97, 1669-1684.	1.7	14
21	Multi-Organ Transcriptome Dynamics in a Mouse Model of Cecal Ligation and Puncture-Induced Polymicrobial Sepsis. Journal of Inflammation Research, 2021, Volume 14, 2377-2388.	1.6	14
22	Integrative genomics reveal a role for MCPIP1 in adipogenesis and adipocyte metabolism. Cellular and Molecular Life Sciences, 2020, 77, 4899-4919.	2.4	13
23	In search for interplay between stool microRNAs, microbiota and short chain fatty acids in Crohn's disease - a preliminary study. BMC Gastroenterology, 2020, 20, 307.	0.8	12
24	Signatures of circulating microRNA in four sarcoma subtypes. Journal of Cancer, 2020, 11, 874-882.	1.2	12
25	Differences between Well-Differentiated Neuroendocrine Tumors and Ductal Adenocarcinomas of the Pancreas Assessed by Multi-Omics Profiling. International Journal of Molecular Sciences, 2020, 21, 4470.	1.8	10
26	Challenges in Cancer Biomarker Discovery Exemplified by the Identification of Diagnostic MicroRNAs in Prostate Tissues. BioMed Research International, 2020, 2020, 1-4.	0.9	9
27	Functional analyses of a low-penetrance risk variant rs6702619/1p21.2 associating with colorectal cancer in Polish population. Acta Biochimica Polonica, 2019, 66, 365-370.	0.3	9
28	Selective Extracellular Signal-Regulated Kinase $1/2$ (ERK1/2) Inhibition by the SCH772984 Compound Attenuates In Vitro and In Vivo Inflammatory Responses and Prolongs Survival in Murine Sepsis Models. International Journal of Molecular Sciences, 2021, 22, 10204.	1.8	8
29	Higher genome variability within metabolism genes associates with recurrent Clostridium difficile infection. BMC Microbiology, 2021, 21, 36.	1.3	7
30	Loss of epidermal MCPIP1 is associated with aggressive squamous cell carcinoma. Journal of Experimental and Clinical Cancer Research, 2021, 40, 391.	3 . 5	6
31	Clinical Applicability of Whole-Exome Sequencing Exemplified by a Study in Young Adults with the Advanced Cryptogenic Cholestatic Liver Diseases. Gastroenterology Research and Practice, 2017, 2017, 1-8.	0.7	5
32	Mutation Profiling of Premalignant Colorectal Neoplasia. Gastroenterology Research and Practice, 2019, 2019, 1-9.	0.7	5
33	Analysis of the Role of FRMD5 in the Biology of Papillary Thyroid Carcinoma. International Journal of Molecular Sciences, 2021, 22, 6726.	1.8	5
34	Oncology Drug Repurposing for Sepsis Treatment. Biomedicines, 2022, 10, 921.	1.4	5
35	Transcriptomes of human mesenchymal cells isolated from the right ventricle and epicardial fat differ strikingly both directly after isolation and longâ€ŧerm culture. ESC Heart Failure, 2019, 6, 351-361.	1.4	4
36	High Prevalence of Genetically Related Clostridium Difficile Strains at a Single Hemato-Oncology Ward Over 10 Years. Frontiers in Microbiology, 2020, 11, 1618.	1.5	4

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37	GWAS Links New Variant in Long Non-Coding RNA LINC02006 with Colorectal Cancer Susceptibility. Biology, 2021, 10, 465.	1.3	4
38	The gastric microbiota in patients with Crohn's disease; a preliminary study. Scientific Reports, 2021, 11, 17866.	1.6	4
39	Mutational landscape of primary and recurrent Ewing sarcoma. Wspolczesna Onkologia, 2021, 25, 241-248.	0.7	4
40	Prediction of fetal blood group antigens from maternal plasma using Ion AmpliSeq HD technology. Transfusion, 2022, 62, 458-468.	0.8	3
41	Limited Practical Utility of Liquid Biopsy in the Treated Patients with Advanced Breast Cancer. Diagnostics, 2020, 10, 523.	1.3	2
42	Cytotoxic Efficacy and Resistance Mechanism of a TRAIL and VEGFA-Peptide Fusion Protein in Colorectal Cancer Models. International Journal of Molecular Sciences, 2021, 22, 3160.	1.8	2
43	IDH mutations to predict shorter survival in chondrosarcoma Journal of Clinical Oncology, 2016, 34, 11024-11024.	0.8	1
44	Gene Expression-Based Functional Differences between the Bladder Body and Trigonal Urothelium in Adolescent Female Patients with Micturition Dysfunction. Biomedicines, 2022, 10, 1435.	1.4	1
45	Peripheral Blood Cells from Patients with Hodgkin's and Diffuse Large B Cell Lymphomas May Be a Better Source of Candidate Diagnostic miRNAs Than Circulating miRNAs. BioMed Research International, 2021, 2021, 1-9.	0.9	O