## Barbara Mouratou

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Application of Affitins for Affinity Purification of Proteins. Methods in Molecular Biology, 2022, 2466, 37-48.	0.9	0
2	Affitins: Ribosome Display for Selection of Aho7c-Based Affinity Proteins. Methods in Molecular Biology, 2020, 2070, 19-41.	0.9	6
3	Multivalent Affidendrons with High Affinity and Specificity toward <i>Staphylococcus aureus</i> as Versatile Tools for Modulating Multicellular Behaviors. ACS Applied Materials & Interfaces, 2019, 11, 21391-21398.	8.0	11
4	Wholeâ€bacterium ribosome display selection for isolation of highly specific anti―Staphyloccocus aureus Affitins for detection―and captureâ€based biomedical applications. Biotechnology and Bioengineering, 2019, 116, 1844-1855.	3.3	9
5	A novel, smaller scaffold for Affitins: Showcase with binders specific for EpCAM. Biotechnology and Bioengineering, 2018, 115, 290-299.	3.3	19
6	The archaeal "7 kDa DNA-binding―proteins: extended characterization of an old gifted family. Scientific Reports, 2016, 6, 37274.	3.3	21
7	Affitins for protein purification by affinity magnetic fishing. Journal of Chromatography A, 2016, 1457, 50-58.	3.7	22
8	Affitins as robust tailored reagents for affinity chromatography purification of antibodies and non-immunoglobulin proteins. Journal of Chromatography A, 2016, 1441, 44-51.	3.7	29
9	Artificial Affinity Proteins as Ligands of Immunoglobulins. Biomolecules, 2015, 5, 60-75.	4.0	24
10	Affinity transfer to the archaeal extremophilic Sac7d protein by insertion of a CDR. Protein Engineering, Design and Selection, 2014, 27, 431-438.	2.1	16
11	Switching an anti-IgG binding site between archaeal extremophilic proteins results in Affitins with enhanced pH stability. Journal of Biotechnology, 2014, 192, 123-129.	3.8	18
12	Potent and Specific Inhibition of Glycosidases by Small Artificial Binding Proteins (Affitins). PLoS ONE, 2014, 9, e97438.	2.5	42
13	Tolerance of the archaeal Sac7d scaffold protein to alternative library designs: characterization of anti-immunoglobulin G Affitins. Protein Engineering, Design and Selection, 2013, 26, 267-275.	2.1	38
14	Ribosome Display for the Selection of Sac7d Scaffolds. Methods in Molecular Biology, 2012, 805, 315-331.	0.9	25
15	Remodeling a DNA-binding protein as a specific <i>in vivo</i> inhibitor of bacterial secretin PulD. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 17983-17988.	7.1	78
16	The domain architecture of large guanine nucleotide exchange factors for the small GTP-binding protein Arf. BMC Genomics, 2005, 6, 20.	2.8	102
17	Application of a modified version of Habeeb's trinitrophenylation method for the characterization of hapten–protein conjugates in a reversed micellar medium. Journal of Immunological Methods, 2002, 263, 75-83.	1.4	12
18	A method for the detection and screening of catalytic anti-DNA antibodies. Journal of Immunological Methods. 2002. 269. 147-155.	1.4	11

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19	Development of nonradioactive microtiter plate assays for nuclease activity. Analytical Biochemistry, 2002, 309, 40-47.	2.4	15
20	Amino Acid Analysis by High-Performance Liquid Chromatography after Derivatization with 1-Fluoro-2,4-dinitrophenyl-5-L-alanine Amide (Marfey's Reagent). , 2000, 159, 049-054.		4
21	Conversion of Tyrosine Phenol-lyase to Dicarboxylic Amino Acid β-Lyase, an Enzyme Not Found in Nature. Journal of Biological Chemistry, 1999, 274, 1320-1325.	3.4	24
22	Transferred Nuclear Overhauser Effect Study of the C-Terminal Helix of Yeast Phosphoglycerate Kinase: NMR Solution Structure of the C-Terminal Bound Peptide. Biochemistry, 1995, 34, 842-846.	2.5	11
23	Role of the C-Terminal Helix in the Folding and Stability of Yeast Phosphoglycerate Kinase. Biochemistry, 1995, 34, 833-841.	2.5	36