Marian Kalocsay

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Quantitative Proteomics of the Cancer Cell Line Encyclopedia. Cell, 2020, 180, 387-402.e16.	28.9	596
2	Mobilization of Processed, Membrane-Tethered SPT23 Transcription Factor by CDC48UFD1/NPL4, a Ubiquitin-Selective Chaperone. Cell, 2001, 107, 667-677.	28.9	421
3	Chromosome-wide Rad51 Spreading and SUMO-H2A.Z-Dependent Chromosome Fixation in Response to a Persistent DNA Double-Strand Break. Molecular Cell, 2009, 33, 335-343.	9.7	317
4	Multidimensional Tracking of GPCR Signaling via Peroxidase-Catalyzed Proximity Labeling. Cell, 2017, 169, 338-349.e11.	28.9	221
5	Mechanism of adrenergic CaV1.2 stimulation revealed by proximity proteomics. Nature, 2020, 577, 695-700.	27.8	163
6	Targeting immunosuppressive macrophages overcomes PARP inhibitor resistance in BRCA1-associated triple-negative breast cancer. Nature Cancer, 2021, 2, 66-82.	13.2	126
7	Nudt21 Controls Cell Fate by Connecting Alternative Polyadenylation to Chromatin Signaling. Cell, 2018, 172, 106-120.e21.	28.9	123
8	Expedited mapping of the ligandable proteome using fully functionalized enantiomeric probe pairs. Nature Chemistry, 2019, 11, 1113-1123.	13.6	93
9	Monitoring Homology Search during DNA Double-Strand Break Repair InÂVivo. Molecular Cell, 2013, 50, 261-272.	9.7	74
10	Structure of the human metapneumovirus polymerase phosphoprotein complex. Nature, 2020, 577, 275-279.	27.8	72
11	Discovery and resistance mechanism of a selective CDK12 degrader. Nature Chemical Biology, 2021, 17, 675-683.	8.0	69
12	Automethylation-induced conformational switch in Clr4 (Suv39h) maintains epigenetic stability. Nature, 2018, 560, 504-508.	27.8	59
13	Time-resolved proteomics profiling of the ciliary Hedgehog response. Journal of Cell Biology, 2021, 220, .	5.2	50
14	Genome-associated RNA Polymerase II Includes the Dissociable Rpb4/7 Subcomplex. Journal of Biological Chemistry, 2008, 283, 26423-26427.	3.4	40
15	RNA helicase DDX21 mediates nucleotide stress responses in neural crest and melanoma cells. Nature Cell Biology, 2020, 22, 372-379.	10.3	37
16	A conserved RNA degradation complex required for spreading and epigenetic inheritance of heterochromatin. ELife, 2020, 9, .	6.0	31
17	YAP regulates cell size and growth dynamics via non-cell autonomous mediators. ELife, 2020, 9, .	6.0	28
18	Discovery of Covalent CDK14 Inhibitors with Pan-TAIRE Family Specificity. Cell Chemical Biology, 2019, 26, 804-817.e12.	5.2	19

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19	Rixosomal RNA degradation contributes to silencing of Polycomb target genes. Nature, 2022, 604, 167-174.	27.8	18
20	Orphan nuclear receptor COUPâ€TFII enhances myofibroblast glycolysis leading to kidney fibrosis. EMBO Reports, 2021, 22, e51169.	4.5	16
21	Timeâ€series transcriptomics and proteomics reveal alternative modes to decode p53 oscillations. Molecular Systems Biology, 2022, 18, e10588.	7.2	16
22	APEX Peroxidase-Catalyzed Proximity Labeling and Multiplexed Quantitative Proteomics. Methods in Molecular Biology, 2019, 2008, 41-55.	0.9	15
23	Detecting Cardiovascular Protein-Protein Interactions by Proximity Proteomics. Circulation Research, 2022, 130, 273-287.	4.5	11
24	Impact of Zika virus on the human type I interferon osteoimmune response. Cytokine, 2021, 137, 155342.	3.2	4
25	860â€Targeting immunosuppressive macrophages overcomes PARP-inhibitor resistance in BRCA1-associated triple-negative breast cancer. , 2020, , .		1
26	Prostaglandin E2 Stimulates CREB-Mediated Modification of Histone Variant Nucleosomes at Enhancers to Promote Hematopoietic Stem Cell Fate. Blood, 2018, 132, 530-530.	1.4	1
27	Abstract P5-04-01: PARP inhibition modulates the infiltration, phenotype and function of tumor-associated macrophages (TAMs) in BRCA-associated breast cancer and can be augmented by harnessing the anti-tumor potential of TAMs. , 2020, , .		0