

# Marianne C Kramer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5124143/publications.pdf>

Version: 2024-02-01

10  
papers

945  
citations

1163117

8  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

1810  
citing authors

#	ARTICLE	IF	CITATIONS
1	Combinatorial control of <i>Drosophila</i> circular RNA expression by intronic repeats, hnRNPs, and SR proteins. <i>Genes and Development</i> , 2015, 29, 2168-2182.	5.9	419
2	Unusual maintenance of X chromosome inactivation predisposes female lymphocytes for increased expression from the inactive X. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E2029-38.	7.1	210
3	N6-Methyladenosine Inhibits Local Ribonucleolytic Cleavage to Stabilize mRNAs in Arabidopsis. <i>Cell Reports</i> , 2018, 25, 1146-1157.e3.	6.4	175
4	N <sup>6</sup> -methyladenosine and RNA secondary structure affect transcript stability and protein abundance during systemic salt stress in Arabidopsis. <i>Plant Direct</i> , 2020, 4, e00239.	1.9	41
5	Messenger RNA 5' NAD <sup>+</sup> Capping Is a Dynamic Regulatory Epitranscriptome Mark That Is Required for Proper Response to Abscisic Acid in Arabidopsis. <i>Developmental Cell</i> , 2021, 56, 125-140.e6.	7.0	40
6	The nucleotides they are a-changing™: function of RNA binding proteins in post-transcriptional messenger RNA editing and modification in Arabidopsis. <i>Current Opinion in Plant Biology</i> , 2018, 45, 88-95.	7.1	20
7	RNA structure, binding, and coordination in <i>Arabidopsis</i> . <i>Wiley Interdisciplinary Reviews RNA</i> , 2017, 8, e1426.	6.4	14
8	Does RNA secondary structure drive translation or vice versa?. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 641-643.	8.2	13
9	Data-Independent Acquisition for the Detection of Mononucleoside RNA Modifications by Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 885-893.	2.8	7
10	Using Protein Interaction Profile Sequencing (PIP-seq) to Identify RNA Secondary Structure and RNA-Protein Interaction Sites of Long Noncoding RNAs in Plants. <i>Methods in Molecular Biology</i> , 2019, 1933, 343-361.	0.9	6