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International Researcher IDs

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Publons / Web Of Science ResearcherID: M-2256-2016

ScopusID: 6602757969 Yoksis Researcher ID: 48646

Education Information

Doctorate, Universite Paris 5, Ufr Mathematique Et Informatique, France 1990 - 1993

Postgraduate, Middle East Technical University, Faculty of Engineering, Elektrik-Elektronik Mühendisliği Bölümü, Turkey 1988 - 1990

Undergraduate, Middle East Technical University, Faculty of Engineering, Elektrik-Elektronik Mühendisliği Bölümü, Turkey 1983 - 1987

Foreign Languages

English, C2 Mastery French, C1 Advanced

Dissertations

Doctorate, Reseaux de Neurones Aleatoires et Textures, Universite Paris 5, Ufr Mathematique Et Informatique, 1993 Postgraduate, Image Processing of Ottoman Documents, Middle East Technical University, Faculty of Engineering, Elektrik-Elektronik Mühendisliği Bölümü, 1990

Academic Titles / Tasks

Professor, Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 2005 - Continues

Associate Professor, Virginia Polytechnic Institute and State University, Virginia Bioinformatics Institute, 2004 - 2005 Associate Professor, Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 1998 - 2005

Assistant Professor, Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 1993 - 1998

Lecturer, Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 1993 - 1993

Academic and Administrative Experience

Middle East Technical University, Presidency Office, 2010 - 2016

Middle East Technical University, Presidency Office, 2008 - 2010

Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 2007 - 2008

Orta Doğu Teknik Üniversitesi, Faculty of Engineering, Department of Computer Engineering, 2001 - 2004

Middle East Technical University, Faculty of Engineering, Department of Computer Engineering, 1996 - 1997

Advising Theses

ATALAY M. V., Distance matrices as protein representations, Postgraduate, M.DİNÇ(Student), 2022

ATALAY M. V., Implementation of crossbar web service for interactive visualizations of biological networks, Postgraduate, A.Atakan(Student), 2020

Atalay M. V., Multi-view subcellular localization prediction of human proteins, Postgraduate, G.ÖZSARI(Student), 2019

Atalay M. V., Enzyme prediction with word embedding approach, Postgraduate, E.AKIN(Student), 2019

Atalay M. V., A customized force-directed layout algorithm with genetic algorithm techniques for biological graphs whose vertices have enzyme commission attributes, Postgraduate, F.AKSOYDAN(Student), 2019

ATALAY M. V., Deep learning for prediction of drug-target interaction space and protein functions, Doctorate, A.Süreyya(Student), 2019

ATALAY M. V., ATALAY R., Prediction of enzymatic properties of protein sequences based on the enzyme commission nomenclature, Postgraduate, A.DALKIRAN(Student), 2017

ATALAY M. V., A customized force-directed layout algorithm for biological graphs whose vertices have enzyme commission attributes, Postgraduate, H.FEHMİ(Student), 2015

ATALAY M. V., ATALAY R., An extension to GOPred to annotate Swiss-Prot and Trembl sequences for all gene ontology categories and EC numbers, Postgraduate, A.SÜREYYA(Student), 2015

ATALAY M. V., Mining microarray data for biologically important gene sets, Doctorate, G.KIRÇİÇEĞİ(Student), 2012 ATALAY M. V., Network structure based pathway enrichment system to analyze pathway activities, Doctorate, Z.IŞIK(Student), 2011

ATALAY M. V., Automatic reconstruction of photorealistic 3-D building models from satellite and ground-level images, Doctorate, E.SÜMER(Student), 2011

ATALAY M. V., Vision-based hand interface systems in human computer interaction, Doctorate, S.GENÇ(Student), 2010 Atalay M. V., Vision based Hand Interface Systems in Human Computer Interaction, Doctorate, S.Genç(Student), 2010 ATALAY M. V., Automated biological data acquisition and integration using machine learning techniques, Doctorate, L.ÇARKACIOĞLU(Student), 2009

ATALAY M. V., Prediction of enzyme classes in a hierarchical approach by using SPMap, Postgraduate, A.GÜL(Student), 2009

ATALAY M. V., Subsequence feature maps for protein function annotation, Doctorate, Ö.SİNAN(Student), 2008

ATALAY M. V., Hybrid CDN P2P architecture for multimedia streaming, Doctorate, K.ÖZTOPRAK(Student), 2008

