VerÃ³nica MixÃ£o

List of Publications by Year
in descending order

16	487	933447 10	940533 16
papers	citations	h-index	g-index

22	22	22	606
all docs	docs citations	times ranked	citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple Hybridization Events Punctuate the Evolutionary Trajectory ofMalassezia furfur. MBio, 2022, 13, e0385321.	4.1	9
2	Genome analysis of five recently described species of the CUG-Ser clade uncovers Candida theae as a new hybrid lineage with pathogenic potential in the Candida parapsilosis species complex. DNA Research, 2022, , .	3.4	4
3	Genome analysis of Candida subhashii reveals its hybrid nature and dual mitochondrial genome conformations. DNA Research, 2021, 28, .	3.4	14
4	Factors enforcing the species boundary between the human pathogens Cryptococcus neoformans and Cryptococcus deneoformans. PLoS Genetics, 2021, 17, e1008871.	3.5	13
5	Extreme diversification driven by parallel events of massive loss of heterozygosity in the hybrid lineage of Candida albicans. Genetics, 2021, 217, .	2.9	16
6	HaploTypo: a variant-calling pipeline for phased genomes. Bioinformatics, 2020, 36, 2569-2571.	4.1	14
7	Differential Expression of Fungal Genes Determines the Lifestyle of Plectosphaerella Strains During Arabidopsis thaliana Colonization. Molecular Plant-Microbe Interactions, 2020, 33, 1299-1314.	2.6	9
8	Genomic evidence for a hybrid origin of the yeast opportunistic pathogen Candida albicans. BMC Biology, 2020, 18, 48.	3.8	46
9	Recent trends in molecular diagnostics of yeast infections: from PCR to NGS. FEMS Microbiology Reviews, 2019, 43, 517-547.	8.6	77
10	Whole-Genome Sequencing of the Opportunistic Yeast Pathogen Candida inconspicua Uncovers Its Hybrid Origin. Frontiers in Genetics, 2019, 10, 383.	2.3	63
11	Genome Assemblies of Two Rare Opportunistic Yeast Pathogens:Diutina rugosa(syn.Candida) Tj ETQq1 1 0.784314 rgBT /Ove Genetics, 2019, 9, 3921-3927.	1.8	6
12	Misidentification of genome assemblies in public databases: The case of Naumovozyma dairenensis and proposal of a protocol to correct misidentifications. Yeast, 2018, 35, 425-429.	1.7	30
13	Hybridization and emergence of virulence in opportunistic human yeast pathogens. Yeast, 2018, 35, 5-20.	1.7	104
14	Comparative morphological and molecular analysis confirms the presence of the West Nile virus mosquito vector, Culex univittatus, in the Iberian Peninsula. Parasites and Vectors, 2016, 9, 601.	2.5	22
15	Molecular detection of Wolbachia pipientis in natural populations of mosquito vectors of Dirofilaria immitis from continental Portugal: first detection in Culex theileri. Medical and Veterinary Entomology, 2016, 30, 301-309.	1.5	13
16	First molecular identification of mosquito vectors of Dirofilaria immitis in continental Portugal. Parasites and Vectors, 2015, 8, 139.	2.5	43