## Monika Gullerova

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4555804/publications.pdf

Version: 2024-02-01

26 papers 1,056 citations

643344 15 h-index 25 g-index

26 all docs

26 docs citations

26 times ranked 1760 citing authors

#	Article	IF	Citations
1	Cohesin Complex Promotes Transcriptional Termination between Convergent Genes in S. pombe. Cell, 2008, 132, 983-995.	13.5	186
2	Human nuclear Dicer restricts the deleterious accumulation of endogenous double-stranded RNA. Nature Structural and Molecular Biology, 2014, 21, 552-559.	3.6	95
3	Swiss army knives: non-canonical functions of nuclear Drosha and Dicer. Nature Reviews Molecular Cell Biology, 2015, 16, 417-430.	16.1	88
4	Noncanonical functions of micro <scp>RNA</scp> pathway enzymes – Drosha, <scp>DGCR</scp> 8, Dicer and Ago proteins. FEBS Letters, 2018, 592, 2973-2986.	1.3	88
5	Nuclear phosphorylated Dicer processes double-stranded RNA in response to DNA damage. Journal of Cell Biology, 2017, 216, 2373-2389.	2.3	73
6	Subcellular RNA profiling links splicing and nuclear DICER1 to alternative cleavage and polyadenylation. Genome Research, 2016, 26, 24-35.	2.4	70
7	Tyrosine kinase c-Abl couples RNA polymerase II transcription to DNA double-strand breaks. Nucleic Acids Research, 2019, 47, 3467-3484.	6.5	68
8	AtCyp59 is a multidomain cyclophilin from Arabidopsis thaliana that interacts with SR proteins and the C-terminal domain of the RNA polymerase II. Rna, 2006, 12, 631-643.	1.6	61
9	Convergent transcription induces transcriptional gene silencing in fission yeast and mammalian cells. Nature Structural and Molecular Biology, 2012, 19, 1193-1201.	3.6	52
10	Beyond the Trinity of ATM, ATR, and DNA-PK: Multiple Kinases Shape the DNA Damage Response in Concert With RNA Metabolism. Frontiers in Molecular Biosciences, 2019, 6, 61.	1.6	44
11	Genome-wide analysis of poly(A) site selection in <i>Schizosaccharomyces pombe</i> . Rna, 2013, 19, 1617-1631.	1.6	37
12	Dicer dependent tRNA derived small RNAs promote nascent RNA silencing. Nucleic Acids Research, 2022, 50, 1734-1752.	6.5	32
13	Jack of all trades? The versatility of RNA in DNA double-strand break repair. Essays in Biochemistry, 2020, 64, 721-735.	2.1	29
14	Autoregulation of convergent RNAi genes in fission yeast. Genes and Development, 2011, 25, 556-568.	2.7	27
15	Nuclear re-localization of Dicer in primary mouse embryonic fibroblast nuclei following DNA damage. PLoS Genetics, 2018, 14, e1007151.	1.5	23
16	Home and Away: The Role of Non-Coding RNA in Intracellular and Intercellular DNA Damage Response. Genes, 2021, 12, 1475.	1.0	15
17	Rct1, a Nuclear RNA Recognition Motif-Containing Cyclophilin, Regulates Phosphorylation of the RNA Polymerase II C-Terminal Domain. Molecular and Cellular Biology, 2007, 27, 3601-3611.	1.1	14
18	Transcription facilitates sister chromatid cohesion on chromosomal arms. Nucleic Acids Research, 2016, 44, 6676-6692.	6.5	13

#	Article	IF	CITATIONS
19	Sweet Melody or Jazz? Transcription Around DNA Double-Strand Breaks. Frontiers in Molecular Biosciences, 2021, 8, 655786.	1.6	12
20	The Methylation Game: Epigenetic and Epitranscriptomic Dynamics of 5-Methylcytosine. Frontiers in Cell and Developmental Biology, 2022, $10$ , .	1.8	10
21	Proximity Ligation Assay for Detection of R-Loop Complexes upon DNA Damage. Methods in Molecular Biology, 2022, , 289-303.	0.4	6
22	Gene Silencing CUTs Both Ways. Cell, 2007, 131, 649-651.	13.5	5
23	Long Non-coding RNA. , 2015, , 83-108.		4
24	p19-Mediated Enrichment and Detection of siRNAs. Methods in Molecular Biology, 2014, 1173, 99-111.	0.4	2
25	Silencing in trans: position matters in fission yeast. EMBO Reports, 2010, 11, 145-146.	2.0	1
26	Understanding non-coding DNA regions in yeast. Biochemical Society Transactions, 2013, 41, 1654-1659.	1.6	1