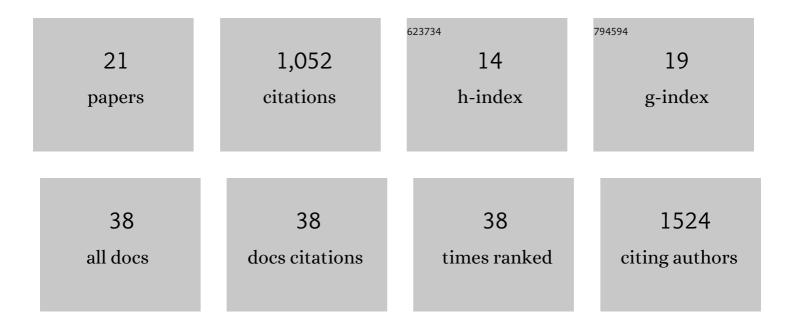
Harpreet Kaur

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7382637/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Computer-aided designing of immunosuppressive peptides based on IL-10 inducing potential. Scientific Reports, 2017, 7, 42851. | 3.3 | 206 |
| 2 | PEPstr: A de novo Method for Tertiary Structure Prediction of Small Bioactive Peptides. Protein and Peptide Letters, 2007, 14, 626-631. | 0.9 | 179 |
| 3 | PEPlife: A Repository of the Half-life of Peptides. Scientific Reports, 2016, 6, 36617. | 3.3 | 108 |
| 4 | Gene expression-based biomarkers for discriminating early and late stage of clear cell renal cancer. Scientific Reports, 2017, 7, 44997. | 3.3 | 92 |
| 5 | Prediction and Analysis of Skin Cancer Progression using Genomics Profiles of Patients. Scientific Reports, 2019, 9, 15790. | 3.3 | 57 |
| 6 | Identification of Platform-Independent Diagnostic Biomarker Panel for Hepatocellular Carcinoma Using Large-Scale Transcriptomics Data. Frontiers in Genetics, 2019, 10, 1306. | 2.3 | 49 |
| 7 | Crowdsourcing assessment of maternal blood multi-omics for predicting gestational age and preterm birth. Cell Reports Medicine, 2021, 2, 100323. | 6.5 | 47 |
| 8 | Classification of early and late stage liver hepatocellular carcinoma patients from their genomics and epigenomics profiles. PLoS ONE, 2019, 14, e0221476. | 2.5 | 46 |
| 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 | CancerPDF: A repository of cancer-associated peptidome found in human biofluids. Scientific Reports, 2017, 7, 1511. | 3.3 | 30 |
| 12 | 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 |
| 13 | CancerLivER: a database of liver cancer gene expression resources and biomarkers. Database: the Journal of Biological Databases and Curation, 2020, 2020, . | 3.0 | 21 |
| 14 | 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 |
| 15 | Expression based biomarkers and models to classify early and late-stage samples of Papillary Thyroid Carcinoma. PLoS ONE, 2020, 15, e0231629. | 2.5 | 13 |
| 16 | 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 |
| 17 | Computational resources for identification of cancer biomarkers from omics data. Briefings in Functional Genomics, 2021, 20, 213-222. | 2.7 | 7 |
| 18 | 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 |

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| # | 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 | Ο |