

Improved prediction of protein-protein interactions features, and an ensemble classifier

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Identifying N6-methyladenosine sites using multi-interval nucleotide pair position specificity and support vector machine. <i>Scientific Reports</i> , 2017, 7, 46757.	1.6	77
2	CPPred-RF: A Sequence-based Predictor for Identifying Cell-Penetrating Peptides and Their Uptake Efficiency. <i>Journal of Proteome Research</i> , 2017, 16, 2044-2053.	1.8	168
3	Integration of data mining classification techniques and ensemble learning to identify risk factors and diagnose ovarian cancer recurrence. <i>Artificial Intelligence in Medicine</i> , 2017, 78, 47-54.	3.8	64
4	iSS-PC: Identifying Splicing Sites via Physical-Chemical Properties Using Deep Sparse Auto-Encoder. <i>Scientific Reports</i> , 2017, 7, 8222.	1.6	21
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7	Drug-Target Interaction Prediction through Label Propagation with Linear Neighborhood Information. <i>Molecules</i> , 2017, 22, 2056.	1.7	68
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