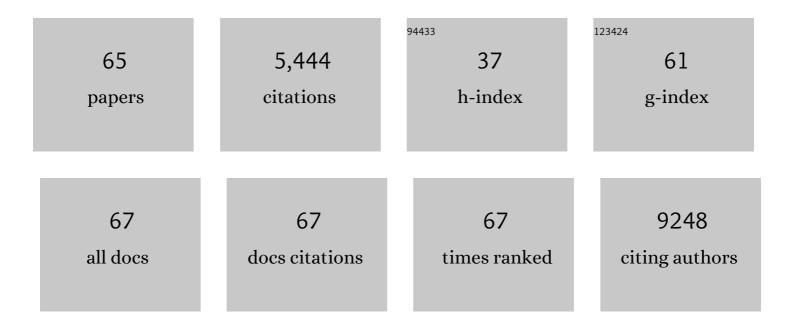
List of Publications by Year in descending order

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DEEDAK NACDATH

#	Article	IF	CITATIONS
1	Tumor microenvironment derived exosomes pleiotropically modulate cancer cell metabolism. ELife, 2016, 5, e10250.	6.0	681
2	Glutaminolysis: A Hallmark of Cancer Metabolism. Annual Review of Biomedical Engineering, 2017, 19, 163-194.	12.3	528
3	Targeting Stromal Glutamine Synthetase in Tumors Disrupts Tumor Microenvironment-Regulated Cancer Cell Growth. Cell Metabolism, 2016, 24, 685-700.	16.2	293
4	Reactive Oxygen Species in the Tumor Microenvironment: An Overview. Cancers, 2019, 11, 1191.	3.7	288
5	The role of stromal cancer-associated fibroblasts in pancreatic cancer. Journal of Hematology and Oncology, 2017, 10, 76.	17.0	281
6	Metabolic shifts toward glutamine regulate tumor growth, invasion and bioenergetics in ovarian cancer. Molecular Systems Biology, 2014, 10, 728.	7.2	255
7	Genomic deletion of malic enzyme 2 confers collateral lethality in pancreatic cancer. Nature, 2017, 542, 119-123.	27.8	209
8	Energy stress-induced lncRNA FILNC1 represses c-Myc-mediated energy metabolism and inhibits renal tumor development. Nature Communications, 2017, 8, 783.	12.8	157
9	Three-Dimensional Primary Hepatocyte Culture in Synthetic Self-Assembling Peptide Hydrogel. Tissue Engineering - Part A, 2008, 14, 227-236.	3.1	144
10	Metabolic preconditioning of donor organs: Defatting fatty livers by normothermic perfusion ex vivo. Metabolic Engineering, 2009, 11, 274-283.	7.0	139
11	The key role of extracellular vesicles in the metastatic process. Biochimica Et Biophysica Acta: Reviews on Cancer, 2018, 1869, 64-77.	7.4	119
12	Tumour-reprogrammed stromal BCAT1 fuels branched-chain ketoacid dependency in stromal-rich PDAC tumours. Nature Metabolism, 2020, 2, 775-792.	11.9	110
13	The glucoseâ€deprivation network counteracts lapatinibâ€induced toxicity in resistant ErbB2â€positive breast cancer cells. Molecular Systems Biology, 2012, 8, 596.	7.2	109
14	Lactate-mediated epigenetic reprogramming regulates formation of human pancreatic cancer-associated fibroblasts. ELife, 2019, 8, .	6.0	103
15	A switch in the source of ATP production and a loss in capacity to perform glycolysis are hallmarks of hepatocyte failure in advance liver disease. Journal of Hepatology, 2014, 60, 1203-1211.	3.7	99
16	Amplification of USP13 drives ovarian cancer metabolism. Nature Communications, 2016, 7, 13525.	12.8	99
17	Human Omental-Derived Adipose Stem Cells Increase Ovarian Cancer Proliferation, Migration, and Chemoresistance. PLoS ONE, 2013, 8, e81859.	2.5	95
18	Generation of systemic antitumour immunity via the in situ modulation of the gut microbiome by an orally administered inulin gel. Nature Biomedical Engineering, 2021, 5, 1377-1388.	22.5	95

