

# Thomas Kaefer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9436012/publications.pdf>

Version: 2024-02-01

7  
papers

501  
citations

1307366

7  
h-index

1719901

7  
g-index

8  
all docs

8  
docs citations

8  
times ranked

791  
citing authors

#	ARTICLE	IF	CITATIONS
1	Unconventional Peptide Presentation by Major Histocompatibility Complex (MHC) Class I Allele HLA-A*02:01. <i>Journal of Biological Chemistry</i> , 2017, 292, 5262-5270.	1.6	57
2	Machine learning reveals a non-canonical mode of peptide binding to <scp>MHC</scp> class II molecules. <i>Immunology</i> , 2017, 152, 255-264.	2.0	27
3	Shared peptide binding of HLA Class I and II alleles associate with cutaneous nevirapine hypersensitivity and identify novel risk alleles. <i>Scientific Reports</i> , 2017, 7, 8653.	1.6	41
4	The Length Distribution of Class I-Restricted T Cell Epitopes Is Determined by Both Peptide Supply and MHC Allele-Specific Binding Preference. <i>Journal of Immunology</i> , 2016, 196, 1480-1487.	0.4	192
5	<i>Toxoplasma gondii</i> peptide ligands open the gate of the HLA class I binding groove. <i>ELife</i> , 2016, 5, .	2.8	88
6	Structural and Functional Characterization of Anti-A33 Antibodies Reveal a Potent Cross-Species Orthopoxviruses Neutralizer. <i>PLoS Pathogens</i> , 2015, 11, e1005148.	2.1	32
7	Using a Combined Computational-Experimental Approach to Predict Antibody-Specific B Cell Epitopes. <i>Structure</i> , 2014, 22, 646-657.	1.6	63