

# Nathaniel A Hathaway

## List of Publications by Year in descending order

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Version: 2024-02-01

29  
papers

2,797  
citations

516681

16  
h-index

477281

29  
g-index

30  
all docs

30  
docs citations

30  
times ranked

4297  
citing authors

#	ARTICLE	IF	CITATIONS
1	Loss of the Lkb1 tumour suppressor provokes intestinal polyposis but resistance to transformation. <i>Nature</i> , 2002, 419, 162-167.	27.8	390
2	Quantitative analysis of in vitro ubiquitinated cyclin B1 reveals complex chain topology. <i>Nature Cell Biology</i> , 2006, 8, 700-710.	10.3	390
3	Dynamics and Memory of Heterochromatin in Living Cells. <i>Cell</i> , 2012, 149, 1447-1460.	28.9	381
4	Deubiquitinating Enzyme Ubp6 Functions Noncatalytically to Delay Proteasomal Degradation. <i>Cell</i> , 2006, 127, 99-111.	28.9	316
5	Pharmacologic Inhibition of the Anaphase-Promoting Complex Induces A Spindle Checkpoint-Dependent Mitotic Arrest in the Absence of Spindle Damage. <i>Cancer Cell</i> , 2010, 18, 382-395.	16.8	285
6	Ubiquitin Chains Are Remodeled at the Proteasome by Opposing Ubiquitin Ligase and Deubiquitinating Activities. <i>Cell</i> , 2006, 127, 1401-1413.	28.9	280
7	Stability of the proteasome can be regulated allosterically through engagement of its proteolytic active sites. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 1180-1188.	8.2	140
8	APC/C-mediated multiple monoubiquitylation provides an alternative degradation signal for cyclin B1. <i>Nature Cell Biology</i> , 2012, 14, 168-176.	10.3	125
9	Histone hyperacetylation disrupts core gene regulatory architecture in rhabdomyosarcoma. <i>Nature Genetics</i> , 2019, 51, 1714-1722.	21.4	113
10	Nucleosome Turnover Regulates Histone Methylation Patterns over the Genome. <i>Molecular Cell</i> , 2019, 73, 61-72.e3.	9.7	69
11	Comprehensive nucleosome interactome screen establishes fundamental principles of nucleosome binding. <i>Nucleic Acids Research</i> , 2020, 48, 9415-9432.	14.5	67
12	Dose-dependent activation of gene expression is achieved using CRISPR and small molecules that recruit endogenous chromatin machinery. <i>Nature Biotechnology</i> , 2020, 38, 50-55.	17.5	51
13	LSH mediates gene repression through macroH2A deposition. <i>Nature Communications</i> , 2020, 11, 5647.	12.8	35
14	Ring finger protein 121 is a potent regulator of adeno-associated viral genome transcription. <i>PLoS Pathogens</i> , 2019, 15, e1007988.	4.7	22
15	Deferiprone: Pan-selective Histone Lysine Demethylase Inhibition Activity and Structure Activity Relationship Study. <i>Scientific Reports</i> , 2019, 9, 4802.	3.3	20
16	Targeted Gene Repression Using Novel Bifunctional Molecules to Harness Endogenous Histone Deacetylation Activity. <i>ACS Synthetic Biology</i> , 2018, 7, 38-45.	3.8	17
17	Epigenetic Control of a Local Chromatin Landscape. <i>International Journal of Molecular Sciences</i> , 2020, 21, 943.	4.1	15
18	A Peptidomimetic Ligand Targeting the Chromodomain of MPP8 Reveals HRP2's Association with the HUSH Complex. <i>ACS Chemical Biology</i> , 2021, 16, 1721-1736.	3.4	12

#	ARTICLE	IF	CITATIONS
19	Design, Implementation, and Outcomes of a Three-week Pharmacy Bridging Course. <i>American Journal of Pharmaceutical Education</i> , 2017, 81, 6313.	2.1	12
20	Cavitation Enhancement Increases the Efficiency and Consistency of Chromatin Fragmentation from Fixed Cells for Downstream Quantitative Applications. <i>Biochemistry</i> , 2018, 57, 2756-2761.	2.5	11
21	Tethering of Lsh at the Oct4 locus promotes gene repression associated with epigenetic changes. <i>Epigenetics</i> , 2018, 13, 173-181.	2.7	10
22	Epigenetic roots of immunologic disease and new methods for examining chromatin regulatory pathways. <i>Immunology and Cell Biology</i> , 2015, 93, 261-270.	2.3	7
23	Dissecting cell biology with chemical scalpels. <i>Current Opinion in Cell Biology</i> , 2005, 17, 12-19.	5.4	5
24	Chemical screen for epigenetic barriers to single allele activation of Oct4. <i>Stem Cell Research</i> , 2019, 38, 101470.	0.7	5
25	Report and Application of a Tool Compound Data Set. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2699-2706.	5.4	4
26	Repressing Gene Transcription by Redirecting Cellular Machinery with Chemical Epigenetic Modifiers. <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	4
27	Contribution of promoter DNA sequence to heterochromatin formation velocity and memory of gene repression in mouse embryo fibroblasts. <i>PLoS ONE</i> , 2019, 14, e0217699.	2.5	4
28	Pathway-Based High-Throughput Chemical Screen Identifies Compounds That Decouple Heterochromatin Transformations. <i>SLAS Discovery</i> , 2019, 24, 802-816.	2.7	3
29	Bioorthogonal Chemical Epigenetic Modifiers Enable Dose-Dependent CRISPR Targeted Gene Activation in Mammalian Cells. <i>ACS Synthetic Biology</i> , 2022, 11, 1397-1407.	3.8	3