

CITATION REPORT

List of articles citing

RNA interactome capture in yeast

DOI: 10.1016/j.ymeth.2016.12.008
Methods, 2017, 118-119, 82-92.

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
18	Protein-RNA: Structure function and recognition. <i>Methods</i> , 2017 , 118-119, 1-2	4.6	
17	Kinetic CRAC uncovers a role for Nab3 in determining gene expression profiles during stress. <i>Nature Communications</i> , 2017 , 8, 12	17.4	49
16	Purification of cross-linked RNA-protein complexes by phenol-toluol extraction. <i>Nature Communications</i> , 2019 , 10, 990	17.4	94
15	Methods to study RNA-protein interactions. <i>Nature Methods</i> , 2019 , 16, 225-234	21.6	126
14	Fast and unbiased purification of RNA-protein complexes after UV cross-linking. <i>Methods</i> , 2020 , 178, 72-82	4.6	13
13	Single and Combined Methods to Specifically or Bulk-Purify RNA-Protein Complexes. <i>Biomolecules</i> , 2020 , 10,	5.9	6
12	Discovering the RNA-Binding Proteome of Plant Leaves with an Improved RNA Interactome Capture Method. <i>Biomolecules</i> , 2020 , 10,	5.9	22
11	The physical basis and practical consequences of biological promiscuity. <i>Physical Biology</i> , 2020 ,	3	5
10	easyCLIP analysis of RNA-protein interactions incorporating absolute quantification. <i>Nature Communications</i> , 2021 , 12, 1569	17.4	2
9	Searching for regulators that interact with BELL1 transcription factor and control the legume-rhizobial symbiosis development. <i>Ecological Genetics</i> , 2021 , 19, 37-45	0.5	
8	Surveying the global landscape of post-transcriptional regulators.		
7	Conserved Methyltransferase Spb1 Targets mRNAs for Regulated Modification with 2'-O-Methyl Ribose.		11
6	Purification of Cross-linked RNA-Protein Complexes by Phenol-Toluol Extraction.		3
5	Fast and unbiased purification of RNA-protein complexes after UV cross-linking.		1
4	Probing the RNA-Binding Proteome from Yeast to Man: Major Advances and Challenges. <i>Methods in Molecular Biology</i> , 2019 , 2049, 213-231	1.4	1
3	Shaping the Innate Immune Response Through Post-Transcriptional Regulation of Gene Expression Mediated by RNA-Binding Proteins.. <i>Frontiers in Immunology</i> , 2021 , 12, 796012	8.4	1
2	The RNA-bound proteome of MRSA reveals post-transcriptional roles for helix-turn-helix DNA-binding and Rossmann-fold proteins. <i>Nature Communications</i> , 2022 , 13,	17.4	1

1 RNA Brotein Interactomics. **2022**, 271-291

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