

Robert N Eisenman

List of Publications by Year in descending order

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22
papers

4,039
citations

567281

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677142

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26
all docs

26
docs citations

26
times ranked

6264
citing authors

#	ARTICLE	IF	CITATIONS
1	Coordinated Cross-Talk Between the Myc and Mlx Networks in Liver Regeneration and Neoplasia. Cellular and Molecular Gastroenterology and Hepatology, 2022, 13, 1785-1804.	4.5	12
2	Loss of MGA repression mediated by an atypical polycomb complex promotes tumor progression and invasiveness. ELife, 2021, 10, .	6.0	26
3	MYC and TFEB Control DNA Methylation and Differentiation in AML. Blood Cancer Discovery, 2021, 2, 116-118.	5.0	4
4	The glucose-sensing transcription factor MLX balances metabolism and stress to suppress apoptosis and maintain spermatogenesis. PLoS Biology, 2021, 19, e3001085.	5.6	7
5	The MNT transcription factor autoregulates its expression and supports proliferation in MYC-associated factor X (MAX)-deficient cells. Journal of Biological Chemistry, 2020, 295, 2001-2017.	3.4	10
6	MAX Functions as a Tumor Suppressor and Rewires Metabolism in Small Cell Lung Cancer. Cancer Cell, 2020, 38, 97-114.e7.	16.8	46
7	Max deletion destabilizes MYC protein and abrogates E μ -Myc lymphomagenesis. Genes and Development, 2019, 33, 1252-1264.	5.9	41
8	Distinct gene-selective roles for a network of core promoter factors in Drosophila neural stem cell identity. Biology Open, 2019, 8, .	1.2	7
9	Pan-cancer Alterations of the MYC Oncogene and Its Proximal Network across the Cancer Genome Atlas. Cell Systems, 2018, 6, 282-300.e2.	6.2	284
10	The MYC transcription factor network: balancing metabolism, proliferation and oncogenesis. Frontiers of Medicine, 2018, 12, 412-425.	3.4	187
11	Myc/Mycn-mediated glycolysis enhances mouse spermatogonial stem cell self-renewal. Genes and Development, 2016, 30, 2637-2648.	5.9	66
12	Quantitative Method to Investigate the Balance between Metabolism and Proteome Biomass: Starting from Glycine. Angewandte Chemie - International Edition, 2016, 55, 15646-15650.	13.8	42
13	Metabolomics method to comprehensively analyze amino acids in different domains. Analyst, The, 2015, 140, 2726-2734.	3.5	39
14	Deregulated Myc Requires MondoA/Mlx for Metabolic Reprogramming and Tumorigenesis. Cancer Cell, 2015, 27, 271-285.	16.8	172
15	The glucose-sensing transcription factor MLX promotes myogenesis via myokine signaling. Genes and Development, 2015, 29, 2475-2489.	5.9	38
16	Functional interactions among members of the MAX and MLX transcriptional network during oncogenesis. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2015, 1849, 484-500.	1.9	91
17	An Overview of MYC and Its Interactome. Cold Spring Harbor Perspectives in Medicine, 2014, 4, a014357-a014357.	6.2	317
18	The Drosophila ubiquitin-specific protease Puffeye regulates dMyc-mediated growth. Development (Cambridge), 2013, 140, 4776-4787.	2.5	15

#	ARTICLE	IF	CITATIONS
19	Myc's broad reach. <i>Genes and Development</i> , 2008, 22, 2755-2766.	5.9	826
20	Mnt-Deficient Mammary Glands Exhibit Impaired Involution and Tumors with Characteristics of Myc Overexpression. <i>Cancer Research</i> , 2006, 66, 5565-5573.	0.9	37
21	The Myc/Max/Mad Network and the Transcriptional Control of Cell Behavior. <i>Annual Review of Cell and Developmental Biology</i> , 2000, 16, 653-699.	9.4	1,182
22	Mad-max transcriptional repression is mediated by ternary complex formation with mammalian homologs of yeast repressor Sin3. <i>Cell</i> , 1995, 80, 767-776.	28.9	585