ATALAY M. V., A programming framework to implement rule-based target detection in images, Postgraduate, Y.ŞAHİN(Student), 2008

CAN T., ATALAY M. V., A classification system for the problem of protein subcellular localization, Postgraduate, G.ALAY(Student), 2007

ATALAY M. V., Improving interactive classification of satellite image content, Postgraduate, G.TEKKAYA(Student), 2007 ATALAY M. V., A clustering method for the problem of protein subcellular localization, Postgraduate, P.BEZEK(Student), 2006

ATALAY M. V., Interactive classification of satellite image content based on query by example, Postgraduate, O.DALAY(Student), 2006

ATALAY M. V., Image-based extraction of material reflectance properties of a 3D object, Postgraduate, M.ERKUT(Student), 2003

ATALAY M. V., Prediction of protein subcellular localization using global protein sequence feature, Postgraduate,

B.BOZKURT(Student), 2003

ATALAY M. V., Texture classification and retrieval using random neural network model, Postgraduate, A.TEKE(Student), 2003

ATALAY M. V., Reconstruction of a 3D human head model from images, Doctorate, R.ZARE(Student), 2003

ATALAY M. V., Fast object tracking by fusing adaptive background models and mean-shift analysis, Postgraduate, Ö.TÜZEL(Student), 2003

ATALAY M. V., Silhouette-based 3D reconstruction by energy minimization, Postgraduate, E.MENGİ (Student), 2003

ATALAY M. V., Prediction of protein subcellular localization based on primary sequence date, Postgraduate,

M.ÖZARAR(Student), 2003

ATALAY M. V., Vision-based human-computer interaction using laser pointer, Postgraduate, İ.AYKUT(Student), 2003

ATALAY M. V., Shortest path analysis in dynamic transportation networks by expert systems: A case study in Ankara, Bahçelievler district by using genetic algorithms, Postgraduate, G.EKE(Student), 2003

ATALAY M. V., Prediction of protein subcellular localization based on primary sequence data, Postgraduate, M.Özarar(Student), 2003

ATALAY M. V., Towards finding optimal mixture of subspaces for data classification, Doctorate, M.ELHAFIZ(Student), 2003

ATALAY M. V., Texture discrimination and segmentation of remotely sensed imagery by using adaptive subband decomposition, Postgraduate, M.YAMAN(Student), 2003

ATALAY M. V., Comparison of photo-consistency measured used in the voxel coloring algorithm, Postgraduate, O.ÖZÜN(Student), 2002

Atalay V., Relative consistency of projective reconstructions obtained from an amage pair., Postgraduate, B.Otlu(Student), 2002

ATALAY M. V., Cognitive and computational aspects of gender estimation from faces, Postgraduate, M.KORAY(Student), 2002

Atalay V., A hybrid approach for Turkish optical character recognition., Postgraduate, G.Kırçiçegi(Student), 2002

ATALAY M. V., Appearance reconstruction of three dimensional models from real images., Postgraduate,

U.YILMAZ(Student), 2002

HALICI U., ATALAY M. V., 3 D reconstruction of rigid objects from multiple calibrated views, Doctorate,

A.YAŞAR(Student), 2002

ATALAY M. V., A Hybrid approach for Turkish optical character recognition, Postgraduate, G.KIRÇİÇEĞİ(Student), 2002 ATALAY M. V., Contextual post processing for Turkish optical character recognition., Postgraduate, S.ULAŞ(Student), 2002

Atalay V., Comparison of photo-consistency measures used in the voxel coloring algorithm., Postgraduate, O.Özün(Student), 2002

ATALAY M. V., Relative consistency of projective reconstructions obtained from an image pair, Postgraduate, B.OTLU(Student), 2002

ATALAY M. V., Image mosaicing, Postgraduate, A.ÖZZEYBEK(Student), 2000

ATALAY M. V., Segmentation of human face, Postgraduate, S.BASKAN(Student), 2000

Atalay V., Texture extraction and reconsturuction from multiple images, Postgraduate, S.Genç(Student), 1999

ATALAY M. V., Texture extraction and reconstruction from multiple images, Postgraduate, S.GENÇ(Student), 1999

ATALAY M. V., An SGML viewer for form document images, Postgraduate, E.ARSLAN(Student), 1998

