Volkan Atalay

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1365118/publications.pdf

Version: 2024-02-01

| | | 567281 | 3 | 345221 | |
|----------|----------------|--------------|---|----------------|--|
| 53 | 1,556 | 15 | | 36 | |
| papers | citations | h-index | | g-index | |
| | | | | | |
| | | | ľ | | |
| 57 | 57 | 57 | | 1631 | |
| 37 | 37 | 37 | | 1031 | |
| all docs | docs citations | times ranked | | citing authors | |
| | | | | | |

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | SLPred: a multi-view subcellular localization prediction tool for multi-location human proteins. Bioinformatics, 2022, 38, 4226-4229. | 4.1 | 3 |
| 2 | Data stream clustering: a review. Artificial Intelligence Review, 2021, 54, 1201-1236. | 15.7 | 65 |
| 3 | Crowdsourced mapping of unexplored target space of kinase inhibitors. Nature Communications, 2021, 12, 3307. | 12.8 | 41 |
| 4 | CROssBAR: comprehensive resource of biomedical relations with knowledge graph representations. Nucleic Acids Research, 2021, 49, e96-e96. | 14.5 | 19 |
| 5 | DEEPScreen: high performance drug–target interaction prediction with convolutional neural networks using 2-D structural compound representations. Chemical Science, 2020, 11, 2531-2557. | 7.4 | 131 |
| 6 | iBioProVis: interactive visualization and analysis of compound bioactivity space. Bioinformatics, 2020, 36, 4227-4230. | 4.1 | 7 |
| 7 | Forecasting of Product Quality Through Anomaly Detection. Communications in Computer and Information Science, 2020, , 357-366. | 0.5 | O |
| 8 | Recent applications of deep learning and machine intelligence on in silico drug discovery: methods, tools and databases. Briefings in Bioinformatics, 2019, 20, 1878-1912. | 6.5 | 310 |
| 9 | DEEPred: Automated Protein Function Prediction with Multi-task Feed-forward Deep Neural Networks. Scientific Reports, 2019, 9, 7344. | 3.3 | 80 |
| 10 | The CAFA challenge reports improved protein function prediction and new functional annotations for hundreds of genes through experimental screens. Genome Biology, 2019, 20, 244. | 8.8 | 261 |
| 11 | Online Embedding and Clustering of Data Streams. , 2019, , . | | 2 |
| 12 | Comparison of Predictive Models for Forecasting Time-series Data. , 2019, , . | | 1 |
| 13 | EClerize: A customized force-directed graph drawing algorithm for biological graphs with EC attributes. Journal of Bioinformatics and Computational Biology, 2018, 16, 1850007. | 0.8 | 1 |
| 14 | Largeâ€scale automated function prediction of protein sequences and an experimental case study validation on PTEN transcript variants. Proteins: Structure, Function and Bioinformatics, 2018, 86, 135-151. | 2.6 | 13 |
| 15 | ECPred: a tool for the prediction of the enzymatic functions of protein sequences based on the EC nomenclature. BMC Bioinformatics, 2018, 19, 334. | 2.6 | 99 |
| 16 | HandVR: a hand-gesture-based interface to a video retrieval system. Signal, Image and Video Processing, 2015, 9, 1717-1726. | 2.7 | 5 |
| 17 | Identification of Novel Reference Genes Based on MeSH Categories. PLoS ONE, 2014, 9, e93341. | 2.5 | 12 |
| 18 | A signal transduction score flow algorithm for cyclic cellular pathway analysis, which combines transcriptome and ChIP-seq data. Molecular BioSystems, 2012, 8, 3224. | 2.9 | 11 |

