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List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

21

papers

94

citations

5

h-index

9

g-index

47

ext. papers

251

ext. citations

4.6

avg, IF

3.02

L-index

#	Paper	IF	Citations
21	Comparison of Machine-Learning Algorithms for the Prediction of Current Procedural Terminology (CPT) Codes from Pathology Reports.. <i>Journal of Pathology Informatics</i> , 2022 , 13, 3	4.4	1
20	Mixed Effects Machine Learning Models for Colon Cancer Metastasis Prediction using Spatially Localized Immuno-Oncology Markers. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2022 , 27, 175-186	1.3	
19	Topological Feature Extraction and Visualization of Whole Slide Images using Graph Neural Networks. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2021 , 26, 285-296	1.3	3
18	Artificial Intelligence in Anatomic Pathology. <i>Advances in Molecular Pathology</i> , 2021 , 4, 145-171	0.3	0
17	Evaluating the mediating effects of perceived vulnerability to disease in the relation between disgust and contamination-based OCD. <i>Journal of Anxiety Disorders</i> , 2021 , 79, 102384	10.9	0
16	Journey across epidemiology's third variables: an anesthesiologist's guide for successfully navigating confounding, mediation, and effect modification. <i>Regional Anesthesia and Pain Medicine</i> , 2021 , 46, 936-940	3.4	2
15	A large-scale internal validation study of unsupervised virtual trichrome staining technologies on nonalcoholic steatohepatitis liver biopsies. <i>Modern Pathology</i> , 2021 , 34, 808-822	9.8	7
14	Atypia of undetermined significance in thyroid cytology: Nuclear atypia and architectural atypia are associated with different molecular alterations and risks of malignancy. <i>Cancer Cytopathology</i> , 2021 , 129, 966-972	3.9	0
13	Using molecular testing to improve the management of thyroid nodules with indeterminate cytology: an institutional experience with review of molecular alterations. <i>Journal of the American Society of Cytopathology</i> , 2021 ,	2.4	2
12	MethylSPWNet and MethylCapsNet: Biologically Motivated Organization of DNAm Neural Networks, Inspired by Capsule Networks. <i>Npj Systems Biology and Applications</i> , 2021 , 7, 33	5	0
11	DNA 5-hydroxymethylcytosine in pediatric central nervous system tumors may impact tumor classification and is a positive prognostic marker. <i>Clinical Epigenetics</i> , 2021 , 13, 176	7.7	0
10	MethylNet: an automated and modular deep learning approach for DNA methylation analysis. <i>BMC Bioinformatics</i> , 2020 , 21, 108	3.6	23
9	Don't dismiss logistic regression: the case for sensible extraction of interactions in the era of machine learning. <i>BMC Medical Research Methodology</i> , 2020 , 20, 171	4.7	10
8	Does the timing of postoperative showering impact infection rates? A systematic review and meta-analysis. <i>Journal of Plastic, Reconstructive and Aesthetic Surgery</i> , 2020 , 73, 1306-1311	1.7	1
7	PathFlowAI: A High-Throughput Workflow for Preprocessing, Deep Learning and Interpretation in Digital Pathology. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2020 , 25, 403-414	1.3	5
6	Gradual polyploid genome evolution revealed by pan-genomic analysis of <i>Brachypodium hybridum</i> and its diploid progenitors. <i>Nature Communications</i> , 2020 , 11, 3670	17.4	22
5	PyMethylProcess-convenient high-throughput preprocessing workflow for DNA methylation data. <i>Bioinformatics</i> , 2019 , 35, 5379-5381	7.2	6

LIST OF PUBLICATIONS

4	PolyCRACKER, a robust method for the unsupervised partitioning of polyploid subgenomes by signatures of repetitive DNA evolution. <i>BMC Genomics</i> , 2019 , 20, 580	4.5	2
3	Preliminary Evaluation of the Utility of Deep Generative Histopathology Image Translation at a Mid-Sized NCI Cancer Center		3
2	PathFlow-MixMatch for Whole Slide Image Registration: An Investigation of a Segment-Based Scalable Image Registration Method		2
1	Comparison of Machine Learning Algorithms for the Prediction of Current Procedural Terminology (CPT) Codes from Pathology Reports		1