ATALAY M. V., A Form document image parser, Postgraduate, E.DİNCEL(Student), 1998

ATALAY M. V., Form document identification based on hierarchical tree representation, Postgraduate, P.DUYGULU(Student), 1998

ATALAY M. V., Two matching algorithms for derial images, Postgraduate, D.ÇULHA(Student), 1998

Atalay V., Two matching algorithms for aerial images., Postgraduate, D.Çulha(Student), 1998

ATALAY M. V., A New method for road finding in aerial images based on self organization and perceptual grouping principles, Postgraduate, A.YAŞAR(Student), 1996

ATALAY M. V., A Matching algorithm based linear features, Postgraduate, M.UĞUR(Student), 1996

Atalay V., A matching algorithm based on linear features, Postgraduate, M.Uğur(Student), 1996

Jury Memberships

Award, TÜSEB Teşvik Ödülü, Middle East Technical University, October, 2020 Award, TÜSEB Teşvik Ödülü, Middle East Technical University, October, 2020

Published journal articles indexed by SCI, SSCI, and AHCI

I. Online embedding and clustering of evolving data streams

Zubaroglu A., Atalay M. V.

STATISTICAL ANALYSIS AND DATA MINING, vol.16, no.1, pp.29-44, 2023 (SCI-Expanded)

II. POWER CONSUMPTION FORECASTING BY HYBRID DEEP ARCHITECTURES WITH DATA FUSION Özen S., ATALAY M. V.

Computing and Informatics, vol.42, no.1, pp.126-156, 2023 (SCI-Expanded)

III. SLPred: a multi-view subcellular localization prediction tool for multi-location human proteins
 ÖZSARI G., Rifaioglu A. S., ATAKAN A., DOĞAN T., Martin M. J., Atalay R. C., ATALAY M. V.
 BIOINFORMATICS, vol.38, no.17, pp.4226-4229, 2022 (SCI-Expanded)

IV. CROssBAR: comprehensive resource of biomedical relations with knowledge graph representations.

DOĞAN T., Atas H., Joshi V., ATAKAN A., Rifaioglu A. S., NALBAT E., Nightingale A., Saidi R., Volynkin V., Zellner H., et

Nucleic acids research, vol.49, no.16, 2021 (SCI-Expanded)

V. Crowdsourced mapping of unexplored target space of kinase inhibitors

Cichonska A., Ravikumar B., Allaway R. J., Wan F., Park S., Isayev O., Li S., Mason M., Lamb A., Tanoli Z., et al. NATURE COMMUNICATIONS, vol.12, no.1, 2021 (SCI-Expanded)

VI. MDeePred: Novel multi-channel protein featurization for deep learning-based binding affinity prediction in drug discovery

Rifaioglu A., Atalay R. C., KAHRAMAN D. C., DOĞAN T., Martin M., ATALAY M. V.

Bioinformatics, vol.37, no.5, pp.693-704, 2021 (SCI-Expanded)

VII. Data stream clustering: a review

Zubaroglu A., Atalay V.

ARTIFICIAL INTELLIGENCE REVIEW, vol.54, no.2, pp.1201-1236, 2021 (SCI-Expanded)

VIII. iBioProVis: interactive visualization and analysis of compound bioactivity space

Dönmez A., Rifaioğlu A. S., Acar A. C., Doğan T., Cetin-Atalay R., Atalay V.

BIOINFORMATICS, vol.36, no.14, pp.4227-4230, 2020 (SCI-Expanded)

IX. DEEPScreen: high performance drug-target interaction prediction with convolutional neural networks using 2-D structural compound representations

RİFAİOĞLU A. S., NALBAT E., Atalay V., Martin M. J., Cetin-Atalay R., DOĞAN T.

CHEMICAL SCIENCE, vol.11, no.9, pp.2531-2557, 2020 (SCI-Expanded)

X. The CAFA challenge reports improved protein function prediction and new functional annotations for hundreds of genes through experimental screens

Zhou N., Jiang Y., Bergquist T., Lee A., Kacsoh B., Crocker A., Lewis K., Georghiou G., Nguyen H., Hamid M., et al. GENOME BIOLOGY, vol.20, no.1, 2019 (SCI-Expanded)

XI. DEEPred: Automated Protein Function Prediction with Multi-task Feed-forward Deep Neural Networks

RİFAİOĞLU A. S., Dogan T., Martin M. J., Cetin-Atalay R., Atalay V.

