

Shao-Qing Zhang

List of Publications by Year in descending order

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16
papers

927
citations

687220

13
h-index

940416

16
g-index

17
all docs

17
docs citations

17
times ranked

1665
citing authors

#	ARTICLE	IF	CITATIONS
1	Designing a Green Fluorogenic Protease Reporter by Flipping a Beta Strand of GFP for Imaging Apoptosis in Animals. <i>Journal of the American Chemical Society</i> , 2019, 141, 4526-4530.	6.6	64
2	<i>De Novo</i> Design of Tetranuclear Transition Metal Clusters Stabilized by Hydrogen-Bonded Networks in Helical Bundles. <i>Journal of the American Chemical Society</i> , 2018, 140, 1294-1304.	6.6	32
3	Visualizing Dynamics of Cell Signaling In Vivo with a Phase Separation-Based Kinase Reporter. <i>Molecular Cell</i> , 2018, 69, 334-346.e4.	4.5	83
4	Spectroscopic and metal binding properties of a <i>de novo</i> metalloprotein binding a tetrazinc cluster. <i>Biopolymers</i> , 2018, 109, e23339.	1.2	15
5	Designed peptides that assemble into cross- β amyloid-like structures. <i>Nature Chemical Biology</i> , 2018, 14, 870-875.	3.9	62
6	De novo design of a hyperstable non-natural protein-ligand complex with sub-Å accuracy. <i>Nature Chemistry</i> , 2017, 9, 1157-1164.	6.6	93
7	Protein-directed self-assembly of a fullerene crystal. <i>Nature Communications</i> , 2016, 7, 11429.	5.8	55
8	Swapped-Domain Constructs of the Glycoprotein-41 Ectodomain Are Potent Inhibitors of HIV Infection. <i>ACS Chemical Biology</i> , 2015, 10, 1247-1257.	1.6	4
9	The Membrane- and Soluble-Protein Helix-Helix Interactome: Similar Geometry via Different Interactions. <i>Structure</i> , 2015, 23, 527-541.	1.6	64
10	A naturally monomeric infrared fluorescent protein for protein labeling in vivo. <i>Nature Methods</i> , 2015, 12, 763-765.	9.0	146
11	Design and characterization of swapped-domain constructs of HIV-1 glycoprotein-41 as receptors for drug discovery. <i>Protein Engineering, Design and Selection</i> , 2015, 28, 107-116.	1.0	4
12	Crystal structure of an amphiphilic foldamer reveals a 48-mer assembly comprising a hollow truncated octahedron. <i>Nature Communications</i> , 2014, 5, 3581.	5.8	14
13	Deciphering Regulatory Mechanism of the Juxtamembrane Region in Thrombopoietin Receptor Activation. <i>Biophysical Journal</i> , 2014, 106, 103a.	0.2	1
14	Structural Stability and Binding Strength of a Designed Peptide-Carbon Nanotube Hybrid. <i>Journal of Physical Chemistry C</i> , 2013, 117, 26255-26261.	1.5	13
15	Manipulating Biopolymer Dynamics by Anisotropic Nanoconfinement. <i>Nano Letters</i> , 2007, 7, 3438-3442.	4.5	31
16	Molecular crowding enhances native structure and stability of β^2 protein flavodoxin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 18976-18981.	3.3	245