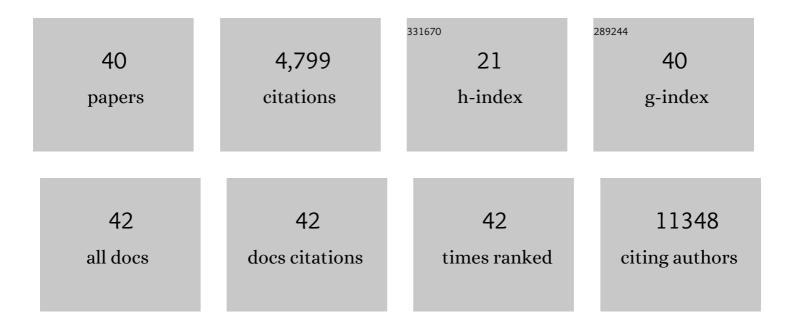
## Syed Haider

List of Publications by Year in descending order

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SVED HAIDER

#	Article	IF	CITATIONS
1	Functional annotation of breast cancer risk loci: current progress and future directions. British Journal of Cancer, 2022, 126, 981-993.	6.4	6
2	Age influences on the molecular presentation of tumours. Nature Communications, 2022, 13, 208.	12.8	26
3	Assessment of the Molecular Heterogeneity of E-Cadherin Expression in Invasive Lobular Breast Cancer. Cancers, 2022, 14, 295.	3.7	5
4	Characterisation of the Stromal Microenvironment in Lobular Breast Cancer. Cancers, 2022, 14, 904.	3.7	13
5	Liver glycogen phosphorylase is upregulated in glioblastoma and provides a metabolic vulnerability to high dose radiation. Cell Death and Disease, 2022, 13, .	6.3	6
6	Increased expression of glutamine transporter SNAT2/SLC38A2 promotes glutamine dependence and oxidative stress resistance, and is associated with worse prognosis in triple-negative breast cancer. British Journal of Cancer, 2021, 124, 494-505.	6.4	62
7	Identifying high-confidence capture Hi-C interactions using CHiCANE. Nature Protocols, 2021, 16, 2257-2285.	12.0	11
8	Predicting the clinical outcome of oral potentially malignant disorders using transcriptomic-based molecular pathology. British Journal of Cancer, 2021, 125, 413-421.	6.4	16
9	Interferon- and STING-independent induction of type I interferon stimulated genes during fractionated irradiation. Journal of Experimental and Clinical Cancer Research, 2021, 40, 161.	8.6	16
10	Impairment of a distinct cancer-associated fibroblast population limits tumour growth and metastasis. Nature Communications, 2021, 12, 3516.	12.8	35
11	PolÎ, inhibitors elicit BRCA-gene synthetic lethality and target PARP inhibitor resistance. Nature Communications, 2021, 12, 3636.	12.8	159
12	Quantitative Assessment and Prognostic Associations of the Immune Landscape in Ovarian Clear Cell Carcinoma. Cancers, 2021, 13, 3854.	3.7	10
13	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. American Journal of Human Genetics, 2021, 108, 1190-1203.	6.2	6
14	The Mutational Concordance of Fixed Formalin Paraffin Embedded and Fresh Frozen Gastro-Oesophageal Tumours Using Whole Exome Sequencing. Journal of Clinical Medicine, 2021, 10, 215.	2.4	5
15	ELTD1 Activation Induces an Endothelial-EMT Transition to a Myofibroblast Phenotype. International Journal of Molecular Sciences, 2021, 22, 11293.	4.1	6
16	Sirtuin inhibition is synthetic lethal with BRCA1 or BRCA2 deficiency. Communications Biology, 2021, 4, 1270.	4.4	4
17	Sex differences in oncogenic mutational processes. Nature Communications, 2020, 11, 4330.	12.8	60
18	Targeting TRIM37-driven centrosome dysfunction in 17q23-amplified breast cancer. Nature, 2020, 585, 447-452.	27.8	63

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19	Systematic Assessment of Tumor Purity and Its Clinical Implications. JCO Precision Oncology, 2020, 4, 995-1005.	3.0	23
20	Landscape of transcriptomic interactions between breast cancer and its microenvironment. Nature Communications, 2019, 10, 3116.	12.8	16
21	BPG: Seamless, automated and interactive visualization of scientific data. BMC Bioinformatics, 2019, 20, 42.	2.6	64
22	Hypoxia-induced switch in SNAT2/SLC38A2 regulation generates endocrine resistance in breast cancer. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 12452-12461.	7.1	86
23	Adaptation to HIF1α Deletion in Hypoxic Cancer Cells by Upregulation of GLUT14 and Creatine Metabolism. Molecular Cancer Research, 2019, 17, 1531-1544.	3.4	22
24	ADGRL4/ELTD1 Silencing in Endothelial Cells Induces ACLY and SLC25A1 and Alters the Cellular Metabolic Profile. Metabolites, 2019, 9, 287.	2.9	14
25	3D Growth of Cancer Cells Elicits Sensitivity to Kinase Inhibitors but Not Lipid Metabolism Modifiers. Molecular Cancer Therapeutics, 2019, 18, 376-388.	4.1	17
26	RASSF1A uncouples Wnt from Hippo signalling and promotes YAP mediated differentiation via p73. Nature Communications, 2018, 9, 424.	12.8	72
27	Development and Validation of a 28-gene Hypoxia-related Prognostic Signature for Localized Prostate Cancer. EBioMedicine, 2018, 31, 182-189.	6.1	132
28	Capture Hi-C identifies putative target genes at 33 breast cancer risk loci. Nature Communications, 2018, 9, 1028.	12.8	98
29	Pathway-based subnetworks enable cross-disease biomarker discovery. Nature Communications, 2018, 9, 4746.	12.8	30
30	Sex Differences in Cancer Driver Genes and Biomarkers. Cancer Research, 2018, 78, 5527-5537.	0.9	108
31	A prosurvival DNA damage-induced cytoplasmic interferon response is mediated by end resection factors and is limited by Trex1. Genes and Development, 2017, 31, 353-369.	5.9	168
32	IGF-1R associates with adverse outcomes after radical radiotherapy for prostate cancer. British Journal of Cancer, 2017, 117, 1600-1606.	6.4	35
33	Developing and validating a multivariable predictive biomarker for treatment selection for oropharyngeal squamous cell carcinoma: The PREDICTR-OPC study Journal of Clinical Oncology, 2017, 35, 6004-6004.	1.6	3
34	Genomic alterations underlie a pan-cancer metabolic shift associated with tumour hypoxia. Genome Biology, 2016, 17, 140.	8.8	67
35	A bedr way of genomic interval processing. Source Code for Biology and Medicine, 2016, 11, 14.	1.7	21
36	ISOpureR: an R implementation of a computational purification algorithm of mixed tumour profiles. BMC Bioinformatics, 2015, 16, 156.	2.6	30

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37	The BioMart community portal: an innovative alternative to large, centralized data repositories. Nucleic Acids Research, 2015, 43, W589-W598.	14.5	682
38	The Molecular Taxonomy of Primary Prostate Cancer. Cell, 2015, 163, 1011-1025.	28.9	2,435
39	Nuclear HER4 mediates acquired resistance to trastuzumab and is associated with poor outcome in HER2 positive breast cancer. Oncotarget, 2014, 5, 5934-5949.	1.8	59
40	Computational purification of individual tumor gene expression profiles leads to significant improvements in prognostic prediction. Genome Medicine, 2013, 5, 29.	8.2	96