SCIENTIFIC REPORTS, vol.9, 2019 (SCI-Expanded)

XII. ECPred: a tool for the prediction of the enzymatic functions of protein sequences based on the EC nomenclature

Dalkıran A., Rifaioğlu A. S., Martin M. J., Cetin-Atalay R., Atalay M. V., Dogan T.

BMC BIOINFORMATICS, vol.19, 2018 (SCI-Expanded)

XIII. EClerize: A customized force-directed graph drawing algorithm for biological graphs with EC

attributes

Danaci H. F., Cetin-Atalay R., ATALAY M. V.

JOURNAL OF BIOINFORMATICS AND COMPUTATIONAL BIOLOGY, vol.16, no.4, 2018 (SCI-Expanded)

XIV. Recent applications of deep learning and machine intelligence on in silico drug discovery: methods, tools and databases.

Rifaioglu A. S., Atas H., Martin M., Cetin-Atalay R., Atalay V., Dogan T.

Briefings in bioinformatics, 2018 (SCI-Expanded)

XV. Large-scale automated function prediction of protein sequences and an experimental case study validation on PTEN transcript variants

RİFAİOĞLU A. S., Dogan T., Sarac O. S., ERSAHİN T., Saidi R., ATALAY M. V., MARTİN M. J., Cetin-Atalay R. PROTEINS-STRUCTURE FUNCTION AND BIOINFORMATICS, vol.86, no.2, pp.135-151, 2018 (SCI-Expanded)

XVI. HandVR: a hand-gesture-based interface to a video retrieval system

Genc S., Bastan M., GÜDÜKBAY U., Atalay V., GÜDÜKBAY U.

SIGNAL IMAGE AND VIDEO PROCESSING, vol.9, no.7, pp.1717-1726, 2015 (SCI-Expanded)

XVII. Identification of Novel Reference Genes Based on MeSH Categories

Ersahin T., ÇARKACIOĞLU L., CAN T., Konu O., Atalay V., Cetin-Atalay R.

PLOS ONE, vol.9, no.3, 2014 (SCI-Expanded)

XVIII. A signal transduction score flow algorithm for cyclic cellular pathway analysis, which combines transcriptome and ChIP-seq data

Isik Z., Ersahin T., Atalay V., AYKANAT C., Cetin-Atalay R.

MOLECULAR BIOSYSTEMS, vol.8, no.12, pp.3224-3231, 2012 (SCI-Expanded)

XIX. GOPred: GO Molecular Function Prediction by Combined Classifiers

Sarac O. S., Atalay V., Cetin-Atalay R.

PLOS ONE, vol.5, no.8, 2010 (SCI-Expanded)

XX. Prediction of enzyme classes in a hierarchical approach by using SPMap

Yaman A., Atalay V., Atalay R.

NEW BIOTECHNOLOGY, vol.27, 2010 (SCI-Expanded)

XXI. Bi-k-bi clustering: mining large scale gene expression data using two-level biclustering

Carkacioglu L., Atalay R. C., KONU KARAKAYALI Ö., Atalay V., CAN T.

INTERNATIONAL JOURNAL OF DATA MINING AND BIOINFORMATICS, vol.4, no.6, pp.701-721, 2010 (SCI-Expanded)

XXII. Subsequence-based feature map for protein function classification

Sarac O. S., Guersoy-Yuezueguellue O., Cetin-Atalay R., Atalay V.

COMPUTATIONAL BIOLOGY AND CHEMISTRY, vol.32, no.2, pp.122-130, 2008 (SCI-Expanded)

XXIII. Experimental study on the sensitivity of autocalibration to projective camera model parameters Hassanpour R., Atalay V.

OPTICAL ENGINEERING, vol.45, no.4, 2006 (SCI-Expanded)

XXIV. Implicit motif distribution based hybrid computational kernel for sequence classification Atalay V., Cetin-Atalay R.

BIOINFORMATICS, vol.21, no.8, pp.1429-1436, 2005 (SCI-Expanded)

XXV. Almost autonomous training of mixtures of principal component analyzers

Musa M., de Ridder D., Duin R., Atalay V.

