Martyna O Urbanek

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3543159/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Nuclear speckles: molecular organization, biological function and role in disease. Nucleic Acids Research, 2017, 45, 10350-10368.	14.5	334
2	Small RNA Detection by in Situ Hybridization Methods. International Journal of Molecular Sciences, 2015, 16, 13259-13286.	4.1	78
3	RNA imaging in living cells $\hat{a} \in $ methods and applications. RNA Biology, 2014, 11, 1083-1095.	3.1	77
4	Triplet repeats in transcripts: structural insights into RNA toxicity. Biological Chemistry, 2012, 393, 1299-1315.	2.5	58
5	Somatic Mutations in miRNA Genes in Lung Cancer—Potential Functional Consequences of Non-Coding Sequence Variants. Cancers, 2019, 11, 793.	3.7	37
6	Nuclear speckles are detention centers for transcripts containing expanded CAG repeats. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2016, 1862, 1513-1520.	3.8	29
7	Pan-cancer analysis of somatic mutations in miRNA genes. EBioMedicine, 2020, 61, 103051.	6.1	29
8	The Role of the Immune System in Triplet Repeat Expansion Diseases. Mediators of Inflammation, 2015, 2015, 1-11.	3.0	28
9	RNA FISH for detecting expanded repeats in human diseases. Methods, 2016, 98, 115-123.	3.8	28
10	A pan-cancer atlas of somatic mutations in miRNA biogenesis genes. Nucleic Acids Research, 2021, 49, 601-620.	14.5	26
11	Sequence-non-specific effects generated by various types of RNA interference triggers. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 306-314.	1.9	19
12	RAN Translation of the Expanded CAG Repeats in the SCA3 Disease Context. Journal of Molecular Biology, 2020, 432, 166699.	4.2	17
13	Reduction of Huntington's Disease RNA Foci by CAG Repeat-Targeting Reagents. Frontiers in Cellular Neuroscience, 2017, 11, 82.	3.7	15
14	miRNAmotif—A Tool for the Prediction of Pre-miRNA–Protein Interactions. International Journal of Molecular Sciences, 2018, 19, 4075.	4.1	10
15	A Comprehensive Transcriptome Analysis Identifies FXN and BDNF as Novel Targets of miRNAs in Friedreich's Ataxia Patients. Molecular Neurobiology, 2020, 57, 2639-2653.	4.0	9
16	Discriminating RNA variants with single-molecule allele-specific FISH. Mutation Research - Reviews in Mutation Research, 2017, 773, 230-241.	5.5	7
17	miRMut: Annotation of mutations in miRNA genes from human whole-exome or whole-genome sequencing. STAR Protocols, 2022, 3, 101023.	1.2	4
18	2D and 3D FISH of expanded repeat RNAs in human lymphoblasts. Methods, 2017, 120, 49-57.	3.8	3

#	Article	IF	CITATIONS
19	Profile of Basal Cell Carcinoma Mutations and Copy Number Alterations - Focus on Gene-Associated Noncoding Variants. Frontiers in Oncology, 2021, 11, 752579.	2.8	1