

DAVID Bioinformatics Resources: expanded annotation better extract biology from large gene lists

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Acclimation of <i>Saccharomyces cerevisiae</i> to Low Temperature: A Chemostat-based Transcriptome Analysis. <i>Molecular Biology of the Cell</i> , 2007, 18, 5100-5112.	0.9	106
2	Integrative decomposition procedure and Kappa statistics set up ATF2 ion binding module in malignant pleural mesothelioma (MPM). <i>Frontiers of Electrical and Electronic Engineering in China: Selected Publications From Chinese Universities</i> , 2008, 3, 381-387.	0.6	23
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4	The molecular basis of genistein-induced mitotic arrest and exit of self-renewal in embryonal carcinoma and primary cancer cell lines. <i>BMC Medical Genomics</i> , 2008, 1, 49.	0.7	34
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7	Defining functional distances over Gene Ontology. <i>BMC Bioinformatics</i> , 2008, 9, 50.	1.2	55
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20	MicroRNA Classifiers for Predicting Prognosis of Squamous Cell Lung Cancer. Cancer Research, 2009, 69, 5776-5783.	0.4	385
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