

Andrea A Duina

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

636
citations

932766

10
h-index

1125271

13
g-index

15
all docs

15
docs citations

15
times ranked

849
citing authors

#	ARTICLE	IF	CITATIONS
1	Budding Yeast for Budding Geneticists: A Primer on the <i>Saccharomyces cerevisiae</i> Model System. <i>Genetics</i> , 2014, 197, 33-48.	1.2	168
2	Requirement for Hsp90 and a CyP-40-type Cyclophilin in Negative Regulation of the Heat Shock Response. <i>Journal of Biological Chemistry</i> , 1998, 273, 18974-18978.	1.6	130
3	Identification of two CyP-40-like cyclophilins in <i>Saccharomyces cerevisiae</i> , one of which is required for normal growth. <i>J Biol Chem</i> , 1996, 271, 943-952.		76
4	The Peptidyl-prolyl Isomerase Domain of the CyP-40 Cyclophilin Homolog Cpr7 Is Not Required to Support Growth or Glucocorticoid Receptor Activity in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 1998, 273, 10819-10822.	1.6	50
5	Evidence that the Localization of the Elongation Factor Spt16 Across Transcribed Genes Is Dependent Upon Histone H3 Integrity in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2007, 177, 101-112.	1.2	46
6	Analysis of a Mutant Histone H3 That Perturbs the Association of Swi/Snf with Chromatin. <i>Molecular and Cellular Biology</i> , 2004, 24, 561-572.	1.1	39
7	Histone Chaperones Spt6 and FACT: Similarities and Differences in Modes of Action at Transcribed Genes. <i>Genetics Research International</i> , 2011, 2011, 1-12.	2.0	38
8	Mutant Versions of the <i>S. cerevisiae</i> Transcription Elongation Factor Spt16 Define Regions of Spt16 That Functionally Interact with Histone H3. <i>PLoS ONE</i> , 2011, 6, e20847.	1.1	24
9	Charged residues on the side of the nucleosome contribute to normal Spt16-gene interactions in budding yeast. <i>Epigenetics</i> , 2018, 13, 1-7.	1.3	19
10	A Nucleosomal Region Important for Ensuring Proper Interactions Between the Transcription Elongation Factor Spt16 and Transcribed Genes in <i>Saccharomyces cerevisiae</i> . <i>G3: Genes, Genomes, Genetics</i> , 2013, 3, 929-940.	0.8	18
11	Uncoupling of the Patterns of Chromatin Association of Different Transcription Elongation Factors by a Histone H3 Mutant in <i>Saccharomyces cerevisiae</i> . <i>Eukaryotic Cell</i> , 2009, 8, 257-260.	3.4	14
12	A Systematic Mutational Analysis of a Histone H3 Residue in Budding Yeast Provides Insights into Chromatin Dynamics. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 741-749.	0.8	10
13	Targeted <i>In Situ</i> Mutagenesis of Histone Genes in Budding Yeast. <i>Journal of Visualized Experiments</i> , 2017, . .	0.2	3
14	Evidence that dissociation of Spt16 from transcribed genes is partially dependent on RNA Polymerase II termination. <i>Transcription</i> , 2019, 10, 195-206.	1.7	1
15	New Roles for Old Characters: An Educational Primer for Use with <i>Vps</i> Factors Are Required for Efficient Transcription Elongation in Budding Yeast. <i>Genetics</i> , 2013, 194, 27-33.	1.2	0