Douda Bensasson

List of Publications by Year
in descending order

25 papers	4,669 citations	394286 19 h-index	552653 26 g-index
32 all docs	32 docs citations	32 times ranked	5921 citing authors

#	Article	IF	Citations
1	Genetic variation in aneuploidy prevalence and tolerance across <i>Saccharomyces cerevisiae</i> lineages. Genetics, 2021, 217, .	1.2	25
2	Sporulation environment drives phenotypic variation in the pathogen <i>Aspergillus fumigatus</i>. G3: Genes, Genomes, Genetics, 2021, 11, .	0.8	11
3	Evolution of Ty1 copy number control in yeast by horizontal transfer and recombination. PLoS Genetics, 2020, 16, e1008632.	1.5	30
4	Diverse Lineages of<i>Candida albicans</i>Live on Old Oaks. Genetics, 2019, 211, 277-288.	1.2	54
5	Phased Diploid Genome Assemblies for Three Strains of <i>Candida albicans</i> from Oak Trees. G3: Genes, Genomes, Genetics, 2019, 9, 3547-3554.	0.8	6
6	mSphere of Influence: the Wild Genetic Diversity of Our Closest Yeast Companions. MSphere, 2019, 4, .	1.3	1
7	Adaptive divergence in wine yeasts and their wild relatives suggests a prominent role for introgressions and rapid evolution at noncoding sites. Molecular Ecology, 2017, 26, 2167-2182.	2.0	44
8	Habitat Predicts Levels of Genetic Admixture in<i>Saccharomyces cerevisiae</i>. G3: Genes, Genomes, Genetics, 2017, 7, 2919-2929.	0.8	19
9	Summer temperature can predict the distribution of wild yeast populations. Ecology and Evolution, 2016, 6, 1236-1250.	0.8	59
10	A population genomics insight into the Mediterranean origins of wine yeast domestication. Molecular Ecology, 2015, 24, 5412-5427.	2.0	186
11	Evolutionary Genomics of Transposable Elements in Saccharomyces cerevisiae. PLoS ONE, 2012, 7, e50978.	1.1	91
12	Evidence for a high mutation rate at rapidly evolving yeast centromeres. BMC Evolutionary Biology, 2011, 11, 211.	3.2	30
13	Population genomics of domestic and wild yeasts. Nature, 2009, 458, 337-341.	13.7	1,391
14	Population genomics of the wild yeast <i>Saccharomyces paradoxus</i> : Quantifying the life cycle. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4957-4962.	3.3	287
15	Rapid Evolution of Yeast Centromeres in the Absence of Drive. Genetics, 2008, 178, 2161-2167.	1.2	57
16	Population genomics of domestic and wild yeasts. Nature Precedings, 2008, , .	0.1	1
17	Transition-Transversion Bias Is Not Universal: A Counter Example from Grasshopper Pseudogenes. PLoS Genetics, 2007, 3, e22.	1.5	128
18	Release and persistence of extracellular DNA in the environment. Environmental Biosafety Research, 2007, 6, 37-53.	1.1	461

#	Article	IF	Citations
19	Recent LTR retrotransposon insertion contrasts with waves of non-LTR insertion since speciation in Drosophila melanogaster. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 11340-11345.	3.3	93
20	Mitochondrial genome sequences and comparative genomics of Phytophthora ramorum and P. sojae. Current Genetics, 2007, 51, 285-296.	0.8	48
21	Phytophthora Genome Sequences Uncover Evolutionary Origins and Mechanisms of Pathogenesis. Science, 2006, 313, 1261-1266.	6.0	1,059
22	Rates of DNA Duplication and Mitochondrial DNA Insertion in the Human Genome. Journal of Molecular Evolution, 2003, 57, 343-354.	0.8	112
23	Size Matters: Non-LTR Retrotransposable Elements and Ectopic Recombination in Drosophila. Molecular Biology and Evolution, 2003, 20, 880-892.	3.5	208
24	Genomic Gigantism: DNA Loss Is Slow in Mountain Grasshoppers. Molecular Biology and Evolution, 2001, 18, 246-253.	3.5	111
25	Frequent Assimilation of Mitochondrial DNA by Grasshopper Nuclear Genomes. Molecular Biology and Evolution, 2000, 17, 406-415.	3.5	147