PATTERN RECOGNITION LETTERS, vol.25, no.9, pp.1085-1095, 2004 (SCI-Expanded)

XXVI. Camera auto-calibration using a sequence of 2D images with small rotations

Hassanpour R., Atalay V.

PATTERN RECOGNITION LETTERS, vol.25, no.9, pp.989-997, 2004 (SCI-Expanded)

XXVII. Vision-based continuous Graffiti (TM)-like text entry system

Erdem I., Erdem M., Atalay V., Cetin A.

OPTICAL ENGINEERING, vol.43, no.3, pp.553-558, 2004 (SCI-Expanded)

XXVIII. A hierarchical representation of form documents for identification and retrieval

Duygulu P., Atalay V.

International Journal on Document Analysis and Recognition, vol.5, no.1, pp.17-27, 2003 (SCI-Expanded)

XXIX. Silhouette-based 3-D model reconstruction from multiple images

Mulayim A., Yilmaz U., Atalay V.

IEEE TRANSACTIONS ON SYSTEMS MAN AND CYBERNETICS PART B-CYBERNETICS, vol.33, no.4, pp.582-591, 2003 (SCI-Expanded)

XXX. Ore-age: a hybrid system for assisting and teaching mining method selection

Guray C., Celebi N., Atalay V., Pasamehmetoglu A.

EXPERT SYSTEMS WITH APPLICATIONS, vol.24, no.3, pp.261-271, 2003 (SCI-Expanded)

XXXI. Prediction of protein subcellular localization based on primary sequence data

Ozarar M., Atalay V., Atalay R.

COMPUTER AND INFORMATION SCIENCES - ISCIS 2003, vol.2869, pp.611-618, 2003 (SCI-Expanded)

XXXII. Texture segmentation using the mixtures of principal component analyzers

Musa M., Duin R., de Ridder D., Atalay V.

COMPUTER AND INFORMATION SCIENCES - ISCIS 2003, vol.2869, pp.505-512, 2003 (SCI-Expanded)

XXXIII. Projection based method for segmentation of human face and its evaluation

Baskan S., Bulut M., Atalay V.

PATTERN RECOGNITION LETTERS, vol.23, no.14, pp.1623-1629, 2002 (SCI-Expanded)

XXXIV. Vision-based single-stroke character recognition for wearable computing

Ozer O., Ozun O., Tuzel C., Atalay V., Cetin A.

IEEE INTELLIGENT SYSTEMS & THEIR APPLICATIONS, vol.16, no.3, pp.33-37, 2001 (SCI-Expanded)

XXXV. Matching algorithm based on Godel coding scheme

Dicle N., Atalay V.

Proceedings of SPIE - The International Society for Optical Engineering, vol.3974, pp.576-583, 2000 (SCI-Expanded)

XXXVI. Subband domain coding of binary textual images for document archiving

Gerek O., Cetin A., Tewfik A., Atalay V.

IEEE TRANSACTIONS ON IMAGE PROCESSING, vol.8, no.10, pp.1438-1446, 1999 (SCI-Expanded)

XXXVII. Repulsive attractive network for baseline extraction on document images

Oztop E., Mulayim A., Atalay V., Yarman-Vural F.

SIGNAL PROCESSING, vol.75, no.1, pp.1-10, 1999 (SCI-Expanded)

XXXVIII. A matching algorithm based on linear features

Atalay V., Yilmaz M.

PATTERN RECOGNITION LETTERS, vol.19, no.9, pp.857-867, 1998 (SCI-Expanded)

Articles Published in Other Journals

I. Data Centric Molecular Analysis and Evaluation of Hepatocellular Carcinoma Therapeutics Using Machine Intelligence-Based Tools

Cetin-Atalay R., Kahraman D. C., Nalbat E., Rifaioğlu A. S., Atakan A., Dönmez A., Atas H., Atalay M. V., Acar A. C., Doğan T.

JOURNAL OF GASTROINTESTINAL CANCER, vol.52, no.4, pp.1266-1276, 2021 (ESCI)

II. Multi-task Deep Neural Networks in Protein Function Prediction

RİFAİOĞLU A. S., doğan t., Martin M. J., ATALAY R., ATALAY M. V.

arXiv, 2017 (Non Peer-Reviewed Journal)

III. An Image-Based Inexpensive 3D Scanner

Yilmaz U., Mülayim A. Y., Atalay V.

