Matthew D Galbraith

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptional regulation by hypoxia inducible factors. Critical Reviews in Biochemistry and Molecular Biology, 2014, 49, 1-15.	5.2	575
2	Mechanisms of transcriptional regulation by p53. Cell Death and Differentiation, 2018, 25, 133-143.	11.2	310
3	HIF1A Employs CDK8-Mediator to Stimulate RNAPII Elongation in Response to Hypoxia. Cell, 2013, 153, 1327-1339.	28.9	300
4	Trisomy 21 consistently activates the interferon response. ELife, 2016, 5, .	6.0	238
5	Global analysis of p53-regulated transcription identifies its direct targets and unexpected regulatory mechanisms. ELife, 2014, 3, e02200.	6.0	205
6	CDK8. Transcription, 2010, 1, 4-12.	3.1	184
7	Identification of a core TP53 transcriptional program with highly distributed tumor suppressive activity. Genome Research, 2017, 27, 1645-1657.	5.5	123
8	Mitogen-induced recruitment of ERK and MSK to SRE promoter complexes by ternary complex factor Elk-1. Nucleic Acids Research, 2008, 36, 2594-2607.	14.5	91
9	The TIP60 Complex Is a Conserved Coactivator of HIF1A. Cell Reports, 2016, 16, 37-47.	6.4	78
10	Mass Cytometry Reveals Global Immune Remodeling with Multi-lineage Hypersensitivity to Type I Interferon in Down Syndrome. Cell Reports, 2019, 29, 1893-1908.e4.	6.4	78
11	Therapeutic targeting of transcriptional cyclin-dependent kinases. Transcription, 2019, 10, 118-136.	3.1	78
12	CDK8 Kinase Activity Promotes Glycolysis. Cell Reports, 2017, 21, 1495-1506.	6.4	67
13	A Kinase-Independent Role for Cyclin-Dependent Kinase 19 in p53 Response. Molecular and Cellular Biology, 2017, 37, .	2.3	57
14	Specialized interferon action in COVID-19. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	56
15	Transcriptional Responses to IFN-Î ³ Require Mediator Kinase-Dependent Pause Release and Mechanistically Distinct CDK8 and CDK19 Functions. Molecular Cell, 2019, 76, 485-499.e8.	9.7	52
16	Biochemical and genetic diversity of pectolytic enterobacteria causing soft rot disease of potatoes in New Zealand. Australasian Plant Pathology, 2008, 37, 559.	1.0	42
17	Seroconversion stages COVID19 into distinct pathophysiological states. ELife, 2021, 10, .	6.0	40
18	ΔNp63α Suppresses TGFB2 Expression and RHOA Activity to Drive Cell Proliferation in Squamous Cell Carcinomas. Cell Reports, 2018, 24, 3224-3236.	6.4	32

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19	Multi-omics analysis reveals contextual tumor suppressive and oncogenic gene modules within the acute hypoxic response. Nature Communications, 2021, 12, 1375.	12.8	31
20	A DR4:tBID axis drives the p53 apoptotic response by promoting oligomerization of poised BAX. EMBO Journal, 2012, 31, 1266-1278.	7.8	29
21	SIX2 Mediates Late-Stage Metastasis via Direct Regulation of <i>SOX2</i> and Induction of a Cancer Stem Cell Program. Cancer Research, 2019, 79, 720-734.	0.9	29
22	Mutual Exclusivity of MED12/MED12L, MED13/13L, and CDK8/19 Paralogs Revealed within the CDK-Mediator Kinase Module. Journal of Proteomics and Bioinformatics, 2013, 01, .	0.4	25
23	Identification of a Small-Molecule Inhibitor That Disrupts the SIX1/EYA2 Complex, EMT, and Metastasis. Cancer Research, 2020, 80, 2689-2702.	0.9	24
24	Lessons on transcriptional control from the serum response network. Current Opinion in Genetics and Development, 2011, 21, 160-166.	3.3	22
25	Nutlin-Induced Apoptosis Is Specified by a Translation Program Regulated by PCBP2 and DHX30. Cell Reports, 2020, 30, 4355-4369.e6.	6.4	18
26	NPM and BRG1 Mediate Transcriptional Resistance to Retinoic Acid in Acute Promyelocytic Leukemia. Cell Reports, 2016, 14, 2938-2949.	6.4	13
27	Precocious clonal hematopoiesis in Down syndrome is accompanied by immune dysregulation. Blood Advances, 2021, 5, 1791-1796.	5.2	13
28	ERK phosphorylation of MED14 in promoter complexes during mitogen-induced gene activation by Elk-1. Nucleic Acids Research, 2013, 41, 10241-10253.	14.5	10
29	Role of glutamine synthetase in phenazine antibiotic production by Pantoea agglomerans Eh1087. Canadian Journal of Microbiology, 2004, 50, 877-881.	1.7	5
30	Global Analyses to Identify Direct Transcriptional Targets of p53. Methods in Molecular Biology, 2021, 2267, 19-56.	0.9	3