

Ana Catarina Gomes

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

1,257
citations

567281

15
h-index

888059

17
g-index

20
all docs

20
docs citations

20
times ranked

1973
citing authors

#	ARTICLE	IF	CITATIONS
1	Understand the Potential Role of <i>Aureobasidium pullulans</i> , a Resident Microorganism From Grapevine, to Prevent the Infection Caused by <i>Diplodia seriata</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 3047.	3.5	45
2	<i>Vitis vinifera</i> microbiome: from basic research to technological development. <i>BioControl</i> , 2016, 61, 243-256.	2.0	44
3	The effects of grapevine trunk diseases (GTDs) on vine physiology. <i>European Journal of Plant Pathology</i> , 2016, 144, 707-721.	1.7	103
4	Wine fermentation microbiome: a landscape from different Portuguese wine appellations. <i>Frontiers in Microbiology</i> , 2015, 6, 905.	3.5	170
5	Exploring the <i>Saccharomyces cerevisiae</i> Volatile Metabolome: Indigenous versus Commercial Strains. <i>PLoS ONE</i> , 2015, 10, e0143641.	2.5	51
6	Establishing the Yeast <i>Kluyveromyces lactis</i> as an Expression Host for Production of the Saposin-Like Domain of the Aspartic Protease Cirsin. <i>Applied and Environmental Microbiology</i> , 2014, 80, 86-96.	3.1	12
7	Inflammatory modulation of stem cells by Magnetic Resonance Imaging (MRI)-detectable nanoparticles. <i>RSC Advances</i> , 2014, 4, 31706-31709.	3.6	9
8	Unravelling the Diversity of Grapevine Microbiome. <i>PLoS ONE</i> , 2014, 9, e85622.	2.5	268
9	Low level genome mistranslations deregulate the transcriptome and translome and generate proteotoxic stress in yeast. <i>BMC Biology</i> , 2012, 10, 55.	3.8	31
10	Genetic Diversity and Population Structure of <i>Saccharomyces cerevisiae</i> Strains Isolated from Different Grape Varieties and Winemaking Regions. <i>PLoS ONE</i> , 2012, 7, e32507.	2.5	81
11	The genetic code of the fungal CTG clade. <i>Comptes Rendus - Biologies</i> , 2011, 334, 607-611.	0.2	90
12	Genotyping of <i>Saccharomyces cerevisiae</i> strains by interdelta sequence typing using automated microfluidics. <i>Electrophoresis</i> , 2011, 32, 1447-1455.	2.4	19
13	Parallel DNA pyrosequencing unveils new zebrafish microRNAs. <i>BMC Genomics</i> , 2009, 10, 195.	2.8	65
14	The Yeast PNC1 Longevity Gene Is Up-Regulated by mRNA Mistranslation. <i>PLoS ONE</i> , 2009, 4, e5212.	2.5	15
15	Comparative genomics of wild type yeast strains unveils important genome diversity. <i>BMC Genomics</i> , 2008, 9, 524.	2.8	111
16	Populational analysis of <i>Saccharomyces cerevisiae</i> strains from different appellations of origin and grape varieties by microsatellite analysis. <i>Nature Precedings</i> , 2008, , .	0.1	0
17	A genetic code alteration generates a proteome of high diversity in the human pathogen <i>Candida albicans</i> . <i>Genome Biology</i> , 2007, 8, R206.	9.6	99
18	Critical roles for a genetic code alteration in the evolution of the genus <i>Candida</i> . <i>EMBO Journal</i> , 2007, 26, 4555-4565.	7.8	43

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19	The Genetic Code of the Candida CTG Clade. , 0, , 45-55.		0