

Katja M Arndt

List of Publications by Year in descending order

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54
papers

2,386
citations

304743

22
h-index

206112

48
g-index

54
all docs

54
docs citations

54
times ranked

2854
citing authors

#	ARTICLE	IF	CITATIONS
1	Strategies for Enzymatic Inactivation of the Veterinary Antibiotic Florfenicol. <i>Antibiotics</i> , 2022, 11, 443.	3.7	2
2	Bacteriophage-templated Assembly of Magnetic Nanoparticles and Their Actuation Potential. <i>ChemNanoMat</i> , 2021, 7, 942-949.	2.8	3
3	Characterizing Transcriptional Interference between Converging Genes in Bacteria. <i>ACS Synthetic Biology</i> , 2019, 8, 466-473.	3.8	14
4	Detection of Incorporation of p-Coumaric Acid into Photoactive Yellow Protein Variants in Vivo. <i>Biochemistry</i> , 2019, 58, 2682-2694.	2.5	6
5	rAAV Engineering for Capsid-Protein Enzyme Insertions and Mosaicism Reveals Resilience to Mutational, Structural and Thermal Perturbations. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5702.	4.1	14
6	Selection of Protein-Protein Interactions of Desired Affinities with a Bandpass Circuit. <i>Journal of Molecular Biology</i> , 2019, 431, 391-400.	4.2	2
7	Strategies for the photo-control of endogenous protein activity. <i>Current Opinion in Structural Biology</i> , 2017, 45, 53-58.	5.7	14
8	A user-friendly, low-cost turbidostat with versatile growth rate estimation based on an extended Kalman filter. <i>PLoS ONE</i> , 2017, 12, e0181923.	2.5	21
9	Long-range transcriptional interference in <i>E. coli</i> used to construct a dual positive selection system for genetic switches. <i>Nucleic Acids Research</i> , 2016, 44, e95-e95.	14.5	18
10	An <i>Escherichia coli</i> system for evolving improved light-controlled DNA-binding proteins. <i>Protein Engineering, Design and Selection</i> , 2015, 28, 293-302.	2.1	8
11	Characterization and inhibition of AF10-mediated interaction. <i>Journal of Peptide Science</i> , 2014, 20, 385-397.	1.4	7
12	Controlling leucine-zipper partner recognition in cells through modification of a ^g interactions. <i>Chemical Communications</i> , 2014, 50, 6364-6367.	4.1	8
13	Analysis of Selected and Designed Chimeric D- and L-Helix Assemblies. <i>Biomacromolecules</i> , 2014, 15, 3296-3305.	5.4	3
14	Improving coiled coil stability while maintaining specificity by a bacterial hitchhiker selection system. <i>Journal of Structural Biology</i> , 2014, 186, 335-348.	2.8	11
15	Modular adeno-associated virus (rAAV) vectors used for cellular virus-directed enzyme prodrug therapy. <i>Scientific Reports</i> , 2014, 4, 3759.	3.3	28
16	Directional cloning of DNA fragments using deoxyinosine-containing oligonucleotides and endonuclease V. <i>BMC Biotechnology</i> , 2013, 13, 81.	3.3	7
17	TAT hitchhiker selection expanded to folding helpers, multimeric interactions and combinations with protein fragment complementation. <i>Protein Engineering, Design and Selection</i> , 2013, 26, 225-242.	2.1	2
18	Free energy calculations of the interactions of c-Jun-based synthetic peptides with the c-Fos protein. <i>Biopolymers</i> , 2012, 97, 899-909.	2.4	10

