

# Harpreet Kaur

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7382637/publications.pdf>

Version: 2024-02-01

21  
papers

1,052  
citations

623734

14  
h-index

794594

19  
g-index

38  
all docs

38  
docs citations

38  
times ranked

1524  
citing authors

#	ARTICLE	IF	CITATIONS
1	Computational resources for identification of cancer biomarkers from omics data. Briefings in Functional Genomics, 2021, 20, 213-222.	2.7	7
2	Crowdsourcing assessment of maternal blood multi-omics for predicting gestational age and preterm birth. Cell Reports Medicine, 2021, 2, 100323.	6.5	47
3	SAPdb: A database of short peptides and the corresponding nanostructures formed by self-assembly. Computers in Biology and Medicine, 2021, 133, 104391.	7.0	10
4	ProCanBio: A Database of Manually Curated Biomarkers for Prostate Cancer. Journal of Computational Biology, 2021, 28, 1248-1257.	1.6	0
5	Community Assessment of the Predictability of Cancer Protein and Phosphoprotein Levels from Genomics and Transcriptomics. Cell Systems, 2020, 11, 186-195.e9.	6.2	19
6	Analysis and prediction of cholangiocarcinoma from transcriptomic profile of patients. Journal of Hepatology, 2020, 73, S16-S17.	3.7	4
7	A Web-Based Platform on Coronavirus Disease-19 to Maintain Predicted Diagnostic, Drug, and Vaccine Candidates. Monoclonal Antibodies in Immunodiagnosis and Immunotherapy, 2020, 39, 204-216.	1.6	25
8	CancerLivER: a database of liver cancer gene expression resources and biomarkers. Database: the Journal of Biological Databases and Curation, 2020, 2020, .	3.0	21
9	Multiple Myeloma DREAM Challenge reveals epigenetic regulator PHF19 as marker of aggressive disease. Leukemia, 2020, 34, 1866-1874.	7.2	36
10	Computing Skin Cutaneous Melanoma Outcome From the HLA-Alleles and Clinical Characteristics. Frontiers in Genetics, 2020, 11, 221.	2.3	33
11	Expression based biomarkers and models to classify early and late-stage samples of Papillary Thyroid Carcinoma. PLoS ONE, 2020, 15, e0231629.	2.5	13
12	Prediction and Analysis of Skin Cancer Progression using Genomics Profiles of Patients. Scientific Reports, 2019, 9, 15790.	3.3	57
13	RareLSD: a manually curated database of lysosomal enzymes associated with rare diseases. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	3.0	4
14	Classification of early and late stage liver hepatocellular carcinoma patients from their genomics and epigenomics profiles. PLoS ONE, 2019, 14, e0221476.	2.5	46
15	Identification of Platform-Independent Diagnostic Biomarker Panel for Hepatocellular Carcinoma Using Large-Scale Transcriptomics Data. Frontiers in Genetics, 2019, 10, 1306.	2.3	49
16	Computer-aided designing of immunosuppressive peptides based on IL-10 inducing potential. Scientific Reports, 2017, 7, 42851.	3.3	206
17	Gene expression-based biomarkers for discriminating early and late stage of clear cell renal cancer. Scientific Reports, 2017, 7, 44997.	3.3	92
18	CancerPDF: A repository of cancer-associated peptidome found in human biofluids. Scientific Reports, 2017, 7, 1511.	3.3	30

#	ARTICLE	IF	CITATIONS
19	PEPlife: A Repository of the Half-life of Peptides. Scientific Reports, 2016, 6, 36617.	3.3	108
20	PEPstr: A de novo Method for Tertiary Structure Prediction of Small Bioactive Peptides. Protein and Peptide Letters, 2007, 14, 626-631.	0.9	179
21	Identification of LncRNAs as Therapeutic Targets in Chronic Lymphocytic Leukemia. Columbia Undergraduate Science Journal, 0, 15, 16-29.	0.0	1