## Kathi Zarnack

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Disruption of Prostaglandin E2 Signaling in Cancer-Associated Fibroblasts Limits Mammary Carcinoma Growth but Promotes Metastasis. Cancer Research, 2022, 82, 1380-1395.	0.9	10
2	Functional RNA Dynamics Are Progressively Governed by RNA Destabilization during the Adaptation to Chronic Hypoxia. International Journal of Molecular Sciences, 2022, 23, 5824.	4.1	3
3	SRSF3 and SRSF7 modulate 3′UTR length through suppression or activation of proximal polyadenylation sites and regulation of CFIm levels. Genome Biology, 2021, 22, 82.	8.8	30
4	Bioinformatics in theory and application– highlights of the 36th German Conference on Bioinformatics. Biological Chemistry, 2021, 402, 869-870.	2.5	1
5	Direct long-read RNA sequencing identifies a subset of questionable exitrons likely arising from reverse transcription artifacts. Genome Biology, 2021, 22, 190.	8.8	20
6	Deep and accurate detection of m6A RNA modifications using miCLIP2 and m6Aboost machine learning. Nucleic Acids Research, 2021, 49, e92-e92.	14.5	50
7	The RNA-binding profile of the splicing factor SRSF6 in immortalized human pancreatic β-cells. Life Science Alliance, 2021, 4, e202000825.	2.8	14
8	iCLIP data analysis: A complete pipeline from sequencing reads to RBP binding sites. Methods, 2020, 178, 49-62.	3.8	45
9	Exon Definition Facilitates Reliable Control of Alternative Splicing in the RON Proto-Oncogene. Biophysical Journal, 2020, 118, 2027-2041.	O.5	7
10	Dynamic mRNP Remodeling in Response to Internal and External Stimuli. Biomolecules, 2020, 10, 1310.	4.0	16
11	An autoinhibitory intramolecular interaction proof-reads RNA recognition by the essential splicing factor U2AF2. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 7140-7149.	7.1	25
12	Muscleblind-like 2 controls the hypoxia response of cancer cells. Rna, 2020, 26, 648-663.	3.5	19
13	Makorin 1 controls embryonic patterning by alleviating Bruno1-mediated repression of oskar translation. PLoS Genetics, 2020, 16, e1008581.	3.5	11
14	Polypyrimidine tract-binding proteins are essential for B cell development. ELife, 2020, 9, .	6.0	25
15	The RNA-binding ubiquitin ligase MKRN1 functions in ribosome-associated quality control of poly(A) translation. Genome Biology, 2019, 20, 216.	8.8	29
16	Phenotypic Plasticity of Fibroblasts during Mammary Carcinoma Development. International Journal of Molecular Sciences, 2019, 20, 4438.	4.1	19
17	A combined computational pipeline to detect circular RNAs in human cancer cells under hypoxic stress. Journal of Molecular Cell Biology, 2019, 11, 829-844.	3.3	21
18	uORF-Tools—Workflow for the determination of translation-regulatory upstream open reading frames. PLoS ONF, 2019, 14, e0222459	2.5	7

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19	The key protein of endosomal mRNP transport Rrm4 binds translational landmark sites of cargo mRNAs. EMBO Reports, 2019, 20, .	4.5	38
20	Membrane-Associated RNA-Binding Proteins Orchestrate Organelle-Coupled Translation. Trends in Cell Biology, 2019, 29, 178-188.	7.9	60
21	In vitro iCLIP-based modeling uncovers how the splicing factor U2AF2 relies on regulation by cofactors. Genome Research, 2018, 28, 699-713.	5.5	62
22	High-Throughput Screens for <i>cis</i> -Acting RNA Sequence Elements That Promote Nuclear Retention. Biochemistry, 2018, 57, 3542-3543.	2.5	1
23	Decoding a cancer-relevant splicing decision in the RON proto-oncogene using high-throughput mutagenesis. Nature Communications, 2018, 9, 3315.	12.8	46
24	Insights into the design and interpretation of iCLIP experiments. Genome Biology, 2017, 18, 7.	8.8	73
25	Cellular differentiation state modulates the mRNA export activity of SR proteins. Journal of Cell Biology, 2017, 216, 1993-2009.	5.2	53
26	Differential Binding of Mitochondrial Transcripts by MRB8170 and MRB4160 Regulates Distinct Editing Fates of Mitochondrial mRNA in Trypanosomes. MBio, 2017, 8, .	4.1	17
27	Interaction profiling of RNA-binding ubiquitin ligases reveals a link between posttranscriptional regulation and the ubiquitin system. Scientific Reports, 2017, 7, 16582.	3.3	17
28	Splicing repression allows the gradual emergence of new Alu-exons in primate evolution. ELife, 2016, 5, .	6.0	57
29	Recognition of distinct RNA motifs by the clustered CCCH zinc fingers of neuronal protein Unkempt. Nature Structural and Molecular Biology, 2016, 23, 16-23.	8.2	23
30	SR proteins are NXF1 adaptors that link alternative RNA processing to mRNA export. Genes and Development, 2016, 30, 553-566.	5.9	242
31	The RNA-binding protein HuR is essential for the B cell antibody response. Nature Immunology, 2015, 16, 415-425.	14.5	125
32	Control of a neuronal morphology program by an RNA-binding zinc finger protein, Unkempt. Genes and Development, 2015, 29, 501-512.	5.9	35
33	Direct Competition between hnRNP C and U2AF65 Protects the Transcriptome from the Exonization of Alu Elements. Cell, 2013, 152, 453-466.	28.9	398
34	Protein–RNA interactions: new genomic technologies and perspectives. Nature Reviews Genetics, 2012, 13, 77-83.	16.3	482
35	The RNA-Binding Protein Rrm4 is Essential for Efficient Secretion of Endochitinase Cts1. Molecular and Cellular Proteomics, 2011, 10, M111.011213.	3.8	48
36	iCLIP reveals the function of hnRNP particles in splicing at individual nucleotide resolution. Nature Structural and Molecular Biology, 2010, 17, 909-915.	8.2	1,026

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37	iCLIP Predicts the Dual Splicing Effects of TIA-RNA Interactions. PLoS Biology, 2010, 8, e1000530.	5.6	226
38	Tandem KH domains of Khd4 recognize AUACCC and are essential for regulation of morphology as well as pathogenicity in <i>Ustilago maydis</i> . Rna, 2009, 15, 2206-2218.	3.5	24
39	The fungal RNA-binding protein Rrm4 mediates long-distance transport of ubi1 and rho3 mRNAs. EMBO Journal, 2009, 28, 1855-1866.	7.8	90
40	Protein–RNA interactions: new genomic technologies and perspectives. , 0, .		1