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19	Oxygen uptake rates and liver-specific functions of hepatocyte and 3T3 fibroblast co-cultures. Biotechnology and Bioengineering, 2007, 97, 188-199.	3.3	86
20	Pyruvate uptake is increased in highly invasive ovarian cancer cells under anoikis conditions for anaplerosis, mitochondrial function, and migration. American Journal of Physiology - Endocrinology and Metabolism, 2012, 303, E1036-E1052.	3.5	83
21	Nitric Oxide: The Forgotten Child of Tumor Metabolism. Trends in Cancer, 2017, 3, 659-672.	7.4	78
22	Mutant Kras- and p16-regulated NOX4 activation overcomes metabolic checkpoints in development of pancreatic ductal adenocarcinoma. Nature Communications, 2017, 8, 14437.	12.8	77
23	Generation of Human Fatty Livers Using Custom-Engineered Induced Pluripotent Stem Cells with Modifiable SIRT1 Metabolism. Cell Metabolism, 2019, 30, 385-401.e9.	16.2	75
24	Nitric Oxide Mediates Metabolic Coupling of Omentum-Derived Adipose Stroma to Ovarian and Endometrial Cancer Cells. Cancer Research, 2015, 75, 456-471.	0.9	70
25	Nitric oxide is a positive regulator of the Warburg effect in ovarian cancer cells. Cell Death and Disease, 2014, 5, e1302-e1302.	6.3	69
26	Effect of pH changes on water release values in hydrophobic interaction chromatographic systems. Journal of Chromatography A, 2005, 1079, 229-235.	3.7	64
27	Exo-MFA – A 13C metabolic flux analysis framework to dissect tumor microenvironment-secreted exosome contributions towards cancer cell metabolism. Metabolic Engineering, 2017, 43, 156-172.	7.0	63
28	Integrated Energy and Flux Balance Based Multiobjective Framework for Large-Scale Metabolic Networks. Annals of Biomedical Engineering, 2007, 35, 863-885.	2.5	62
29	Electron transport chain activity is a predictor and target for venetoclax sensitivity in multiple myeloma. Nature Communications, 2020, 11, 1228.	12.8	62
30	Metabolomics for mitochondrial and cancer studies. Biochimica Et Biophysica Acta - Bioenergetics, 2011, 1807, 650-663.	1.0	60
31	Use and Optimization of a Dual-Flowrate Loading Strategy To Maximize Throughput in Protein-A Affinity Chromatography. Biotechnology Progress, 2004, 20, 830-840.	2.6	57
32	Microfluidic device for high-throughput affinity-based isolation of extracellular vesicles. Lab on A Chip, 2020, 20, 1762-1770.	6.0	57
33	Role of Increased n-acetylaspartate Levels in Cancer. Journal of the National Cancer Institute, 2016, 108, djv426.	6.3	51
34	Evaluation of selectivity changes in HIC systems using a preferential interaction based analysis. Biotechnology and Bioengineering, 2004, 87, 354-363.	3.3	48
35	Circulating tumor cells in precision medicine: challenges and opportunities. Trends in Pharmacological Sciences, 2022, 43, 378-391.	8.7	47
36	Evolution of intrahepatic carbon, nitrogen, and energy metabolism in a D-galactosamine-induced rat liver failure model. Metabolic Engineering, 2005, 7, 88-103.	7.0	40

