Baihan Lin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6213986/publications.pdf

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		1478505	1372567	
15	315	6	10	
papers	citations	h-index	g-index	
17	17	17	575	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	Citations
1	Regularity Normalization: Neuroscience-Inspired Unsupervised Attention across Neural Network Layers. Entropy, 2022, 24, 59.	2.2	3
2	Predicting human decision making in psychological tasks with recurrent neural networks. PLoS ONE, 2022, 17, e0267907.	2.5	6
3	Models of Human Behavioral Agents inÂBandits, Contextual Bandits and RL. Communications in Computer and Information Science, 2021, , 14-33.	0.5	7
4	Influence of ocular dominance columns and patchy callosal connections on binocularity in lateral striate cortex: Long Evans versus albino rats. Journal of Comparative Neurology, 2020, 528, 650-663.	1.6	7
5	Online Semi-supervised Learning in Contextual Bandits with Episodic Reward. Lecture Notes in Computer Science, 2020, , 407-419.	1.3	5
6	Keep It Real: a Window to Real Reality in Virtual Reality. , 2020, , .		1
7	Self-Assembling 2D Arrays with <i>de Novo</i> Protein Building Blocks. Journal of the American Chemical Society, 2019, 141, 8891-8895.	13.7	37
8	Split Q Learning: Reinforcement Learning with Two-Stream Rewards. , 2019, , .		9
9	Neural Networks as Model Selection withÂIncremental MDL Normalization. Communications in Computer and Information Science, 2019, , 195-208.	0.5	0
10	Visualizing Representational Dynamics with Multidimensional Scaling Alignment., 2019,,.		2
11	Contextual Bandit with Adaptive Feature Extraction. , 2018, , .		10
12	Model-informed risk assessment for Zika virus outbreaks in the Asia-Pacific regions. Journal of Infection, 2017, 74, 484-491.	3.3	23
13	An Integrative Analysis Reveals a Central Role of P53 Activation via MDM2 in Zika Virus Infection Induced Cell Death. Frontiers in Cellular and Infection Microbiology, 2017, 7, 327.	3.9	23
14	Dynamic Forecasting of Zika Epidemics Using Google Trends. PLoS ONE, 2017, 12, e0165085.	2.5	137
15	Systematic Genome-wide Screening and Prediction of microRNAs in EBOV During the 2014 Ebolavirus Outbreak. Scientific Reports, 2015, 5, 9912.	3.3	44