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19	Exploring the Molecular Linkage of Protein Stability Traits for Enzyme Optimization by Iterative Truncation and Evolution. <i>Biochemistry</i> , 2012, 51, 4850-4867.	2.5	13
20	Standardization in Synthetic Biology. <i>Methods in Molecular Biology</i> , 2012, 813, 23-43.	0.9	38
21	Light-Controlled Gene Switches in Mammalian Cells. <i>Methods in Molecular Biology</i> , 2012, 813, 195-210.	0.9	4
22	Nucleotide Exchange and Excision Technology DNA Shuffling and Directed Evolution. <i>Methods in Molecular Biology</i> , 2011, 687, 333-344.	0.9	7
23	Efficient phage display of intracellularly folded proteins mediated by the TAT pathway. <i>Protein Engineering, Design and Selection</i> , 2011, 24, 473-484.	2.1	27
24	Photocontrol of Coiled-Coil Proteins in Living Cells. <i>Angewandte Chemie - International Edition</i> , 2010, 49, 3943-3946.	13.8	108
25	SynBioWave—a real-time communication platform for molecular and synthetic biology. <i>Bioinformatics</i> , 2010, 26, 2782-2783.	4.1	7
26	Improving the interaction of Myc-interfering peptides with Myc using molecular dynamics simulations. <i>Journal of Peptide Science</i> , 2009, 15, 5-15.	1.4	11
27	Role of Hydrophobic and Electrostatic Interactions in Coiled Coil Stability and Specificity. <i>Biochemistry</i> , 2009, 48, 10380-10388.	2.5	34
28	Selection of Peptides Interfering with Protein-Protein Interaction. <i>Methods in Molecular Biology</i> , 2009, 535, 263-291.	0.9	7
29	Targeting the c-Myc coiled coil with interfering peptides. <i>Journal of Peptide Science</i> , 2008, 14, 1022-1031.	1.4	20
30	Selectional and Mutational Scope of Peptides Sequestering the Jun-Fos Coiled-Coil Domain. <i>Journal of Molecular Biology</i> , 2008, 381, 73-88.	4.2	49
31	iPEP: peptides designed and selected for interfering with protein interaction and function. <i>Biochemical Society Transactions</i> , 2008, 36, 1442-1447.	3.4	13
32	Protein Engineering. <i>Springer Protocols</i> , 2008, , 587-629.	0.3	2
33	Directed Protein Evolution. <i>Springer Protocols</i> , 2008, , 631-656.	0.3	3
34	Considerations in the Design and Optimization of Coiled Coil Structures. , 2007, 352, 35-70.		29
35	A General Method of Terminal Truncation, Evolution, and Re-Elongation to Generate Enzymes of Enhanced Stability. , 2007, 352, 275-304.		6
36	Versatile DNA Fragmentation and Directed Evolution With Nucleotide Exchange and Excision Technology. , 2007, 352, 167-190.		1

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37	Improved Stability of the Jun-Fos Activator Protein-1 Coiled Coil Motif. <i>Journal of Biological Chemistry</i> , 2007, 282, 23015-23024.	3.4	39
38	Positive Aspects of Negative Design: Simultaneous Selection of Specificity and Interaction Stability. <i>Biochemistry</i> , 2007, 46, 4804-4814.	2.5	55
39	Semirational design of Jun-Fos coiled coils with increased affinity: Universal implications for leucine zipper prediction and design. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 8989-8994.	7.1	127
40	Nucleotide exchange and excision technology (NExT) DNA shuffling: a robust method for DNA fragmentation and directed evolution. <i>Nucleic Acids Research</i> , 2005, 33, e117-e117.	14.5	36
41	Coiled Coil Domains: Stability, Specificity, and Biological Implications. <i>ChemBioChem</i> , 2004, 5, 170-176.	2.6	611
42	Coiled Coil Domains: Stability, Specificity, and Biological Implications. <i>ChemInform</i> , 2004, 35, no.	0.0	0
43	Comparison of In Vivo Selection and Rational Design of Heterodimeric Coiled Coils. <i>Structure</i> , 2002, 10, 1235-1248.	3.3	51
44	Helix-stabilized fv (hsfv) antibody fragments: substituting the constant domains of a fab fragment for a heterodimeric coiled-coil domain 1 Edited by I. A. Wilson. <i>Journal of Molecular Biology</i> , 2001, 312, 221-228.	4.2	62
45	[22] Selectively infective phage technology. <i>Methods in Enzymology</i> , 2000, 328, 364-388.	1.0	6
46	A heterodimeric coiled-coil peptide pair selected in vivo from a designed library- versus -library ensemble 1 Edited by A. R. Fersht. <i>Journal of Molecular Biology</i> , 2000, 295, 627-639.	4.2	101
47	[17] Protein fusions to coiled-coil domains. <i>Methods in Enzymology</i> , 2000, 328, 261-282.	1.0	25
48	An in vivo library-versus-library selection of optimized protein-protein interactions. <i>Nature Biotechnology</i> , 1999, 17, 683-690.	17.5	182
49	Selectively infective phage (SIP) technology: scope and limitations. <i>Journal of Immunological Methods</i> , 1999, 231, 93-104.	1.4	35
50	Tandem Immobilized Metal-Ion Affinity Chromatography/Immunoaffinity Purification of His-tagged Proteins Evaluation of Two Anti-His-Tag Monoclonal Antibodies. <i>Analytical Biochemistry</i> , 1998, 259, 54-61.	2.4	75
51	Model and Simulation of Multivalent Binding to Fixed Ligands. <i>Analytical Biochemistry</i> , 1998, 261, 149-158.	2.4	135
52	The first constant domain (CH1 and CL) of an antibody used as heterodimerization domain for bispecific miniantibodies. <i>FEBS Letters</i> , 1998, 422, 259-264.	2.8	76
53	A dimeric bispecific miniantibody combines two specificities with avidity. <i>FEBS Letters</i> , 1998, 432, 45-49.	2.8	69
54	Factors Influencing the Dimer to Monomer Transition of an Antibody Single-Chain Fv Fragment. <i>Biochemistry</i> , 1998, 37, 12918-12926.	2.5	144