

# Timothy R Keiffer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1007335/publications.pdf>

Version: 2024-02-01

10  
papers

588  
citations

1307594

7  
h-index

1474206

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

861  
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein Stability and Surface Electrostatics: A Charged Relationship. <i>Biochemistry</i> , 2006, 45, 2761-2766.	2.5	285
2	Incoming human papillomavirus type 16 genome resides in a vesicular compartment throughout mitosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 6289-6294.	7.1	81
3	Topography of the Human Papillomavirus Minor Capsid Protein L2 during Vesicular Trafficking of Infectious Entry. <i>Journal of Virology</i> , 2015, 89, 10442-10452.	3.4	56
4	A new cell culture model to genetically dissect the complete human papillomavirus life cycle. <i>PLoS Pathogens</i> , 2018, 14, e1006846.	4.7	48
5	Human Papillomavirus Major Capsid Protein L1 Remains Associated with the Incoming Viral Genome throughout the Entry Process. <i>Journal of Virology</i> , 2017, 91, .	3.4	45
6	Meprin Metalloproteases Inactivate Interleukin 6. <i>Journal of Biological Chemistry</i> , 2014, 289, 7580-7588.	3.4	36
7	Incoming human papillomavirus 16 genome is lost in PML protein-deficient HaCaT keratinocytes. <i>Cellular Microbiology</i> , 2017, 19, e12708.	2.1	26
8	Recent Advances in Our Understanding of the Infectious Entry Pathway of Human Papillomavirus Type 16. <i>Microorganisms</i> , 2021, 9, 2076.	3.6	7
9	Amino acid substitutions affecting protein dynamics in eglin C do not affect heat capacity change upon unfolding. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 64, 295-300.	2.6	4
10	Pericellular Proteolysis. , 2011, , 75-94.		0