

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	Computer-aided designing of immunosuppressive peptides based on IL-10 inducing potential. <i>Scientific Reports</i> , 2017, 7, 42851.	3.3	206
2	PEPstr: A de novo Method for Tertiary Structure Prediction of Small Bioactive Peptides. <i>Protein and Peptide Letters</i> , 2007, 14, 626-631.	0.9	179
3	PEPlife: A Repository of the Half-life of Peptides. <i>Scientific Reports</i> , 2016, 6, 36617.	3.3	108
4	Gene expression-based biomarkers for discriminating early and late stage of clear cell renal cancer. <i>Scientific Reports</i> , 2017, 7, 44997.	3.3	92
5	Prediction and Analysis of Skin Cancer Progression using Genomics Profiles of Patients. <i>Scientific Reports</i> , 2019, 9, 15790.	3.3	57
6	Identification of Platform-Independent Diagnostic Biomarker Panel for Hepatocellular Carcinoma Using Large-Scale Transcriptomics Data. <i>Frontiers in Genetics</i> , 2019, 10, 1306.	2.3	49
7	Crowdsourcing assessment of maternal blood multi-omics for predicting gestational age and preterm birth. <i>Cell Reports Medicine</i> , 2021, 2, 100323.	6.5	47
8	Classification of early and late stage liver hepatocellular carcinoma patients from their genomics and epigenomics profiles. <i>PLoS ONE</i> , 2019, 14, e0221476.	2.5	46
9	Multiple Myeloma DREAM Challenge reveals epigenetic regulator PHF19 as marker of aggressive disease. <i>Leukemia</i> , 2020, 34, 1866-1874.	7.2	36
10	Computing Skin Cutaneous Melanoma Outcome From the HLA-Alleles and Clinical Characteristics. <i>Frontiers in Genetics</i> , 2020, 11, 221.	2.3	33
11	CancerPDF: A repository of cancer-associated peptidome found in human biofluids. <i>Scientific Reports</i> , 2017, 7, 1511.	3.3	30
12	A Web-Based Platform on Coronavirus Disease-19 to Maintain Predicted Diagnostic, Drug, and Vaccine Candidates. <i>Monoclonal Antibodies in Immunodiagnosis and Immunotherapy</i> , 2020, 39, 204-216.	1.6	25
13	CancerLivER: a database of liver cancer gene expression resources and biomarkers. <i>Database: the Journal of Biological Databases and Curation</i> , 2020, 2020, .	3.0	21
14	Community Assessment of the Predictability of Cancer Protein and Phosphoprotein Levels from Genomics and Transcriptomics. <i>Cell Systems</i> , 2020, 11, 186-195.e9.	6.2	19
15	Expression based biomarkers and models to classify early and late-stage samples of Papillary Thyroid Carcinoma. <i>PLoS ONE</i> , 2020, 15, e0231629.	2.5	13
16	SAPdb: A database of short peptides and the corresponding nanostructures formed by self-assembly. <i>Computers in Biology and Medicine</i> , 2021, 133, 104391.	7.0	10
17	Computational resources for identification of cancer biomarkers from omics data. <i>Briefings in Functional Genomics</i> , 2021, 20, 213-222.	2.7	7
18	RareLSD: a manually curated database of lysosomal enzymes associated with rare diseases. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	3.0	4

#	ARTICLE	IF	CITATIONS
19	Analysis and prediction of cholangiocarcinoma from transcriptomic profile of patients. Journal of Hepatology, 2020, 73, S16-S17.	3.7	4
20	Identification of LncRNAs as Therapeutic Targets in Chronic Lymphocytic Leukemia. Columbia Undergraduate Science Journal, 0, 15, 16-29.	0.0	1
21	ProCanBio: A Database of Manually Curated Biomarkers for Prostate Cancer. Journal of Computational Biology, 2021, 28, 1248-1257.	1.6	0