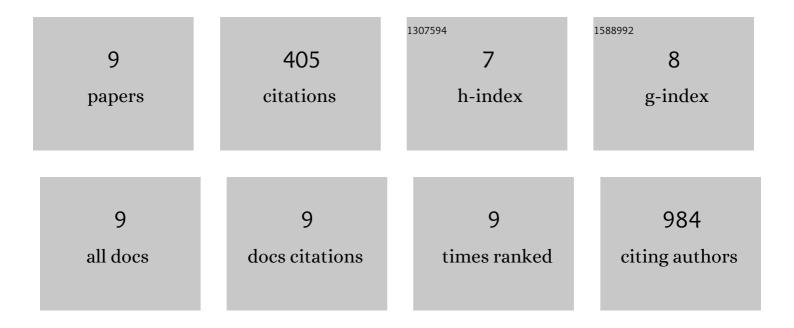
## Wenguang Shao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7614701/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The SysteMHC Atlas project. Nucleic Acids Research, 2018, 46, D1237-D1247.	14.5	119
2	Systematic proteome and proteostasis profiling in human Trisomy 21 fibroblast cells. Nature Communications, 2017, 8, 1212.	12.8	112
3	DPHL: A DIA Pan-human Protein Mass Spectrometry Library for Robust Biomarker Discovery. Genomics, Proteomics and Bioinformatics, 2020, 18, 104-119.	6.9	51
4	A tissue-based draft map of the murine MHC class I immunopeptidome. Scientific Data, 2018, 5, 180157.	5.3	45
5	Comparative analysis of mRNA and protein degradation in prostate tissues indicates high stability of proteins. Nature Communications, 2019, 10, 2524.	12.8	35
6	Refining similarity scoring to enable decoyâ€free validation in spectral library searching. Proteomics, 2013, 13, 3273-3283.	2.2	23
7	The SysteMHC Atlas: a Computational Pipeline, a Website, and a Data Repository for Immunopeptidomic Analyses. Methods in Molecular Biology, 2020, 2120, 173-181.	0.9	10
8	Differential Diagnosis of Preeclampsia Based on Urine Peptidome Features Revealed by High Resolution Mass Spectrometry. Diagnostics, 2020, 10, 1039.	2.6	9
9	PCfun: a hybrid computational framework for systematic characterization of protein complex function. Briefings in Bioinformatics, 0, , .	6.5	1