

Robert J Pantazes

List of Publications by Year in descending order

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13
papers

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citations

932766

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citing authors

#	ARTICLE	IF	CITATIONS
1	Nanobody-based CTLA4 inhibitors for immune checkpoint blockade therapy of canine cancer patients. <i>Scientific Reports</i> , 2021, 11, 20763.	1.6	10
2	Antibody epitope repertoire analysis enables rapid antigen discovery and multiplex serology. <i>Scientific Reports</i> , 2020, 10, 5294.	1.6	31
3	Development and Analyses of a Database of Antibody – Antigen Complexes. <i>Computer Aided Chemical Engineering</i> , 2018, 44, 2113-2118.	0.3	3
4	De novo design of antibody complementarity determining regions binding a FLAG tetra-peptide. <i>Scientific Reports</i> , 2017, 7, 10295.	1.6	27
5	Identification of disease-specific motifs in the antibody specificity repertoire via next-generation sequencing. <i>Scientific Reports</i> , 2016, 6, 30312.	1.6	35
6	Engineering pH responsive fibronectin domains for biomedical applications. <i>Journal of Biological Engineering</i> , 2015, 9, 6.	2.0	9
7	The Iterative Protein Redesign and Optimization (IPRO) suite of programs. <i>Journal of Computational Chemistry</i> , 2015, 36, 251-263.	1.5	34
8	OptMAVE – A New Framework for the de novo Design of Antibody Variable Region Models Targeting Specific Antigen Epitopes. <i>PLoS ONE</i> , 2014, 9, e105954.	1.1	59
9	MAPs: a database of modular antibody parts for predicting tertiary structures and designing affinity matured antibodies. <i>BMC Bioinformatics</i> , 2013, 14, 168.	1.2	24
10	OptZyme: Computational Enzyme Redesign Using Transition State Analogues. <i>PLoS ONE</i> , 2013, 8, e75358.	1.1	22
11	Recent advances in computational protein design. <i>Current Opinion in Structural Biology</i> , 2011, 21, 467-472.	2.6	78
12	Computational design of <i>Candida boidinii</i> xylose reductase for altered cofactor specificity. <i>Protein Science</i> , 2009, 18, 2125-2138.	3.1	84
13	Optimal protein library design using recombination or point mutations based on sequence-based scoring functions. <i>Protein Engineering, Design and Selection</i> , 2007, 20, 361-373.	1.0	37