Andre Azevedo Reis Teixeira

List of Publications by Year
in descending order

12	179	1305906	1336881
papers	citations	8	12
		h-index	g-index

12	12	12	346
all docs	docs citations	times ranked	citing authors

#	ARTICLE	IF	CITATIONS
1	A pandemic-enabled comparison of discovery platforms demonstrates a naiÌ^ve antibody library can match the best immune-sourced antibodies. Nature Communications, 2022, 13, 462.	5.8	17
2	A single donor is sufficient to produce a highly functional in vitro antibody library. Communications Biology, 2021, 4, 350.	2.0	12
3	A refined genome phage display methodology delineates the human antibody response in patients with Chagas disease. IScience, 2021, 24, 102540.	1.9	10
4	Protocol for design, construction, and selection of genome phage (gPhage) display libraries. STAR Protocols, 2021, 2, 100936.	0.5	2
5	Drug-like antibodies with high affinity, diversity and developability directly from next-generation antibody libraries. MAbs, 2021, 13, 1980942.	2.6	24
6	Exploiting next-generation sequencing in antibody selections â€" a simple PCR method to recover binders. MAbs, 2020, 12, 1701792.	2.6	7
7	Recombinant Antibodies against Mycolactone. Toxins, 2019, 11, 346.	1.5	9
8	A ligand motif enables differential vascular targeting of endothelial junctions between brain and retina. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 2300-2305.	3.3	14
9	CD100/Sema4D Increases Macrophage Infection by Leishmania (Leishmania) amazonensis in a CD72 Dependent Manner. Frontiers in Microbiology, 2018, 9, 1177.	1.5	8
10	Going viral? Linking the etiology of human prostate cancer to the *PCA 3* long noncoding *RNA* and oncogenic viruses. EMBO Molecular Medicine, 2017, 9, 1327-1330.	3.3	10
11	Trypanosoma cruzi Binds to Cytokeratin through Conserved Peptide Motifs Found in the Laminin-G-Like Domain of the gp85/Trans-sialidase Proteins. PLoS Neglected Tropical Diseases, 2015, 9, e0004099.	1.3	20
12	Fast Proton Titration Scheme for Multiscale Modeling of Protein Solutions. Journal of Chemical Theory and Computation, 2010, 6, 3259-3266.	2.3	46