International Journal of Image and Graphics, vol.3, no.2, pp.235-263, 2003 (ESCI)

IV. Computer vision based mouse

Erdem A., Erdem E., Yardimci Y., Atalay V., Cetin A. E.

ICASSP, IEEE International Conference on Acoustics, Speech and Signal Processing - Proceedings, vol.4, pp.4178,

Books & Book Chapters

I. Forecasting of Product Quality Through Anomaly Detection

Dinc M., Ertekin Bolelli Ş., Ozkan H., Meydanli C., Atalay M. V.

in: Machine Learning and Knowledge Discovery in Databases, Cellier P., Driessens K., Editor, Springer-Verlag, Basel, pp.357-366, 2020

Refereed Congress / Symposium Publications in Proceedings

I. Novel isoxazole- piperazine derivatives inhibit the stemness of HCC cells

NALBAT E., İBİŞ K., ÇALIŞKAN B., BANOĞLU E., ATALAY M. V., ATALAY R.

AASLD-TASL Digital Hepatology Meeting, Ankara, Turkey, 15 January 2022, pp.5-6

II. In vitro validation of drug-target interactions revealed in silico by Comprehensive Resource of Biomedical Relations with Network Representations and Deep Learning (CROssBAR) in HCC

NALBAT E., RİFAİOĞLU A. S., DOĞAN T., Martin M. J., Cetin-Atalay R., ATALAY M. V.

AACR Annual Meeting, ELECTR NETWORK, 22 - 24 June 2020, vol.80

III. Comparison of predictive models for forecasting timeseries data

Özen S., Atalay V., YAZICI A.

3rd International Conference on Big Data Research, ICBDR 2019, Cergy-Pontoise, France, 20 - 21 November 2019, pp.172-176

IV. Online embedding and clustering of data streams

Zubaroğlu A., Atalay V.

3rd International Conference on Big Data Research, ICBDR 2019, Cergy-Pontoise, France, 20 - 21 November 2019, pp.142-146

V. Forecasting of Product Quality Through Anomaly DetectionMehmet Dinç,

Dinc M., ERTEKİN BOLELLİ Ş., Ozkan H., Meydanli C., ATALAY M. V.

2nd International Conference on Digital Transformation and Smart Systems, Ankara, Turkey, 23 - 25 October 2019

VI. Receptor-Ligand Binding Affinity Prediction via Multi-Channel Deep Chemogenomic Modeling

Atalay M. V., Rifaioğlu A. S., Atalay R., Doğan T., Martin M.

The International Symposium on Health Informatics and Bioinformatics (HIBIT 2019), İzmir, Turkey, 17 - 18 October 2019

VII. Forecasting of Product Quality Through Anomaly Detection

Dinc M., ERTEKİN BOLELLİ Ş., Ozkan H., Meydanli C., ATALAY M. V.

ECML/PKDD 2019 Workshop on Io Stream for Data Driven Predictive Maintenance, Würzburg, Germany, 16 September 2019

VIII. iBioProVis: Interactive Visualization and Analysis of Compound Bioactivity Space

Dönmez A., RİFAİOĞLU A. S., ACAR A. C., DOĞAN T., Martin M. J., ATALAY R., ATALAY M. V.

27th Conference on Intellient Systems for Molecular Biology and 18th European Conference on Computational Biology, Basel, Switzerland, 21 - 25 July 2019

IX. CROssBAR: Comprehensive Resource of Biomedical Relations with Network Representations and Deep Learning", Oral presentation

Atalay M. V., Doğan T., Atalay R., Nalbat E., Rifaioğlu A. S., Ataş H., Martin M.

27th Conference on Intelligent Systems for Molecular Biology and 18th European Conference on Computational Biology, Basel, Switzerland, 21 - 25 July 2019

X. Drug-Target Interaction Prediction with Deep Convolutional Neural Networks Using Compound Images

RİFAİOĞLU A. S., ATALAY M. V., Martin M. J., ATALAY R., Doğan T.

26th Conference on Intelligent Systems for Molecular Biology, Chicago, United States Of America, 6 - 10 July 2018

XI. Automated Negative Gene Ontology Based Functional Predictions for Proteins with UniGOPred Doğan T., RİFAİOĞLU A. S., Saidi R., Martin M. J., ATALAY M. V., ATALAY R.

