

# Muhammad Arif

## List of Publications by Year in descending order

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Version: 2024-02-01

30  
papers

4,040  
citations

430874

18  
h-index

454955

30  
g-index

46  
all docs

46  
docs citations

46  
times ranked

7510  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiomics Analysis Reveals the Impact of Microbiota on Host Metabolism in Hepatic Steatosis. <i>Advanced Science</i> , 2022, 9, e2104373.	11.2	23
2	Machine Learning Analysis Reveals Biomarkers for the Detection of Neurological Diseases. <i>Frontiers in Molecular Neuroscience</i> , 2022, 15, .	2.9	3
3	Discovery of Functional Alternatively Spliced PKM Transcripts in Human Cancers. <i>Cancers</i> , 2021, 13, 348.	3.7	8
4	Spatiotemporal dissection of the cell cycle with single-cell proteogenomics. <i>Nature</i> , 2021, 590, 649-654.	27.8	104
5	iNetModels 2.0: an interactive visualization and database of multi-omics data. <i>Nucleic Acids Research</i> , 2021, 49, W271-W276.	14.5	25
6	Molecular pathways behind acquired obesity: Adipose tissue and skeletal muscle multiomics in monozygotic twin pairs discordant for BMI. <i>Cell Reports Medicine</i> , 2021, 2, 100226.	6.5	31
7	Integrative transcriptomic analysis of tissue-specific metabolic crosstalk after myocardial infarction. <i>ELife</i> , 2021, 10, .	6.0	20
8	Combined Metabolic Activators Accelerates Recovery in Mild to Moderate COVID-19. <i>Advanced Science</i> , 2021, 8, e2101222.	11.2	49
9	A single-cell type transcriptomics map of human tissues. <i>Science Advances</i> , 2021, 7, .	10.3	632
10	Stratification of patients with clear cell renal cell carcinoma to facilitate drug repositioning. <i>IScience</i> , 2021, 24, 102722.	4.1	8
11	Glucosylceramide synthase deficiency in the heart compromises $\beta$ 1-adrenergic receptor trafficking. <i>European Heart Journal</i> , 2021, 42, 4481-4492.	2.2	14
12	Systems Analysis Reveals Ageing-Related Perturbations in Retinoids and Sex Hormones in Alzheimer's and Parkinson's Diseases. <i>Biomedicines</i> , 2021, 9, 1310.	3.2	8
13	Combined metabolic activators therapy ameliorates liver fat in nonalcoholic fatty liver disease patients. <i>Molecular Systems Biology</i> , 2021, 17, e10459.	7.2	22
14	A network-based approach reveals the dysregulated transcriptional regulation in non-alcoholic fatty liver disease. <i>IScience</i> , 2021, 24, 103222.	4.1	14
15	Revealing the Molecular Mechanisms of Alzheimer's Disease Based on Network Analysis. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11556.	4.1	10
16	Next generation plasma proteome profiling of COVID-19 patients with mild to moderate symptoms. <i>EBioMedicine</i> , 2021, 74, 103723.	6.1	26
17	Skeletal Muscle Transcriptomic Comparison between Long-Term Trained and Untrained Men and Women. <i>Cell Reports</i> , 2020, 31, 107808.	6.4	39
18	Integration of molecular profiles in a longitudinal wellness profiling cohort. <i>Nature Communications</i> , 2020, 11, 4487.	12.8	66

#	ARTICLE	IF	CITATIONS
19	Classification of clear cell renal cell carcinoma based on PKM alternative splicing. <i>Heliyon</i> , 2020, 6, e03440.	3.2	9
20	Boosting Natural Killer Cell-Mediated Targeting of Sarcoma Through DNAM-1 and NKG2D. <i>Frontiers in Immunology</i> , 2020, 11, 40.	4.8	40
21	The acute effect of metabolic cofactor supplementation: a potential therapeutic strategy against non-alcoholic fatty liver disease. <i>Molecular Systems Biology</i> , 2020, 16, e9495.	7.2	39
22	Elucidating the Reprogramming of Colorectal Cancer Metabolism Using Genome-Scale Metabolic Modeling. <i>Frontiers in Oncology</i> , 2019, 9, 681.	2.8	40
23	Cardiac expression of the microsomal triglyceride transport protein protects the heart function during ischemia. <i>Journal of Molecular and Cellular Cardiology</i> , 2019, 137, 1-8.	1.9	3
24	Characterization of heterogeneous redox responses in hepatocellular carcinoma patients using network analysis. <i>EBioMedicine</i> , 2019, 40, 471-487.	6.1	38
25	Pyruvate kinase L/R is a regulator of lipid metabolism and mitochondrial function. <i>Metabolic Engineering</i> , 2019, 52, 263-272.	7.0	37
26	TCSBN: a database of tissue and cancer specific biological networks. <i>Nucleic Acids Research</i> , 2018, 46, D595-D600.	14.5	55
27	Characterization of different fat depots in NAFLD using inflammation-associated proteome, lipidome and metabolome. <i>Scientific Reports</i> , 2018, 8, 14200.	3.3	28
28	ESS: A Tool for Genome-Scale Quantification of Essentiality Score for Reaction/Genes in Constraint-Based Modeling. <i>Frontiers in Physiology</i> , 2018, 9, 1355.	2.8	8
29	Metabolic Network-Based Identification and Prioritization of Anticancer Targets Based on Expression Data in Hepatocellular Carcinoma. <i>Frontiers in Physiology</i> , 2018, 9, 916.	2.8	38
30	A pathology atlas of the human cancer transcriptome. <i>Science</i> , 2017, 357, .	12.6	2,570