

Shannon M Lauberth

List of Publications by Year in descending order

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9
papers

903
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1306789

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1473754

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times ranked

1698
citing authors

#	ARTICLE	IF	CITATIONS
1	H3K4me3 Interactions with TAF3 Regulate Preinitiation Complex Assembly and Selective Gene Activation. <i>Cell</i> , 2013, 152, 1021-1036.	13.5	353
2	Enhancer RNAs are an important regulatory layer of the epigenome. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 521-528.	3.6	214
3	RNAs interact with BRD4 to promote enhanced chromatin engagement and transcription activation. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 687-697.	3.6	123
4	Mutant p53 shapes the enhancer landscape of cancer cells in response to chronic immune signaling. <i>Nature Communications</i> , 2017, 8, 754.	5.8	71
5	The Basal Transcription Complex Component TAF3 Transduces Changes in Nuclear Phosphoinositides into Transcriptional Output. <i>Molecular Cell</i> , 2015, 58, 453-467.	4.5	67
6	A Phosphomimetic Mutation in the Sall1 Repression Motif Disrupts Recruitment of the Nucleosome Remodeling and Deacetylase Complex and Repression of Gbx2. <i>Journal of Biological Chemistry</i> , 2007, 282, 34858-34868.	1.6	33
7	Mutant p53 regulates enhancer-associated H3K4 monomethylation through interactions with the methyltransferase MLL4. <i>Journal of Biological Chemistry</i> , 2018, 293, 13234-13246.	1.6	22
8	It's a DoG-eat-DoG world" altered transcriptional mechanisms drive downstream-of-gene (DoG) transcript production. <i>Molecular Cell</i> , 2022, 82, 1981-1991.	4.5	12
9	The role of enhancer RNAs in epigenetic regulation of gene expression. <i>Transcription</i> , 2020, 11, 19-25.	1.7	8