

# Chia-yi Chang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6529594/publications.pdf>

Version: 2024-02-01

11  
papers

321  
citations

1040056

9  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

332  
citing authors

#	ARTICLE	IF	CITATIONS
1	Phylogenetic analysis of classical swine fever virus isolated from Taiwan. <i>Veterinary Microbiology</i> , 2005, 106, 187-193.	1.9	72
2	The challenges of classical swine fever control: Modified live and E2 subunit vaccines. <i>Virus Research</i> , 2014, 179, 1-11.	2.2	69
3	Structures and Functions of Pestivirus Glycoproteins: Not Simply Surface Matters. <i>Viruses</i> , 2015, 7, 3506-3529.	3.3	36
4	The role of porcine teschovirus in causing diseases in endemically infected pigs. <i>Veterinary Microbiology</i> , 2012, 161, 88-95.	1.9	32
5	Antigenic domains analysis of classical swine fever virus E2 glycoprotein by mutagenesis and conformation-dependent monoclonal antibodies. <i>Virus Research</i> , 2010, 149, 183-189.	2.2	29
6	Identification of antigen-specific residues on E2 glycoprotein of classical swine fever virus. <i>Virus Research</i> , 2010, 152, 65-72.	2.2	21
7	Antigenic mimicking with cysteine-based cyclized peptides reveals a previously unknown antigenic determinant on E2 glycoprotein of classical swine fever virus. <i>Virus Research</i> , 2012, 163, 190-196.	2.2	20
8	Identification of conformational epitopes and antigen-specific residues at the D/A domains and the extramembrane C-terminal region of E2 glycoprotein of classical swine fever virus. <i>Virus Research</i> , 2012, 168, 56-63.	2.2	16
9	Multiple models of porcine teschovirus pathogenesis in endemically infected pigs. <i>Veterinary Microbiology</i> , 2014, 168, 69-77.	1.9	12
10	Identification of a Common Conformational Epitope on the Glycoprotein E2 of Classical Swine Fever Virus and Border Disease Virus. <i>Viruses</i> , 2021, 13, 1655.	3.3	9
11	The urinary shedding of porcine teschovirus in endemic field situations. <i>Veterinary Microbiology</i> , 2016, 182, 150-155.	1.9	5