Prediction of Recombination Spots Using Novel Hybrid Learning Approach

Frontiers in Genetics

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Citation Report

#	Article	IF	CITATIONS
1	Epigenetic Marks and Variation of Sequence-Based Information Along Genomic Regions Are Predictive of Recombination Hot/Cold Spots in Saccharomyces cerevisiae. Frontiers in Genetics, 2021, 12, 705038.	1.1	0
2	Recent Advancements in Fruit Detection and Classification Using Deep Learning Techniques. Mathematical Problems in Engineering, 2022, 2022, 1-29.	0.6	58
3	Deep-piRNA: Bi-Layered Prediction Model for PIWI-Interacting RNA Using Discriminative Features. Computers, Materials and Continua, 2022, 72, 2243-2258.	1.5	5
4	Knowledge structure and emerging trends in the application of deep learning in genetics research: A bibliometric analysis [2000–2021]. Frontiers in Genetics, 0, 13, .	1.1	2
5	Information Theory for Biological Sequence Classification: A Novel Feature Extraction Technique Based on Tsallis Entropy. Entropy, 2022, 24, 1398.	1.1	2
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7	Semantic Segmentation of Fish and Underwater Environments Using Deep Convolutional Neural Networks and Learned Active Contours. IEEE Access, 2023, 11, 33652-33665.	2.6	3