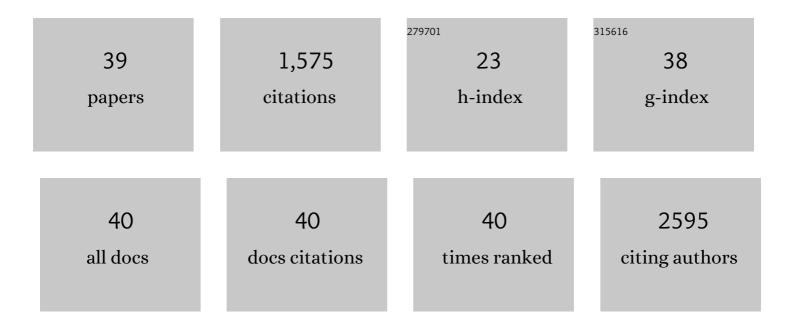
Yan Chai

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	An engineered bispecific human monoclonal antibody against SARS-CoV-2. Nature Immunology, 2022, 23, 423-430.	7.0	38
2	PD-1 N58-Glycosylation-Dependent Binding of Monoclonal Antibody Cemiplimab for Immune Checkpoint Therapy. Frontiers in Immunology, 2022, 13, 826045.	2.2	13
3	A broadly protective antibody that targets the flavivirus NS1 protein. Science, 2021, 371, 190-194.	6.0	66
4	Stability and Structure of Bat Major Histocompatibility Complex Class I with Heterologous β ₂ -Microglobulin. Journal of Visualized Experiments, 2021, , .	0.2	2
5	Protective Zika vaccines engineered to eliminate enhancement of dengue infection via immunodominance switch. Nature Immunology, 2021, 22, 958-968.	7.0	23
6	Structural basis of malarial parasite RIFIN-mediated immune escape against LAIR1. Cell Reports, 2021, 36, 109600.	2.9	7
7	Structural insights into the cis and trans assembly of human trophoblast cell surface antigen 2. IScience, 2021, 24, 103190.	1.9	15
8	Atypical TNF-TNFR superfamily binding interface in the GITR-GITRL complex for TÂcell activation. Cell Reports, 2021, 36, 109734.	2.9	3
9	Peptide Presentations of Marsupial MHC Class I Visualize Immune Features of Lower Mammals Paralleled with Bats. Journal of Immunology, 2021, 207, 2167-2178.	0.4	3
10	Mooring stone-like Arg 114 pulls diverse bulged peptides: first insight into African swine fever virus-derived T cell epitopes presented by swine MHC class I. Journal of Virology, 2021, , JVI0137821.	1.5	3
11	Molecular basis of Coxsackievirus A10 entry using the two-in-one attachment and uncoating receptor KRM1. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 18711-18718.	3.3	18
12	Molecular basis of EphA2 recognition by gHgL from gammaherpesviruses. Nature Communications, 2020, 11, 5964.	5.8	22
13	Structure-Based Modification of an Anti-neuraminidase Human Antibody Restores Protection Efficacy against the Drifted Influenza Virus. MBio, 2020, 11, .	1.8	12
14	Nâ€glycosylation of PDâ€1 promotes binding of camrelizumab. EMBO Reports, 2020, 21, e51444.	2.0	47
15	The identification of a CD47-blocking "hotspot―and design of a CD47/PD-L1 dual-specific antibody with limited hemagglutination. Signal Transduction and Targeted Therapy, 2020, 5, 16.	7.1	29
16	The structural basis of African swine fever virus pA104R binding to DNA and its inhibition by stilbene derivatives. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 11000-11009.	3.3	30
17	Divergent Peptide Presentations of HLA-A*30 Alleles Revealed by Structures With Pathogen Peptides. Frontiers in Immunology, 2019, 10, 1709.	2.2	12
18	Peptide presentation by bat MHC class I provides new insight into the antiviral immunity of bats. PLoS Biology, 2019, 17, e3000436.	2.6	23

