Kahraman Tanriverdi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Stimulation of Toll-Like Receptor 2 in Human Platelets Induces a Thromboinflammatory Response Through Activation of Phosphoinositide 3-Kinase. Circulation Research, 2009, 104, 346-354.	2.0	231
2	Diverse human extracellular RNAs are widely detected in human plasma. Nature Communications, 2016, 7, 11106.	5.8	170
3	The Extracellular RNA Communication Consortium: Establishing Foundational Knowledge and Technologies for Extracellular RNA Research. Cell, 2019, 177, 231-242.	13.5	152
4	Integrated genome-wide analysis of expression quantitative trait loci aids interpretation of genomic association studies. Genome Biology, 2017, 18, 16.	3.8	151
5	Interleukin 1 Receptor 1 and Interleukin 1β Regulate Megakaryocyte Maturation, Platelet Activation, and Transcript Profile During Inflammation in Mice and Humans. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 552-564.	1.1	136
6	Genome-wide identification of microRNA expression quantitative trait loci. Nature Communications, 2015, 6, 6601.	5.8	134
7	Comprehensive multi-center assessment of small RNA-seq methods for quantitative miRNA profiling. Nature Biotechnology, 2018, 36, 746-757.	9.4	134
8	SARS-CoV-2 Initiates Programmed Cell Death in Platelets. Circulation Research, 2021, 129, 631-646.	2.0	126
9	Relation of Platelet and Leukocyte Inflammatory Transcripts to Body Mass Index in the Framingham Heart Study. Circulation, 2010, 122, 119-129.	1.6	121
10	Ageâ€associated micro <scp>RNA</scp> expression in human peripheral blood is associated with allâ€cause mortality and ageâ€related traits. Aging Cell, 2018, 17, e12687.	3.0	114
11	Gene Expression Signatures of Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 1418-1426.	1.1	105
12	Plasma microRNAs are associated with atrial fibrillation and change after catheter ablation (the) Tj ETQq0 0 0 rgB	T /Oyerloc 0.3	k 10 Tf 50 3 101
13	Regulatory effects of TLR2 on megakaryocytic cell function. Blood, 2011, 117, 5963-5974.	0.6	91
14	Sex Differences in Platelet Toll-Like Receptors and Their Association With Cardiovascular Risk Factors. Arteriosclerosis, Thrombosis, and Vascular Biology, 2015, 35, 1030-1037.	1.1	91
15	Relations between circulating microRNAs and atrial fibrillation: Data from the Framingham Offspring Study. Heart Rhythm, 2014, 11, 663-669.	0.3	80

17	Extracellular RNAs Are Associated With Insulin Resistance and Metabolic Phenotypes. Diabetes Care, 2017, 40, 546-553.	4.3	73

18The role of RNA uptake in platelet heterogeneity. Thrombosis and Haemostasis, 2017, 117, 948-961.1.868

Reduced Adipose Tissue Inflammation Represents an Intermediate Cardiometabolic Phenotype in Obesity. Journal of the American College of Cardiology, 2011, 58, 232-237.

