## Ulisses Braga-Neto

List of Publications by Year in descending order

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Version: 2024-02-01

471509 526287 1,039 38 17 27 citations h-index g-index papers 39 39 39 1041 docs citations times ranked citing authors all docs

| #  | Article  | IF           | Citations |
|----|--|--------------|-----------|
| 1  | Bolstered error estimation. Pattern Recognition, 2004, 37, 1267-1281.  | 8.1          | 108       |
| 2  | Alternative Complement Pathway Deregulation Is Correlated with Dengue Severity. PLoS ONE, 2009, 4, e6782.  | 2.5          | 95        |
| 3  | Gene Expression Profiling during Early Acute Febrile Stage of Dengue Infection Can Predict the Disease<br>Outcome. PLoS ONE, 2009, 4, e7892.   | 2.5          | 77        |
| 4  | A Theoretical Tour of Connectivity in Image Processing and Analysis. Journal of Mathematical Imaging and Vision, $2003,19,5\text{-}31.$  | 1.3          | 58        |
| 5  | Connectivity on Complete Lattices: New Results. Computer Vision and Image Understanding, 2002, 85, 22-53.  | 4.7          | 57        |
| 6  | Fads and fallacies in the name of small-sample microarray classification - A highlight of misunderstanding and erroneous usage in the applications of genomic signal processing. IEEE Signal Processing Magazine, 2007, 24, 91-99. | 5 <b>.</b> 6 | 54        |
| 7  | Exact performance of error estimators for discrete classifiers. Pattern Recognition, 2005, 38, 1799-1814.  | 8.1          | 51        |
| 8  | Severe Dengue Prognosis Using Human Genome Data and Machine Learning. IEEE Transactions on Biomedical Engineering, 2019, 66, 2861-2868.  | 4.2          | 50        |
| 9  | Superior feature-set ranking for small samples using bolstered error estimation. Bioinformatics, 2005, 21, 1046-1054.  | 4.1          | 49        |
| 10 | EPISTEMOLOGY OF COMPUTATIONAL BIOLOGY: MATHEMATICAL MODELS AND EXPERIMENTAL PREDICTION AS THE BASIS OF THEIR VALIDITY. Journal of Biological Systems, 2006, 14, 65-90.   | 1.4          | 46        |
| 11 | Reliable Classifier to Differentiate Primary and Secondary Acute Dengue Infection Based on IgG ELISA.<br>PLoS ONE, 2009, 4, e4945.   | 2.5          | 42        |
| 12 | Boolean Kalman filter and smoother under model uncertainty. Automatica, 2020, 111, 108609.   | 5 <b>.</b> 0 | 40        |
| 13 | Grayscale Level Connectivity: Theory and Applications. IEEE Transactions on Image Processing, 2004, 13, 1567-1580.   | 9.8          | 32        |
| 14 | A multiscale approach to connectivity. Computer Vision and Image Understanding, 2003, 89, 70-107.  | 4.7          | 27        |
| 15 | Optimal state estimation for Boolean dynamical systems. , 2011, , .  |              | 26        |
| 16 | Confidence Intervals for the True Classification Error Conditioned on the Estimated Error.<br>Technology in Cancer Research and Treatment, 2006, 5, 579-589.   | 1.9          | 21        |
| 17 | Complement factor H gene (CFH) polymorphisms C-257T, G257A and haplotypes are associated with protection against severe dengue phenotype, possible related with high CFH expression. Human Immunology, 2013, 74, 1225-1230.        | 2.4          | 21        |
| 18 | Exact representation of the second-order moments for resubstitution and leave-one-out error estimation for linear discriminant analysis in the univariate heteroskedastic Gaussian model. Pattern Recognition, 2012, 45, 908-917.  | 8.1          | 20        |

| #  | Article  | IF   | Citations |
|----|--|------|-----------|
| 19 | Object-based image analysis using multiscale connectivity. IEEE Transactions on Pattern Analysis and Machine Intelligence, 2005, 27, 892-907.                              | 13.9 | 19        |
| 20 | Fundamentals of Pattern Recognition and Machine Learning. , 2020, , .  |      | 18        |
| 21 | Multiscale Connected Operators. Journal of Mathematical Imaging and Vision, 2005, 22, 199-216.   | 1.3  | 17        |
| 22 | Scalable optimal Bayesian classification of single-cell trajectories under regulatory model uncertainty. BMC Genomics, 2019, 20, 435.                                      | 2.8  | 16        |
| 23 | Gene regulatory network state estimation from arbitrary correlated measurements. Eurasip Journal on Advances in Signal Processing, 2018, 2018, .                           | 1.7  | 15        |
| 24 | Supremal Multiscale Signal Analysis. SIAM Journal on Mathematical Analysis, 2004, 36, 94-120.  | 1.9  | 13        |
| 25 | Classification of State Trajectories in Gene Regulatory Networks. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018, 15, 68-82.                      | 3.0  | 11        |
| 26 | Classification and Error Estimation for Discrete Data. Current Genomics, 2009, 10, 446-462.  | 1.6  | 9         |
| 27 | Nutrient optimization for plant growth in Aquaponic irrigation using Machine Learning for small training datasets. Artificial Intelligence in Agriculture, 2022, 6, 68-76. | 6.0  | 9         |
| 28 | Exact Performance of CoD Estimators in Discrete Prediction. Eurasip Journal on Advances in Signal Processing, 2010, 2010, .  | 1.7  | 7         |
| 29 | Constructing multiscale connectivities. Computer Vision and Image Understanding, 2005, 99, 126-150.  | 4.7  | 6         |
| 30 | Studying the possibility of peaking phenomenon in linear support vector machines with non-separable data. , $2011,  ,  .$  |      | 6         |
| 31 | Small-sample error estimation: mythology versus mathematics. , 2005, , .   |      | 4         |
| 32 | Maximum likelihood estimation of the binary Coefficient of Determination. , 2011, , .  |      | 3         |
| 33 | Optimal Bayesian MMSE estimation of the coefficient of determination for discrete prediction. , 2013, ,  |      | 3         |
| 34 | A Naive-Bayes approach to Bolstered error estimation in high-dimensional spaces. , 2014, , .   |      | 3         |
| 35 | An asymptotically-exact expression for the variance of classification error for the discrete histogram rule. , 2008, , .   |      | 2         |
| 36 | Sample-based estimators for the instrinsically multivariate prediction score., 2011,,.   |      | 2         |

| #  | Article   | IF | CITATIONS |
|----|---|----|-----------|
| 37 | A Semi-Supervised Generative Adversarial Network for Prediction of Genetic Disease Outcomes. , 2021, , .  |    | 2         |
| 38 | Approximate expressions for the variances of non-randomized error estimators and CoD estimators for the discrete histogram rule. , $2010$ , , . |    | 0         |