Ficha de investigador

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Grupo de Investigación: RAZONAMIENTO APROXIMADO E INTELIGENCIA ARTIFICIAL (Cod.: TIC111)
Departamento: Universidad de Granada. Ciencias de la Computación e Inteligencia Artificial
Citas en Google Scholar:https://scholar.google.es/citations?user=r8GiY20AAAAJ&hl=es
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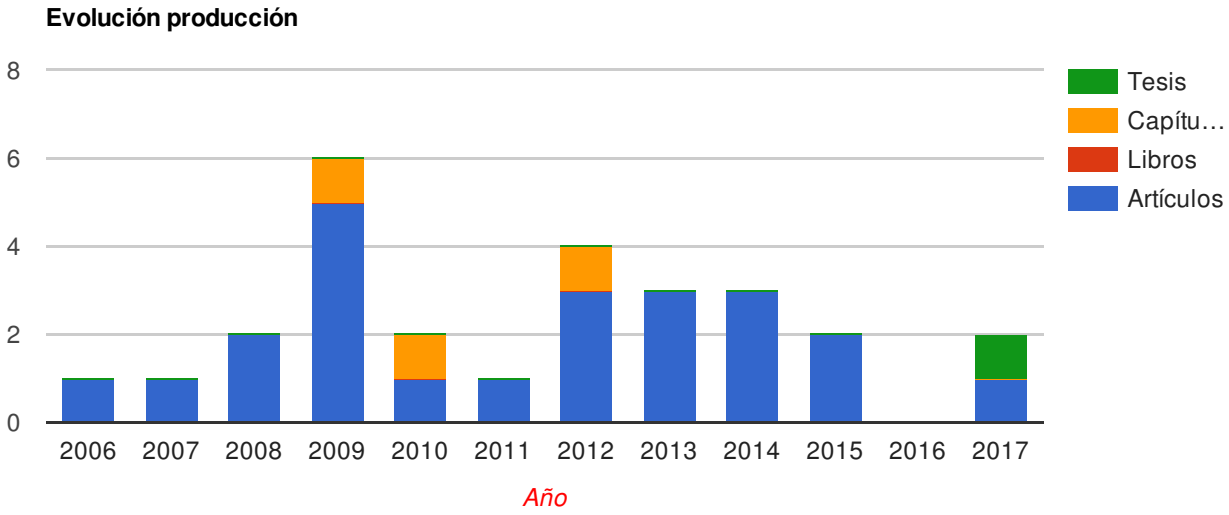
Producción 27

Artículos (23) Libros (0) Capítulos de Libros (3) Tesis dirigidas (1)



Proyectos dirigidos 0

Proyectos (0) Contratos (0) Convenios (0)

Actividades 0

Titulo publicación	Fuente	Tipo	Fecha
Aproximación a la medicina personalizada mediante el desarrollo de nuevas metodologías en inteligencia artificial	Universidad de granada. ciencias de la computación e inteligencia artificial, granada, españa.	Tesis doctoral	2017
Preoperative chemoradiotherapy for rectal cancer: the sensitizer role of the association between mir-375 and c-myc	Oncotarget	Articulo	2017
Drugnet: network-based drug-disease prioritization by integrating heterogeneous data	Artificial intelligence in medicine	Articulo	2015
Predictive biomarkers to chemoradiation in locally advanced rectal cancer	Biomed research international	Articulo	2015
Cisminer: genome-wide in silico cis-regulatory module prediction by fuzzy itemset mining.	Plos one	Articulo	2014
Expression profiling of rectal tumors defines response to neoadjuvant treatment related genes	Plos one	Articulo	2014
Prophnet: a generic network-based method of prioritization through propagation of information	Bmc bioinformatics	Articulo	2014
Expression profiling of breast tumors based on human epidermal growth factor receptor 2 status defines migration-related genes	Pathobiology	Articulo	2013
Microarray profiling of mononuclear peripheral blood cells identifies novel candidate genes related to chemoradiation response in rectal cancer	Plos one	Articulo	2013
Systems biology as a comparative approach to understand complex gene expression in neurological diseases.	Behavioral sciences	Articulo	2013
A public resource facilitating clinical use of genomes	Proceedings of the national academy of sciences of the united states of america	Articulo	2012
Biomedical application of fuzzy association rules for identifying breast cancer biomarkers	Medical & biological engineering & computing	Articulo	2012
Computer-assisted annotation of snps in regulatory regions of whole-genome sequencing data	Principles and applications of biomedical engineering	Capítulo de libro	2012
Fuzzy association rules. a new tool for identifying breast cancer biomarkers	Computer methods and programs in biomedicine	Articulo	2012
Genome-wide differential genetic profiling characterizes colorectal cancers with genetic instability and specific routes to hla class i loss and immune escape	Cancer immunology, immunotherapy	Articulo	2011
Her2 status in breast cancer: experience of a spanish national reference centre	Clinical & translational oncology	Articulo	2010
Social and semantic web technologies for the text-to-knowledge translation process in biomedicine	Biomedical engineering, trends, researches and technologies	Capítulo de libro	2010
A fuzzy approach for studying combinatorial regulatory actions of transcription factors in yeast	Lecture notes in computer science	Articulo	2009
Automated identification of diagnosis and co-morbidity in clinical records	Methods of information in medicine	Articulo	2009
Collaborative text-annotation resource for disease-centered relation extraction from biomedical text	Journal of biomedical informatics	Articulo	2009
Extracting biological knowledge by association rule mining	Data mining in biomedicine using ontologies	Capítulo de libro	2009
Fisim: a new similarity measure between transcription factor binding sites based on the fuzzy integral	Bmc bioinformatics	Articulo	2009
Intelligent system for the analysis of microarray data using principal components and estimation of distribution	Expert systems with applications	Articulo	2009

algorithms			
Análisis bioinformático de datos: aplicación en microarrays	Cuadernos de medicina reproductiva	Articulo	2008
Fuzzy association rules for biological data analysis: a case study on yeast	Bmc bioinformatics	Articulo	2008
Possibilistic approach for biclustering microarray data	Computers in biology and medicine	Articulo	2007
Evolutionary algorithms for finding interpretable patterns in gene expression data	Iadis international journal on computer science and information systems	Articulo	2006

	Titulo proyecto	Tipo	Inicio	Fin

Actividades 0

Titulo actividad	Fuente	Tipo	Fecha

Colaboradores

- ARMANDO BLANCO MORON (18)
- MARTA EUGENIA CUADROS CELORRIO (9)
- Carmen Navarro Luzón (2)
- PEDRO PABLO MEDINA VICO (2)
- ANTONIO MARIN RODRIGUEZ (1)
- Juan Antonio Morente Molinera (1)
- WALDO FAJARDO CONTRERAS (1)