

Michael J Law

List of Publications by Year in descending order

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Version: 2024-02-01

9
papers

107
citations

1478505

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1474206

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144
citing authors

#	ARTICLE	IF	CITATIONS
1	The conserved histone deacetylase Rpd3 and its DNA binding subunit Ume6 control dynamic transcript architecture during mitotic growth and meiotic development. <i>Nucleic Acids Research</i> , 2015, 43, 115-128.	14.5	29
2	Acetylation of the Transcriptional Repressor Ume6p Allows Efficient Promoter Release and Timely Induction of the Meiotic Transient Transcription Program in Yeast. <i>Molecular and Cellular Biology</i> , 2014, 34, 631-642.	2.3	22
3	Fine-Tuning of Histone H3 Lys4 Methylation During Pseudohyphal Differentiation by the CDK Submodule of RNA Polymerase II. <i>Genetics</i> , 2015, 199, 435-453.	2.9	19
4	Global alterations of the transcriptional landscape during yeast growth and development in the absence of Ume6-dependent chromatin modification. <i>Molecular Genetics and Genomics</i> , 2015, 290, 2031-2046.	2.1	11
5	The conserved histone deacetylase <i>Rpd3</i> and the <i>DNA</i> binding regulator <i>Ume6</i> repress <i>BOI1</i> 's meiotic transcript isoform during vegetative growth in <i>Saccharomyces cerevisiae</i> . <i>Molecular Microbiology</i> , 2015, 96, 861-874.	2.5	10
6	The <i>Saccharomyces cerevisiae</i> Cdk8 Mediator Represses <i>AQY1</i> Transcription by Inhibiting Set1p-Dependent Histone Methylation. <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 1001-1010.	1.8	6
7	Integrated RNA- and protein profiling of fermentation and respiration in diploid budding yeast provides insight into nutrient control of cell growth and development. <i>Journal of Proteomics</i> , 2015, 119, 30-44.	2.4	5
8	Distinct requirements for the COMPASS core subunits Set1, Swd1, and Swd3 during meiosis in the budding yeast <i>Saccharomyces cerevisiae</i> . <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	3
9	The histone deacetylase Rpd3/Sin3/Ume6 complex represses an acetate-inducible isoform of <i>VTH2</i> in fermenting budding yeast cells. <i>FEBS Letters</i> , 2015, 589, 924-932.	2.8	2