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19	Circulating microRNAs miR-331 and miR-195 differentiate local luminal a from metastatic breast cancer. BMC Cancer, 2019, 19, 436.	1.1	68
20	PRAME mRNA levels in cases with acute leukemia: Clinical importance and future prospects. American Journal of Hematology, 2005, 79, 257-261.	2.0	66
21	Micro-RNA (miRNA) profile in Hodgkin lymphoma: association between clinical and pathological variables. Medical Oncology, 2016, 33, 34.	1.2	60
22	Circulating Cell and Plasma microRNA Profiles Differ between Non-STSegment and ST-Segment-Elevation Myocardial Infarction. Family Medicine & Medical Science Research, 2013, 02, 108.	0.1	58
23	Relationship Among Circulating Inflammatory Proteins, Platelet Gene Expression, and Cardiovascular Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 2666-2673.	1.1	56
24	Dissecting the Roles of MicroRNAs in Coronary Heart Disease via Integrative Genomic Analyses. Arteriosclerosis, Thrombosis, and Vascular Biology, 2015, 35, 1011-1021.	1.1	53
25	Plasma Circulating Extracellular RNAs in Left Ventricular Remodeling Post-Myocardial Infarction. EBioMedicine, 2018, 32, 172-181.	2.7	52
26	Meeting report: discussions and preliminary findings on extracellular RNA measurement methods from laboratories in the NIH Extracellular RNA Communication Consortium. Journal of Extracellular Vesicles, 2015, 4, 26533.	5.5	51
27	Genetic associations with expression for genes implicated in GWAS studies for atherosclerotic cardiovascular disease and blood phenotypes. Human Molecular Genetics, 2014, 23, 782-795.	1.4	49
28	MicroRNA Signature of Cigarette Smoking and Evidence for a Putative Causal Role of MicroRNAs in Smoking-Related Inflammation and Target Organ Damage. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	45
29	The distribution of circulating microRNA and their relation to coronary disease. F1000Research, 2012, 1, 50.	0.8	40
30	Stroke and Circulating Extracellular RNAs. Stroke, 2017, 48, 828-834.	1.0	35
31	Small RNA-seq during acute maximal exercise reveal RNAs involved in vascular inflammation and cardiometabolic health: brief report. American Journal of Physiology - Heart and Circulatory Physiology, 2017, 313, H1162-H1167.	1.5	34
32	Messenger RNA and MicroRNA transcriptomic signatures of cardiometabolic risk factors. BMC Genomics, 2017, 18, 139.	1.2	33
33	Circulating MicroRNAs as Potential Biomarkers for Traumatic Brain Injury-Induced Hypopituitarism. Journal of Neurotrauma, 2016, 33, 1818-1825.	1.7	32
34	PRAME mRNA levels in cases with chronic leukemia: Clinical importance and review of the literature. Leukemia Research, 2007, 31, 365-369.	0.4	30
35	Discordant Expression of Circulating microRNA from Cellular and Extracellular Sources. PLoS ONE, 2016, 11, e0153691.	1.1	30
36	Comparison of RNA isolation and associated methods for extracellular RNA detection by high-throughput quantitative polymerase chain reaction. Analytical Biochemistry, 2016, 501, 66-74.	1.1	17

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#	Article	IF	CITATIONS
37	Micro-RNAs Are Related to Epicardial Adipose Tissue in Participants With Atrial Fibrillation: Data From the MiRhythm Study. Frontiers in Cardiovascular Medicine, 2019, 6, 115.	1.1	17
38	Unique Circulating MicroRNA Profiles in HIV Infection. Journal of Acquired Immune Deficiency Syndromes (1999), 2018, 79, 644-650.	0.9	16
39	Epigenome-wide association study of DNA methylation and microRNA expression highlights novel pathways for human complex traits. Epigenetics, 2020, 15, 183-198.	1.3	15
40	The Dynamic Platelet Transcriptome in Obesity and Weight Loss. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 854-864.	1.1	15
41	Crossâ€sectional relations of wholeâ€blood mi <scp>RNA</scp> expression levels and hand grip strength in a community sample. Aging Cell, 2017, 16, 888-894.	3.0	13
42	The role of the blood transcriptome in innate inflammation and stroke. Annals of the New York Academy of Sciences, 2010, 1207, 41-45.	1.8	12
43	Relations between plasma microRNAs, echocardiographic markers of atrial remodeling, and atrial fibrillation: Data from the Framingham Offspring study. PLoS ONE, 2020, 15, e0236960.	1.1	10
44	Specific Inflammatory Stimuli Lead to Distinct Platelet Responses in Mice and Humans. PLoS ONE, 2015, 10, e0131688.	1.1	8
45	Pollen-derived RNAs Are Found in the Human Circulation. IScience, 2019, 19, 916-926.	1.9	7
46	Platelet functional and transcriptional changes induced by intralipid infusion. Thrombosis and Haemostasis, 2016, 115, 1147-1156.	1.8	6
47	Micro RNAs from DNA Viruses are Found Widely in Plasma in a Large Observational Human Population. Scientific Reports, 2018, 8, 6397.	1.6	6
48	A Translational Model for Venous Thromboembolism: MicroRNA Expression in Hibernating Black Bears. Journal of Surgical Research, 2021, 257, 203-212.	0.8	6
49	Plasma MicroRNAs Relate to Atrial Fibrillation Recurrence after Catheter Ablation: Longitudinal Findings from the MiRhythm Study. Journal of Clinical & Experimental Cardiology, 2017, 08, .	0.0	5
50	Identifying miRNA Biomarkers and Predicted Targets Associated with Venous Thromboembolism in Colorectal Cancer Patients. Blood, 2019, 134, 3643-3643.	0.6	5
51	Circulating extracellular RNAs, myocardial remodeling, and heart failure in patients with acute coronary syndrome. Journal of Clinical and Translational Research, 2019, 5, 33-43.	0.3	4