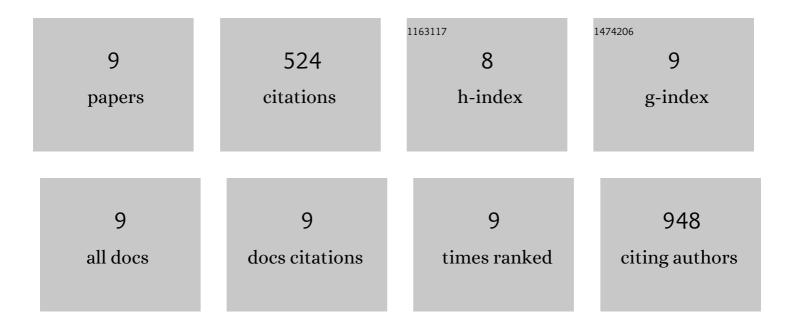
## Tao Wu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11694128/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Enzymatic Characterization of In Vitro Activity of RNA Methyltransferase PCIF1 on DNA. Biochemistry, 2022, 61, 1005-1013.	2.5	6
2	Mammalian ALKBH1 serves as an N6-mA demethylase of unpairing DNA. Cell Research, 2020, 30, 197-210.	12.0	71
3	Extensive Nuclear Reprogramming Underlies Lineage Conversion into Functional Trophoblast Stem-like Cells. Cell Stem Cell, 2015, 17, 543-556.	11.1	80
4	The Developmental Potential of iPSCs Is Greatly Influenced by Reprogramming Factor Selection. Cell Stem Cell, 2014, 15, 295-309.	11.1	137
5	Histone Variant H2A.X Deposition Pattern Serves as a Functional Epigenetic Mark for Distinguishing the Developmental Potentials of iPSCs. Cell Stem Cell, 2014, 15, 281-294.	11.1	58
6	Genome-scale identification of Caenorhabditis elegans regulatory elements by tiling-array mapping of DNase I hypersensitive sites. BMC Genomics, 2009, 10, 92.	2.8	13
7	Mapping the <i>C. elegans</i> noncoding transcriptome with a whole-genome tiling microarray. Genome Research, 2007, 17, 1471-1477.	5.5	53
8	Identifying Hfq-binding small RNA targets in Escherichia coli. Biochemical and Biophysical Research Communications, 2006, 343, 950-955.	2.1	13
9	NPInter: the noncoding RNAs and protein related biomacromolecules interaction database. Nucleic Acids Research, 2006, 34, D150-D152.	14.5	93