

ilastik: interactive machine learning for (bio)image anal

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Mass Cytometry Imaging for the Study of Human Diseases—Applications and Data Analysis Strategies. <i>Frontiers in Immunology</i> , 2019, 10, 2657.	2.2	139
2	QUINT: Workflow for Quantification and Spatial Analysis of Features in Histological Images From Rodent Brain. <i>Frontiers in Neuroinformatics</i> , 2019, 13, 75.	1.3	51
3	BactMAP: An R package for integrating, analyzing and visualizing bacterial microscopy data. <i>Molecular Microbiology</i> , 2020, 113, 297-308.	1.2	26
4	Mapping Heterogeneous Buried Archaeological Features Using Multisensor Data from Unmanned Aerial Vehicles. <i>Remote Sensing</i> , 2020, 12, 41.	1.8	25
5	Application of automated electron microscopy imaging and machine learning to characterise and quantify nanoparticle dispersion in aqueous media. <i>Journal of Microscopy</i> , 2020, 279, 177-184.	0.8	21
6	Examining the Progressive Behavior and Neuropathological Outcomes Associated with Chronic Repetitive Mild Traumatic Brain Injury in Rats. <i>Cerebral Cortex Communications</i> , 2020, 1, tgaa002.	0.7	6
7	Interactive machine learning for fast and robust cell profiling. <i>PLoS ONE</i> , 2020, 15, e0237972.	1.1	1
8	The use and limitations of single-cell mass cytometry for studying human microglia function. <i>Brain Pathology</i> , 2020, 30, 1178-1191.	2.1	18
9	MitoSegNet: Easy-to-use Deep Learning Segmentation for Analyzing Mitochondrial Morphology. <i>IScience</i> , 2020, 23, 101601.	1.9	44
10	Osteopontin Expression Identifies a Subset of Recruited Macrophages Distinct from Kupffer Cells in the Fatty Liver. <i>Immunity</i> , 2020, 53, 641-657.e14.	6.6	287
11	Amoeboid Swimming Is Propelled by Molecular Paddling in Lymphocytes. <i>Biophysical Journal</i> , 2020, 119, 1157-1177.	0.2	26
12	AnnotatorJ: an ImageJ plugin to ease hand annotation of cellular compartments. <i>Molecular Biology of the Cell</i> , 2020, 31, 2179-2186.	0.9	30
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14	A Plasma Membrane Nanodomain Ensures Signal Specificity during Osmotic Signaling in Plants. <i>Current Biology</i> , 2020, 30, 4654-4664.e4.	1.8	40
15	Large-scale characterization of the microvascular geometry in development and disease by tissue clearing and quantitative ultramicroscopy. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2020, 41, 0271678X2096185.	2.4	10
16	High-content, label-free analysis of proplatelet production from megakaryocytes. <i>Journal of Thrombosis and Haemostasis</i> , 2020, 18, 2701-2711.	1.9	11
17	Apical Relaxation during Mitotic Rounding Promotes Tension-Oriented Cell Division. <i>Developmental Cell</i> , 2020, 55, 695-706.e4.	3.1	20
18	Energy Sources of the Depth-Generalist Mixotrophic Coral <i>Stylophora pistillata</i> . <i>Frontiers in Marine Science</i> , 2020, 7, 988.	1.2	36

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20	Warburg-like Metabolic Reprogramming in Aging Intestinal Stem Cells Contributes to Tissue Hyperplasia. <i>Cell Reports</i> , 2020, 33, 108423.	2.9	36
21	Single Cell Characterization of a Synthetic Bacterial Clock with a Hybrid Feedback Loop Containing dCas9-sgRNA. <i>ACS Synthetic Biology</i> , 2020, 9, 3377-3387.	1.9	13
22	OpSeF: Open Source Python Framework for Collaborative Instance Segmentation of Bioimages. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 558880.	2.0	9
23	FUS is lost from nuclei and gained in neurites of motor neurons in a human stem cell model of VCP-related ALS. <i>Brain</i> , 2020, 143, e103-e103.	3.7	15
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29	Epigenomic State Transitions Characterize Tumor Progression in Mouse Lung Adenocarcinoma. <i>Cancer Cell</i> , 2020, 38, 212-228.e13.	7.7	140
30	BIAFLOWS: A Collaborative Framework to Reproducibly Deploy and Benchmark Bioimage Analysis Workflows. <i>Patterns</i> , 2020, 1, 100040.	3.1	25
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