

Foglio1

Risultati Seminari
 Corso Riconoscimento e Recupero dell'informazione per Bioinformatica

Titolo	Voto
Learning from healthy and stable eyes: a new approach for detection of glaucomatous progression	14.5/15
Identifying representative drug resistant mutants of HIV	14/15
A computational method for drug repositioning using publicly available gene expression data	14.5/15
Automated identification of copepods using digital image processing and artificial neural network	14/15
Gene selection for the reconstruction of stem cell differentiation trees: a linear programming approach	15/15
DINGO: Differential Network Analysis in Genomics	14.5/15
Pattern recognition methods to relate time profiles of gene expression with phenotypic data: a comparative study	14.5/15
Hierarchical boosting: a machine-learning framework to detect and classify hard selective sweeps in human populations	14/15
Differential diagnosis of pleural mesothelioma using Logic Learning Machine	13/15
HMMvar-func: a new method for predicting the functional outcome of genetic variants	15/15
Identification of a small set of plasma signalling proteins using neural network for prediction of Alzheimer's disease	15/15
Data-intensive analysis of HIV mutations	13.5/15
Inferring dynamic gene regulatory networks in cardiac differentiation through the integration of multi-dimensional data	14.5/15
Probabilistic models of genetic variation in structured populations applied to global human studies	14.5/15
An auditory feature detection circuit for sound pattern recognition	12.5/15
viralmiR: a support vector machine based method for predicting viral microRNA precursors	13.5/15
In-vitro diagnosis of single and poly microbial species targeted for diabetic foot infection using e-nose technology"	14.5/15
Robust feature selection to predict tumor treatment outcome	14/15
A Gaussian mixture model based cost function for parameter estimation of chaotic biological systems	14/15