

# Laure Teyssset

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5943073/publications.pdf>

Version: 2024-02-01

12  
papers

662  
citations

932766

10  
h-index

1281420

11  
g-index

16  
all docs

16  
docs citations

16  
times ranked

719  
citing authors

#	ARTICLE	IF	CITATIONS
1	Paramutation in <i>Drosophila</i> linked to emergence of a piRNA-producing locus. <i>Nature</i> , 2012, 490, 112-115.	13.7	216
2	RNA 2â€™-O-Methylation (Nm) Modification in Human Diseases. <i>Genes</i> , 2019, 10, 117.	1.0	126
3	Telomeric Trans-Silencing: An Epigenetic Repression Combining RNA Silencing and Heterochromatin Formation. <i>PLoS Genetics</i> , 2007, 3, e158.	1.5	93
4	Paramutation in <i>Drosophila</i> Requires Both Nuclear and Cytoplasmic Actors of the piRNA Pathway and Induces Cis-spreading of piRNA Production. <i>Genetics</i> , 2015, 201, 1381-1396.	1.2	43
5	The Epigenetic Trans-Silencing Effect in <i>Drosophila</i> Involves Maternally-Transmitted Small RNAs Whose Production Depends on the piRNA Pathway and HP1. <i>PLoS ONE</i> , 2010, 5, e11032.	1.1	42
6	tRNA 2â€™-O-methylation by a duo of TRM7/FTSJ1 proteins modulates small RNA silencing in <i>Drosophila</i> . <i>Nucleic Acids Research</i> , 2020, 48, 2050-2072.	6.5	30
7	Short and long-term evolutionary dynamics of subtelomeric piRNA clusters in <i>Drosophila</i> . <i>DNA Research</i> , 2017, 24, 459-472.	1.5	28
8	Environmentally-Induced Transgenerational Epigenetic Inheritance: Implication of PIWI Interacting RNAs. <i>Cells</i> , 2019, 8, 1108.	1.8	27
9	Environmentally-induced epigenetic conversion of a piRNA cluster. <i>ELife</i> , 2019, 8, .	2.8	26
10	Telomeric Trans-Silencing in <i>Drosophila melanogaster</i> : Tissue Specificity, Development and Functional Interactions between Non-Homologous Telomeres. <i>PLoS ONE</i> , 2008, 3, e3249.	1.1	19
11	Homology-Dependent Silencing by an Exogenous Sequence in the <i>Drosophila</i> Germline. <i>G3: Genes, Genomes, Genetics</i> , 2012, 2, 331-338.	0.8	11
12	Comparative genomic and transcriptomic analyses of transposable elements in polychaetous annelids highlight LTR retrotransposon diversity and evolution. <i>Mobile DNA</i> , 2021, 12, 24.	1.3	1