A NOVEL PREDICTIVE ANALYSER FOR DATA INTENSIVE APPLICATION

by Vignesh Voleti Samudra

Submission date: 17- Apr- 2019 04:26PM (UT C+0530)

Submission ID: 1114207594

File name: 15BCE_0076_82_93.pdf (4.14M)

Word count: 5623 Character count: 29886

A NOVEL PREDICTIVE ANALYSER FOR DATA INTENSIVE APPLICATION

ORIGINALITY REPORT PUBLICAT IONS ST UDENT PAPERS SIMILARITY INDEX INT ERNET SOURCES PRIMARY SOURCES arxiv.org Internet Source Alexander Rosenberg Johansen, Casper Kaae Sønderby, Søren Kaae Sønderby, Ole Winther. "Deep Recurrent Conditional Random Field Network for Protein Secondary Prediction", Proceedings of the 8th ACM International Conference on Bioinformatics, Computational Biology, and Health Informatics - ACM-BCB '17, 2017 Publicat ion Johansen, Alexander Rosenberg, Jing Jin, 2% Tomasz Maszczyk, Justin Dauwels, Sydney S. Cash, and M. Brandon Westover. "Epileptiform spike detection via convolutional neural networks", 2016 IEEE International Conference on Acoustics Speech and Signal Processing (ICASSP), 2016.

Publicat ion

4	Int ernet Source	1%
5	Int ernet Source	1%
6	www.doc.ic.ac.uk Int ernet Source	1%
7	en.wikipedia.org Int ernet Source	1%
8	S. Muggleton, R.D. King, M.J.E. Sternberg. "Using logic for protein structure prediction", Proceedings of the Twenty-Fifth Hawaii International Conference on System Sciences, 1992 Publicat ion	1%
9	Submitted to Trinity College Dublin St udent Paper	<1%
10	Qian, N "Predicting the secondary structure of globular proteins using neural network models", Journal of Molecular Biology, 19880820 Publicat ion	<1%
11	LEMOINE, E., D. MERCERON, J. SALLANTIN, and E. MEPHU NGUIFO. "IMPROVING THE EFFICIENCY OF A USER-DRIVEN LEARNING	<1%

SYSTEM WITH RECONFIGURABLE

HARDWARE.: APPLICATION TO DNA SPLICING.", Biocomputing 99, 1998.

Publicat ion

12	krishikosh.egranth.ac.in Int ernet Source	<1%
13	amitpednekar971992.blogspot.com Int ernet Source	<1%
14	www.educationaldatamining.org Int ernet Source	<1%
15	mti.binus.ac.id Int ernet Source	<1%
16	Shweta Yadav, Asif Ekbal, Sriparna Saha, Ankit Kumar, Pushpak Bhattacharyya. "Feature assisted stacked attentive shortest dependency path based Bi-LSTM model for protein—protein interaction", Knowledge-Based Systems, 2019 Publicat ion	<1%

Exclude quotes On

Exclude bibliography On

Exclude matches

< 10 words