26th Conference on Intelligent Systems for Molecular Biology, Chicago, United States Of America, 6 - 10 July 2018

XII. Investigation of Multi-task Deep Neural Networks in Automated Protein Function Prediction RİFAİOĞLU A. S., Martin M. J., ATALAY R., ATALAY M. V., Doğan T.

ISMB/ECCB 2017: 25th Annual International Conference on Intelligent Systems for Molecular Biology, Prag, Czech Republic, 21 - 25 July 2017

XIII. Prediction of Enzymatic Properties of Protein Sequences Based on the EC Nomenclature

DALKIRAN A., Doğan T., RİFAİOĞLU A. S., ATALAY M. V., Martin M. J., ATALAY R.

ISMB /FCCR 2017: 25th Appeal International Conference on Intelligent Systems for Mal

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XIV. Prediction of Enzymatic Properties of Protein Sequences Based on the EC Nomenclature DALKIRAN A., RİFAİOĞLU A. S., Doğan T., ATALAY M. V., Martin M. J., ATALAY R.

10th The International Symposium on Health Informatics and Bioinformatics (HIBIT 2017), Kalkanlı, Cyprus (Kktc), 28 - 30 June 2017, pp.37

XV. Investigation of Multitask Deep Neural Networks for Protein Function Prediction

RİFAİOĞLU A. S., Martin M. J., ATALAY R., ATALAY M. V., Doğan T.

10th The International Symposium on Health Informatics and Bioinformatics (HIBIT 2017), Erdemli, Cyprus (Kktc), 28 - 30 June 2017

XVI. EClerize: A Customized Force-Directed Layout Algorithm for Biological Networks with EC Attributes
Danacı H. F., ATALAY R., ATALAY M. V.

GLBIO 2017: Great Lakes Bioinformatics Conference, Şikago, United States Of America, 15 - 17 May 2017

XVII. UniGOPred: A Large Scale Automated GO Term Annotation System for UniProtKB
RİFAİOĞLU A. S., Doğan T., SARAÇ Ö. S., Saidi R., ATALAY M. V., Martin M. J., ATALAY R.
GLBIO 2017: Great Lakes Bioinformatics Conference, Şikago, United States Of America, 15 - 17 May 2017

XVIII. ECPred: An Automated Enzymatic Function Prediction Tool Based on Combination of Different Classifiers

DALKIRAN A., RİFAİOĞLU A. S., Doğan T., ATALAY M. V., Martin M. J., ATALAY R.

GLBIO 2017: Great Lakes Bioinformatics Conference, May 15-17, Şikago, United States Of America, 15 - 17 May 2017

XIX. UniGOPred and ECPred Automated Function Prediction Tools Based on A Combination of Different Classifiers

Rifaioglu A., Dogan T., SARAÇ Ö. S., ATALAY M. V., Martin M., ATALAY R.

ISMB Automated FunctionPrediction-SIG, 11 July 2015

XX. Discriminative Modeling of Cell Signaling as Bayesian Networks

Yaman A. G., ACAR A. F., Atalay V., Cetin-Atalay R.

2nd International Work-Conference on Bioinformatics and Biomedical Engineering (IWBBIO), Granada, Nicaragua, 7 - 09 April 2014, pp.1617

XXI. Genetic variants within endotoxin signaling pathway association with asthma and atopy phenotypes ŞAHİNER Ü. M., YAVUZ S., Buyuktiryaki B., BİRBEN E., Topaloglu E., TUNCER A., Yaman A., Atalay R., Atalay V., Sackesen C., et al.

31st Congress of the European-Academy-of-Allergy-and-Clinical-Immunology (EAACI), Geneva, Switzerland, 16 - 20 June 2012, vol.67, pp.469

XXII. Association of asthma with polymorphisms in the endotoxin signaling pathway

ŞAHİNER Ü. M., YAVUZ S., Buyuktiryaki B., BİRBEN E., Topaloglu E., TUNCER A., Yaman A., Atalay R., Atalay V., Sackesen C., et al.

31st Congress of the European-Academy-of-Allergy-and-Clinical-Immunology (EAACI), Geneva, Switzerland, 16 - 20 June 2012, vol.67, pp.469-470

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Non Academic Experience

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