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19	Disrupting LILRB4/APOE Interaction by an Efficacious Humanized Antibody Reverses T-cell Suppression and Blocks AML Development. Cancer Immunology Research, 2019, 7, 1244-1257.	1.6	51
20	Molecular Basis of Arthritogenic Alphavirus Receptor MXRA8 Binding to Chikungunya Virus Envelope Protein. Cell, 2019, 177, 1714-1724.e12.	13.5	75
21	The FG Loop of PD-1 Serves as a "Hotspot―for Therapeutic Monoclonal Antibodies in Tumor Immune Checkpoint Therapy. IScience, 2019, 14, 113-124.	1.9	34
22	Neutralization mechanism of human monoclonal antibodies against Rift Valley fever virus. Nature Microbiology, 2019, 4, 1231-1241.	5.9	39
23	Crystal Structure of African Swine Fever Virus dUTPase Reveals a Potential Drug Target. MBio, 2019, 10, .	1.8	24
24	Glycosylation-independent binding of monoclonal antibody toripalimab to FG loop of PD-1 for tumor immune checkpoint therapy. MAbs, 2019, 11, 681-690.	2.6	30
25	Distinct PD-L1 binding characteristics of therapeutic monoclonal antibody durvalumab. Protein and Cell, 2018, 9, 135-139.	4.8	107
26	The Postfusion Structure of the Heartland Virus Gc Glycoprotein Supports Taxonomic Separation of the Bunyaviral Families Phenuiviridae and Hantaviridae. Journal of Virology, 2018, 92, .	1.5	24
27	Limited Cross-Linking of 4-1BB by 4-1BB Ligand and the Agonist Monoclonal Antibody Utomilumab. Cell Reports, 2018, 25, 909-920.e4.	2.9	33
28	An Invariant Arginine in Common with MHC Class II Allows Extension at the C-Terminal End of Peptides Bound to Chicken MHC Class I. Journal of Immunology, 2018, 201, 3084-3095.	0.4	19
29	Structural and functional insights into MCR-2 mediated colistin resistance. Science China Life Sciences, 2018, 61, 1432-1436.	2.3	5
30	Heterosubtypic Protections against Human-Infecting Avian Influenza Viruses Correlate to Biased Cross-T-Cell Responses. MBio, 2018, 9, .	1.8	25
31	An unexpected N-terminal loop in PD-1 dominates binding by nivolumab. Nature Communications, 2017, 8, 14369.	5.8	192
32	The crystal structure of Zika virus <scp>NS</scp> 5 reveals conserved drug targets. EMBO Journal, 2017, 36, 919-933.	3.5	107
33	Protective T Cell Responses Featured by Concordant Recognition of Middle East Respiratory Syndrome Coronavirus–Derived CD8+ T Cell Epitopes and Host MHC. Journal of Immunology, 2017, 198, 873-882.	0.4	42
34	Alternate binding modes of anti-CRISPR viral suppressors AcrF1/2 to Csy surveillance complex revealed by cryo-EM structures. Cell Research, 2017, 27, 853-864.	5.7	64
35	Structures of phlebovirus glycoprotein Gn and identification of a neutralizing antibody epitope. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7564-E7573.	3.3	98
36	Structural basis of anti-PD-L1 monoclonal antibody avelumab for tumor therapy. Cell Research, 2017, 27, 151-153.	5.7	116

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8	37	Two classes of protective antibodies against Pseudorabies virus variant glycoprotein B: Implications for vaccine design. PLoS Pathogens, 2017, 13, e1006777.	2.1	34
ę	38	Remarkably similar CTLA-4 binding properties of therapeutic ipilimumab and tremelimumab antibodies. Oncotarget, 2017, 8, 67129-67139.	0.8	65
8	39	Diversified Anchoring Features the Peptide Presentation of DLA-88*50801: First Structural Insight into Domestic Dog MHC Class I. Journal of Immunology, 2016, 197, 2306-2315.	0.4	25