

Lina R Riego-Ruiz

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Two alpha isopropylmalate synthase isozymes with similar kinetic properties are extant in the yeast <i>Lachancea kluyveri</i> . FEMS Yeast Research, 2022, 22, .	2.3	2
2	Synthetic negative genome screen of the GPN-loop GTPase NPA3 in <i>Saccharomyces cerevisiae</i> . Current Genetics, 2022, 68, 343-360.	1.7	3
3	Long-insert clone experimental evidence for assembly improvement and chimeric chromosomes detection in an allopolyploid beer yeast. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	2
4	In <i>Kluyveromyces lactis</i> a Pair of Paralogous Isozymes Catalyze the First Committed Step of Leucine Biosynthesis in Either the Mitochondria or the Cytosol. Frontiers in Microbiology, 2020, 11, 1843.	3.5	4
5	A Lineage of Begomoviruses Encode Rep and AC4 Proteins of Enigmatic Ancestry: Hints on the Evolution of Geminiviruses in the New World. Viruses, 2019, 11, 644.	3.3	13
6	Whole-Genome Duplication and Yeast's Fruitful Way of Life. Trends in Genetics, 2019, 35, 42-54.	6.7	23
7	Diversification of the kinetic properties of yeast NADP-glutamate dehydrogenase isozymes proceeds independently of their evolutionary origin. MicrobiologyOpen, 2017, 6, e00419.	3.0	9
8	Npa3/ScGpn1 carboxy-terminal tail is dispensable for cell viability and RNA polymerase II nuclear targeting but critical for microtubule stability and function. Biochimica Et Biophysica Acta - Molecular Cell Research, 2017, 1864, 451-462.	4.1	15
9	Diversification of Transcriptional Regulation Determines Subfunctionalization of Paralogous Branched Chain Aminotransferases in the Yeast <i>Saccharomyces cerevisiae</i> . Genetics, 2017, 207, 975-991.	2.9	27
10	Haemosporidian parasite prevalence, parasitemia, and diversity in three resident bird species at a shrubland dominated landscape of the Mexican highland plateau. Parasites and Vectors, 2016, 9, 307.	2.5	19
11	Functional roles of a predicted branched chain aminotransferase encoded by the LkBAT1 gene of the yeast <i>Lachancea kluyveri</i> . Fungal Genetics and Biology, 2015, 85, 71-82.	2.1	9
12	Genetic diversity of the Mycobacterium tuberculosis Complex in San Luis Potosí, México. BMC Research Notes, 2013, 6, 172.	1.4	24
13	DNA-binding specificity determinants of replication proteins encoded by eukaryotic ssDNA viruses are adjacent to widely separated RCR conserved motifs. Archives of Virology, 2010, 155, 1033-1046.	2.1	46
14	Ancestry informative markers and admixture proportions in northeastern Mexico. Journal of Human Genetics, 2009, 54, 504-509.	2.3	40
15	High Resistance to Oxidative Stress in the Fungal Pathogen <i>Candida glabrata</i> Is Mediated by a Single Catalase, Cta1p, and Is Controlled by the Transcription Factors Yap1p, Skn7p, Msn2p, and Msn4p. Eukaryotic Cell, 2008, 7, 814-825.	3.4	147