Justin T Roberts

List of Publications by Year in Descending Order

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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.

17	309	8	17
papers	citations	h-index	g-index
20	417	5.7	3.35
ext. papers	ext. citations	avg, IF	L-index

#	Paper	IF	Citations
17	Identification of mA residues at single-nucleotide resolution using eCLIP and an accessible custom analysis pipeline. <i>Rna</i> , 2021 , 27, 527-541	5.8	3
16	Establishing RNA-RNA interactions remodels lncRNA structure and promotes PRC2 activity. <i>Science Advances</i> , 2021 , 7,	14.3	7
15	A Positive Feedback Loop Between TGFIand Androgen Receptor Supports Triple-negative Breast Cancer Anoikis Resistance. <i>Endocrinology</i> , 2021 , 162,	4.8	6
14	RNA matchmaking in chromatin regulation. <i>Biochemical Society Transactions</i> , 2020 , 48, 2467-2481	5.1	4
13	Characterization of novel small RNAs (sRNAs) contributing to the desiccation response of serovar Typhimurium. <i>RNA Biology</i> , 2019 , 16, 1643-1657	4.8	4
12	ADAR Mediated RNA Editing Modulates MicroRNA Targeting in Human Breast Cancer. <i>Processes</i> , 2018 , 6,	2.9	8
11	Global profiling of hnRNP A2/B1-RNA binding on chromatin highlights LncRNA interactions. <i>RNA Biology</i> , 2018 , 15, 901-913	4.8	17
10	Computational Prediction of MicroRNA Target Genes, Target Prediction Databases, and Web Resources. <i>Methods in Molecular Biology</i> , 2017 , 1617, 109-122	1.4	23
9	Human snoRNA-93 is processed into a microRNA-like RNA that promotes breast cancer cell invasion. <i>Npj Breast Cancer</i> , 2017 , 3, 25	7.8	31
8	Novel small RNA (sRNA) landscape of the starvation-stress response transcriptome of Salmonella enterica serovar typhimurium. <i>RNA Biology</i> , 2016 , 13, 331-42	4.8	14
7	An oxidative DNA "damage" and repair mechanism localized in the VEGF promoter is important for hypoxia-induced VEGF mRNA expression. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2015 , 309, L1367-75	5.8	94
6	Burgeoning evidence indicates that microRNAs were initially formed from transposable element sequences. <i>Mobile Genetic Elements</i> , 2014 , 4, e29255		62
5	Continuing analysis of microRNA origins: Formation from transposable element insertions and noncoding RNA mutations. <i>Mobile Genetic Elements</i> , 2013 , 3, e27755		31
4	Identification of m6A residues at single-nucleotide resolution using eCLIP and an accessible custom analysis pipeline		1
3	RNA matchmaking remodels lncRNA structure and promotes PRC2 activity		2
2	A single N6-methyladenosine site in lncRNA HOTAIR regulates its function in breast cancer cells		2
1	Evolutionary Origin of MicroRNAs1-8		