| # | Article | IF | Citations |
|----|--|-----|-----------|
| 19 | Bi-k-bi clustering: mining large scale gene expression data using two-level biclustering. International Journal of Data Mining and Bioinformatics, 2010, 4, 701. | 0.1 | 5 |
| 20 | GOPred: GO Molecular Function Prediction by Combined Classifiers. PLoS ONE, 2010, 5, e12382. | 2.5 | 22 |
| 21 | Which Shape Representation Is the Best for Real-Time Hand Interface System?. Lecture Notes in Computer Science, 2009, , 1-11. | 1.3 | 2 |
| 22 | Subsequence-based feature map for protein function classification. Computational Biology and Chemistry, 2008, 32, 122-130. | 2.3 | 19 |
| 23 | Short time series microarray data analysis and biological annotation. , 2008, , . | | 0 |
| 24 | Texture Classification and Retrieval Using the Random Neural Network Model. Computational Management Science, 2006, 3, 193-205. | 1.3 | 11 |
| 25 | Experimental study on the sensitivity of autocalibration to projective camera model parameters. Optical Engineering, 2006, 45, 047002. | 1.0 | 1 |
| 26 | Implicit motif distribution based hybrid computational kernel for sequence classification. Bioinformatics, 2005, 21, 1429-1436. | 4.1 | 15 |
| 27 | Vision-based continuous Graffitiâ,,¢-like text entry system. Optical Engineering, 2004, 43, 553. | 1.0 | 1 |
| 28 | Camera auto-calibration using a sequence of 2D images with small rotations. Pattern Recognition Letters, 2004, 25, 989-997. | 4.2 | 8 |
| 29 | Almost autonomous training of mixtures of principal component analyzers. Pattern Recognition Letters, 2004, 25, 1085-1095. | 4.2 | 6 |
| 30 | Ore-age: a hybrid system for assisting and teaching mining method selection. Expert Systems With Applications, 2003, 24, 261-271. | 7.6 | 16 |
| 31 | Silhouette-based 3-D model reconstruction from multiple images. IEEE Transactions on Systems, Man, and Cybernetics, 2003, 33, 582-591. | 5.0 | 87 |
| 32 | AN IMAGE-BASED INEXPENSIVE 3D SCANNER. International Journal of Image and Graphics, 2003, 03, 235-263. | 1.5 | 2 |
| 33 | Computer vision based unistroke keyboard system and mouse for the handicapped., 2003,,. | | 7 |
| 34 | Texture Segmentation Using the Mixtures of Principal Component Analyzers. Lecture Notes in Computer Science, 2003, , 505-512. | 1.3 | 3 |
| 35 | Prediction of Protein Subcellular Localization Based on Primary Sequence Data. Lecture Notes in Computer Science, 2003, , 611-618. | 1.3 | 0 |
| 36 | Computer vision based mouse. , 2002, , . | | 10 |

| # | Article | IF | Citations |
|----|---|-----|-----------|
| 37 | A hierarchical representation of form documents for identification and retrieval. International Journal on Document Analysis and Recognition, 2002, 5, 17-27. | 3.4 | 31 |
| 38 | Projection based method for segmentation of human face and its evaluation. Pattern Recognition Letters, 2002, 23, 1623-1629. | 4.2 | 80 |
| 39 | Vision-based single-stroke character recognition for wearable computing. IEEE Intelligent Systems, 2001, 16, 33-37. | 4.0 | 15 |
| 40 | <title>Segmentation of human face using gradient-based approach</title> ., 2001, , . | | 1 |
| 41 | Automated contour detection in blood cell images by an efficient snake algorithm. Nonlinear Analysis: Theory, Methods & Applications, 2001, 47, 5839-5847. | 1.1 | 12 |
| 42 | $$ $$ $$ $$ $$ $$ $$ $$ $$ | | 1 |
| 43 | <title>Matching algorithm based on Godel coding scheme</title> ., 2000, 3974, 576. | | 0 |
| 44 | Repulsive attractive network for baseline extraction on document images. Signal Processing, 1999, 75, 1-10. | 3.7 | 26 |
| 45 | Subband domain coding of binary textual images for document archiving. IEEE Transactions on Image Processing, 1999, 8, 1438-1446. | 9.8 | 10 |
| 46 | A matching algorithm based on linear features. Pattern Recognition Letters, 1998, 19, 857-867. | 4.2 | 6 |
| 47 | <title>Estimation of rotation between two frames of a scene</title> ., 1998, 3387, 252. | | 0 |
| 48 | THE RANDOM NEURAL NETWORK MODEL FOR TEXTURE GENERATION. International Journal of Pattern Recognition and Artificial Intelligence, 1992, 06, 131-141. | 1.2 | 63 |
| 49 | Reconstruction of three dimensional models from real images. , 0, , . | | 10 |
| 50 | A character recognizer for Turkish language. , 0, , . | | 3 |
| 51 | Change detection in aerial images. , 0, , . | | 3 |
| 52 | A Novel Model-based Method for Feature Extraction from Protein Sequences for Classification. , 0, , . | | 0 |
| 53 | Online embedding and clustering of evolving data streams. Statistical Analysis and Data Mining, 0, , . | 2.8 | 2 |