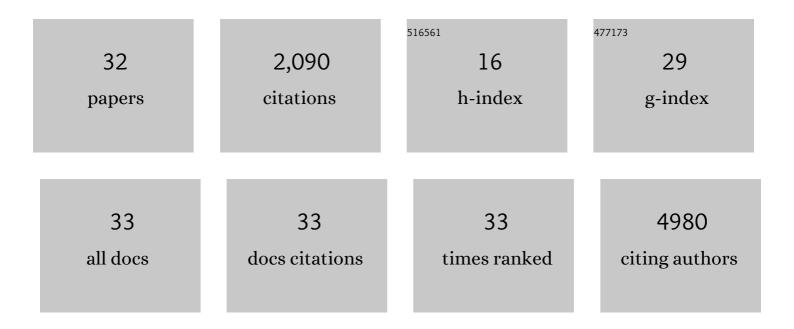
Yasser Riazalhosseini

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Targeting HIF1-alpha/miR-326/ITGA5 axis potentiates chemotherapy response in triple-negative breast cancer. Breast Cancer Research and Treatment, 2022, 193, 331-348.	1.1	18
2	Cell–Cell Contact Mediates Gene Expression and Fate Choice of Human Neural Stem/Progenitor Cells. Cells, 2022, 11, 1741.	1.8	2
3	Morphological findings in frozen non-neoplastic kidney tissues of patients with kidney cancer from large-scale multicentric studies on renal cancer. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2021, 478, 1099-1107.	1.4	1
4	Best Practices for Spatial Profiling for Breast Cancer Research with the GeoMx® Digital Spatial Profiler. Cancers, 2021, 13, 4456.	1.7	50
5	A Preclinical Trial and Molecularly Annotated Patient Cohort Identify Predictive Biomarkers in Homologous Recombination–deficient Pancreatic Cancer. Clinical Cancer Research, 2020, 26, 5462-5476.	3.2	20
6	Whole genome sequencing identifies a duplicated region encompassing Xq13.2q13.3 in a large Iranian family with intellectual disability. Molecular Genetics & Genomic Medicine, 2020, 8, e1418.	0.6	1
7	Targeting lysyl oxidase (LOX) overcomes chemotherapy resistance in triple negative breast cancer. Nature Communications, 2020, 11, 2416.	5.8	179
8	Latency and interval therapy affect the evolution in metastatic colorectal cancer. Scientific Reports, 2020, 10, 581.	1.6	6
9	The potential of dopamine receptor D2 (DRD2) as a therapeutic target for tackling pancreatic cancer. Expert Opinion on Therapeutic Targets, 2019, 23, 365-367.	1.5	12
10	Age-related variations in gene expression patterns of renal cell carcinoma. Urologic Oncology: Seminars and Original Investigations, 2019, 37, 166-175.	0.8	8
11	Consequences of VHL Loss on Global DNA Methylome. Scientific Reports, 2018, 8, 3313.	1.6	16
12	Reactivation of cAMP Pathway by PDE4D Inhibition Represents a Novel Druggable Axis for Overcoming Tamoxifen Resistance in ER-positive Breast Cancer. Clinical Cancer Research, 2018, 24, 1987-2001.	3.2	37
13	The genomic landscape of two Burkitt lymphoma cases and derived cell lines: comparison between primary and relapse samples. Leukemia and Lymphoma, 2018, 59, 2159-2174.	0.6	6
14	Modules of Correlated Genes in a Gene Expression Regulatory Network of CDDP-Resistant Cancer Cells. , 2018, , .		0
15	A General Framework for Interrogation of mRNA Stability Programs Identifies RNA-Binding Proteins that Govern Cancer Transcriptomes. Cell Reports, 2018, 23, 1639-1650.	2.9	56
16	Aberration hubs in protein interaction networks highlight actionable targets in cancer. Oncotarget, 2018, 9, 25166-25180.	0.8	6
17	Loss of chromosome Y leads to down regulation of KDM5D and KDM6C epigenetic modifiers in clear cell renal cell carcinoma. Scientific Reports, 2017, 7, 44876.	1.6	42
18	Epigenome Aberrations: Emerging Driving Factors of the Clear Cell Renal Cell Carcinoma. International Journal of Molecular Sciences, 2017, 18, 1774	1.8	46

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#	Article	IF	CITATIONS
19	Precision medicine from the renal cancer genome. Nature Reviews Nephrology, 2016, 12, 655-666.	4.1	64
20	Expression of DRD2 Is Increased in Human Pancreatic Ductal Adenocarcinoma and Inhibitors Slow Tumor Growth in Mice. Gastroenterology, 2016, 151, 1218-1231.	0.6	100
21	RNA-Seq as a Tool to Study the Tumor Microenvironment. Methods in Molecular Biology, 2016, 1458, 311-337.	0.4	7
22	Genetic Landscapes of Relapsed and Refractory Diffuse Large B-Cell Lymphomas. Clinical Cancer Research, 2016, 22, 2290-2300.	3.2	186
23	<i>CHSR</i> hypermethylation: a promising pan-cancer marker. Cell Cycle, 2015, 14, 689-690.	1.3	10
24	TCF12 is mutated in anaplastic oligodendroglioma. Nature Communications, 2015, 6, 7207.	5.8	42
25	The Prognostic Ease and Difficulty of Invasive Breast Carcinoma. Cell Reports, 2014, 9, 129-142.	2.9	64
26	Rare missense variants in POT1 predispose to familial cutaneous malignant melanoma. Nature Genetics, 2014, 46, 482-486.	9.4	283
27	Variation in genomic landscape of clear cell renal cell carcinoma across Europe. Nature Communications, 2014, 5, 5135.	5.8	158
28	Germline and somatic SMARCA4 mutations characterize small cell carcinoma of the ovary, hypercalcemic type. Nature Genetics, 2014, 46, 438-443.	9.4	383
29	Diagnostic values of GHSR DNA methylation pattern in breast cancer. Breast Cancer Research and Treatment, 2012, 135, 705-713.	1.1	16
30	Epigenetically Deregulated microRNA-375 Is Involved in a Positive Feedback Loop with Estrogen Receptor α in Breast Cancer Cells. Cancer Research, 2010, 70, 9175-9184.	0.4	260
31	Do we use the appropriate controls for the identification of informative methylation markers for early cancer detection?. Genome Biology, 2008, 9, 405.	13.9	9
32	A multi-layer post-transcriptional gene regulatory program fuels cancer angiogenesis and metastasis. Biotarget, 0, 3, 11-11.	0.5	0