Michael J Law

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Distinct requirements for the COMPASS core subunits Set1, Swd1, and Swd3 during meiosis in the budding yeast <i>Saccharomyces cerevisiae</i> . G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	3
2	The <i>Saccharomyces cerevisiae</i> Cdk8 Mediator Represses <i>AQY1</i> Transcription by Inhibiting Set1p-Dependent Histone Methylation. G3: Genes, Genomes, Genetics, 2017, 7, 1001-1010.	1.8	6
3	The conserved histone deacetylase Rpd3 and its DNA binding subunit Ume6 control dynamic transcript architecture during mitotic growth and meiotic development. Nucleic Acids Research, 2015, 43, 115-128.	14.5	29
4	Global alterations of the transcriptional landscape during yeast growth and development in the absence of Ume6-dependent chromatin modification. Molecular Genetics and Genomics, 2015, 290, 2031-2046.	2.1	11
5	Integrated RNA- and protein profiling of fermentation and respiration in diploid budding yeast provides insight into nutrient control of cell growth and development. Journal of Proteomics, 2015, 119, 30-44.	2.4	5
6	Fine-Tuning of Histone H3 Lys4 Methylation During Pseudohyphal Differentiation by the CDK Submodule of RNA Polymerase II. Genetics, 2015, 199, 435-453.	2.9	19
7	The histone deacetylase Rpd3/Sin3/Ume6 complex represses an acetateâ€inducible isoform of <i>VTH2</i> in fermenting budding yeast cells. FEBS Letters, 2015, 589, 924-932.	2.8	2
8	The conserved histone deacetylase <scp>R</scp> pd3 and the <scp>DNA</scp> binding regulator <scp>U</scp> me6 repress <scp><i>BOI</i></scp> <i>1</i> 's meiotic transcript isoform during vegetative growth in <scp><i>S</i></scp> <i>accharomyces cerevisiae</i> . Molecular Microbiology, 2015, 96, 861-874.	2.5	10
9	Acetylation of the Transcriptional Repressor Ume6p Allows Efficient Promoter Release and Timely Induction of the Meiotic Transient Transcription Program in Yeast. Molecular and Cellular Biology, 2014, 34, 631-642.	2.3	22