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CÓRDOBA, 11 AL 13 DE MAYO

FDMDE, a free software framework for protein structure research using discrete- events molecular dynamics

Pablo Ybalo, Nicolás Brailovsky, Hernán Curras, Marcos Villarreal, Daniel Gutson

Feature analysis for activity identification in p53 protein mutant segments

Simonetti Franco L, Hadad Alejandro

First application of a high density microarray for Sunflower: Methodological aspects of the design and statistical analysis in the study of the tolerance to Sclerotinia sclerotiorum

Julio A. Di Rienzo, Lucila Peluffo, Darío Principi, Paula Fernández, Verónica Lía, Federico Ehrenbolger, Ana Conesa, Francisco García, David Blesa, Joaquín Dopazo, Norma Paniego y Ruth A. Heinz

Functional diversity in plant HD-Zip subfamily I transcription factors. Using phylogenetic and motif discovery approaches to look beyond conserved domains

Agustín L Arce, Jesica Raineri, Matías Capella, Julieta V Cabello, Raquel L Chan

Fuzzy Multiobjetive Optimization for Chemotherapy Schedules

Andrés Barrea and Matías Hernández

Gender-specific tumor pathways in colon cancer revealed by ontology analysis of tissue by gender gene interaction patterns.

Fernandez, EA, Salvatierra E, Gidekel M, Fresno C, Llera A, Balzarini M, Podhajcer O

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