Accepted Manuscript

Machine learning techniques to discover genes with Potential Prognosis Role in Alzheimer's Disease using different biological sources

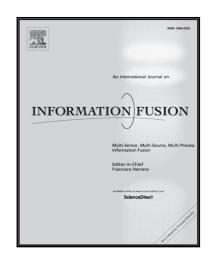
María Martínez-Ballesteros, José M. García-Heredia, Isabel A. Nepomuceno-Chamorro, José C. Riquelme-Santos

PII: S1566-2535(16)30130-0 DOI: 10.1016/j.inffus.2016.11.005

Reference: INFFUS 816

To appear in: Information Fusion

Received date: 22 October 2015
Revised date: 10 September 2016
Accepted date: 3 November 2016



Please cite this article as: María Martínez-Ballesteros, José M. García-Heredia, Isabel A. Nepomuceno-Chamorro, José C. Riquelme-Santos, Machine learning techniques to discover genes with Potential Prognosis Role in Alzheimer's Disease using different biological sources, *Information Fusion* (2016), doi: 10.1016/j.inffus.2016.11.005

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

ACCEPTED MANUSCRIPT

Highlights

- Analyze Alzheimer disease gene expression profiles by changes in expression levels
- Integration of machine learning methods: decision tree, associations and clustering
- Fusion of external information sources: microarray, PubMed, GO and PPI network
- Significant set of down/up regulated genes highly related with Alzheimer
- Gene expression patterns and deep knowledge into relevant biological functions

Download English Version:

https://daneshyari.com/en/article/4969142

Download Persian Version:

https://daneshyari.com/article/4969142

<u>Daneshyari.com</u>