Guillaume Lesage

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12021006/publications.pdf

Version: 2024-02-01

759233 1125743 3,362 13 12 13 h-index citations g-index papers 13 13 13 4271 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Global Mapping of the Yeast Genetic Interaction Network. Science, 2004, 303, 808-813.	12.6	1,908
2	Cell Wall Assembly in Saccharomyces cerevisiae. Microbiology and Molecular Biology Reviews, 2006, 70, 317-343.	6.6	673
3	Combining biological networks to predict genetic interactions. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 15682-15687.	7.1	225
4	Motifs, themes and thematic maps of an integrated Saccharomyces cerevisiae interaction network. Journal of Biology, 2005, 4, 6.	2.7	154
5	Analysis of \hat{l}^2 -1,3-Glucan Assembly in Saccharomyces cerevisiae Using a Synthetic Interaction Network and Altered Sensitivity to Caspofungin. Genetics, 2004, 167, 35-49.	2.9	149
6	An interactional network of genes involved in chitin synthesis in Saccharomyces cerevisiae. BMC Genetics, 2005, 6, 8.	2.7	105
7	A synthetic analysis of the Saccharomyces cerevisiae stress sensor Mid2p, and identification of a Mid2p-interacting protein, Zeo1p, that modulates the PKC1–MPK1 cell integrity pathway. Microbiology (United Kingdom), 2003, 149, 2487-2499.	1.8	46
8	The Kex2p Proregion Is Essential for the Biosynthesis of an Active Enzyme and Requires a C-terminal Basic Residue for Its Function. Molecular Biology of the Cell, 2000, 11, 1947-1957.	2.1	26
9	Large-Scale Analysis of Genes that Alter Sensitivity to the Anticancer Drug Tirapazamine inSaccharomyces cerevisiae. Molecular Pharmacology, 2005, 68, 1365-1375.	2.3	22
10	Integrative studies put cell wall synthesis on the yeast functional map. Current Opinion in Microbiology, 2004, 7, 617-623.	5.1	21
11	Mechanism of Kex2p inhibition by its proregion. FEBS Letters, 2001, 508, 332-336.	2.8	17
12	Genetic interaction network of the Saccharomyces cerevisiae type 1 phosphatase Glc7. BMC Genomics, 2008, 9, 336.	2.8	14
13	trans-Complementation assay establishes the role of proregion hydrophobic amino acid residues in the biosynthesis of Saccharomyces cerevisiae Kex2p endoprotease. Yeast, 2003, 20, 397-406.	1.7	2