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37	Modeling of adsorption in hydrophobic interaction chromatography systems using a preferential interaction quadratic isotherm. Journal of Chromatography A, 2003, 989, 47-54.	3.7	38
38	A Hybrid Model Framework for the Optimization of Preparative Chromatographic Processes. Biotechnology Progress, 2008, 20, 162-178.	2.6	34
39	Oncosecretomics coupled to bioenergetics identifies α-amino adipic acid, isoleucine and GABA as potential biomarkers of cancer: Differential expression of c-Myc, Oct1 and KLF4 coordinates metabolic changes. Biochimica Et Biophysica Acta - Bioenergetics, 2012, 1817, 2060-2071.	1.0	34
40	Soft constraints-based multiobjective framework for flux balance analysis. Metabolic Engineering, 2010, 12, 429-445.	7.0	33
41	A model predictive formulation for control of open-loop unstable cascade systems. Chemical Engineering Science, 2002, 57, 365-378.	3.8	31
42	Characterization and modeling of nonlinear hydrophobic interaction chromatographic systems. Journal of Chromatography A, 2011, 1218, 1219-1226.	3.7	29
43	Targeting integrated epigenetic and metabolic pathways in lethal childhood PFA ependymomas. Science Translational Medicine, 2021, 13, eabc0497.	12.4	29
44	ITLN1 modulates invasive potential and metabolic reprogramming of ovarian cancer cells in omental microenvironment. Nature Communications, 2020, 11, 3546.	12.8	28
45	HSulf-1 deficiency dictates a metabolic reprograming of glycolysis and TCA cycle in ovarian cancer. Oncotarget, 2015, 6, 33705-33719.	1.8	28
46	Multiobjective optimization strategies for linear gradient chromatography. AICHE Journal, 2005, 51, 511-525.	3.6	26
47	Adipocyteâ€derived basement membrane extract with biological activity: applications in hepatocyte functional augmentation <i>in vitro</i> . FASEB Journal, 2010, 24, 2364-2374.	0.5	24
48	The Lung Microbiome: A Central Mediator of Host Inflammation and Metabolism in Lung Cancer Patients?. Cancers, 2021, 13, 13.	3.7	21
49	Evolutionary operation and control of chromatographic processes. AICHE Journal, 2003, 49, 82-95.	3.6	17
50	Radiotherapy-induced metabolic hallmarks in the tumor microenvironment. Trends in Cancer, 2022, 8, 855-869.	7.4	17
51	Modeling Integrated Cellular Machinery Using Hybrid Petri-Boolean Networks. PLoS Computational Biology, 2013, 9, e1003306.	3.2	14
52	Metabolic regulation of collagen gel contraction by porcine aortic valvular interstitial cells. Journal of the Royal Society Interface, 2014, 11, 20140852.	3.4	14
53	Cellular Location of HNF4α is Linked With Terminal Liver Failure in Humans. Hepatology Communications, 2020, 4, 859-875.	4.3	12
54	Metabolic Profiling Based Quantitative Evaluation of Hepatocellular Metabolism in Presence of Adipocyte Derived Extracellular Matrix. PLoS ONE, 2011, 6, e20137.	2.5	11

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55	Regulation of protein metabolism in cancer. Molecular and Cellular Oncology, 2018, 5, e1285384.	0.7	9
56	Metabolic Reprogramming and Vulnerabilities in Cancer. Cancers, 2020, 12, 90.	3.7	8
57	Liver Tissue Engineering. , 2011, , 389-419.		7
58	Synthesis and Biological Evaluation of Dimeric Furanoid Macroheterocycles: Discovery of New Anticancer Agents. Journal of the American Chemical Society, 2015, 137, 4766-4770.	13.7	7
59	Quantifying Metabolic Transfer Mediated by Extracellular Vesicles Using Exo-MFA: An Integrated Empirical and Computational Platform. Methods in Molecular Biology, 2020, 2088, 205-221.	0.9	5
60	Linking omentum and ovarian cancer: NO. Oncoscience, 2015, 2, 797-798.	2.2	5
61	Optimality and thermodynamics determine the evolution of transcriptional regulatory networks. Molecular BioSystems, 2012, 8, 511-530.	2.9	4
62	Liquid Chromatography Methods for Separation of Polar and Charged Intracellular Metabolites for 13C Metabolic Flux Analysis. Methods in Molecular Biology, 2020, 2088, 33-50.	0.9	4
63	Reply to: "ls the pathway of energy metabolism modified in advanced cirrhosis?― Journal of Hepatology, 2014, 61, 453.	3.7	0
64	Mitochondrial Electron Transport Chain Inhibition Promotes Resistance to Proteasome Inhibitors in Multiple Myeloma. Blood, 2021, 138, 1611-1611.	1.4	0
65	Driving with Both Feet: Supplementing AKG While Inhibiting BCAT1 Leads to Synthetic Lethality in GBM. Cancer Research, 2022, 82, 2354-2356.	0.9	0