Maria Rosa Felice

List of Publications by Year in descending order

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Version: 2024-02-01

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#	Article	IF	CITATIONS
1	Cytotoxicity, haemolymphatic parameters, and oxidative stress following exposure to sub-lethal concentrations of quaternium-15 in Mytilus galloprovincialis. Aquatic Toxicology, 2016, 180, 258-265.	4.0	182
2	The essential role of Glu-185 and Tyr-354 residues in the ferroxidase activity of Saccharomyces cerevisiae Fet3. FEBS Letters, 2000, 472, 283-286.	2.8	35
3	Whole Genome-Based Amplified Fragment Length Polymorphism Analysis Reveals Genetic Diversity in Candida africana. Frontiers in Microbiology, 2017, 8, 556.	3.5	19
4	Copper chelation and interleukin-6 proinflammatory cytokine effects on expression of different proteins involved in iron metabolism in HepG2 cell line. BMC Biochemistry, 2017, 18, 1.	4.4	18
5	Mutational analysis of the iron binding site of Saccharomyces cerevisiae ferroxidase Fet 3. An in vivo study. FEBS Letters, 2001, 508, 475-478.	2.8	14
6	Genetic diversity of Candida albicans isolates recovered from hospital environments and patients with severe acquired brain injuries. Infection, Genetics and Evolution, 2019, 76, 104068.	2.3	10
7	Molecular Characterization of the N-Acetylglucosamine Catabolic Genes in Candida africana, a Natural N-Acetylglucosamine Kinase (HXK1) Mutant. PLoS ONE, 2016, 11, e0147902.	2.5	10
8	Specific aspartate residues in FET3 control high-affinity iron transport inSaccharomyces cerevisiae. Yeast, 2005, 22, 677-687.	1.7	9
9	Regulation of prostaglandin generation in carrageenan-induced pleurisy by inducible nitric oxide synthase in knockout mice. Life Sciences, 2003, 72, 1199-1208.	4.3	8
10	Whole RNA-Sequencing and Transcriptome Assembly of Candida albicans and Candida africana under Chlamydospore-Inducing Conditions. Genome Biology and Evolution, 2017, 9, 1971-1977.	2.5	8
11	Looking for New Antifungal Drugs from Flavonoids: Impact of the Genetic Diversity of Candida albicans on the in-vitro Response. Current Medicinal Chemistry, 2019, 26, 5108-5123.	2.4	6