Sosuke Yoshinaga

List of Publications by Year in descending order

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1478505 1372567 12 141 10 6 citations h-index g-index papers 12 12 12 174 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Targeting FROUNT with disulfiram suppresses macrophage accumulation and its tumor-promoting properties. Nature Communications, 2020, 11 , 609.	12.8	57
2	Application of spin-crossover water soluble nanoparticles for use as MRI contrast agents. Scientific Reports, 2018, 8, 14911.	3.3	23
3	Structure of the Mouse Sex Peptide Pheromone ESP1 Reveals a Molecular Basis for Specific Binding to the Class C G-protein-coupled Vomeronasal Receptor. Journal of Biological Chemistry, 2013, 288, 16064-16072.	3.4	17
4	Identification of a binding element for the cytoplasmic regulator FROUNT in the membrane-proximal C-terminal region of chemokine receptors CCR2 and CCR5. Biochemical Journal, 2014, 457, 313-322.	3.7	13
5	Expression and purification of human FROUNT, a common cytosolic regulator of CCR2 and CCR5. Protein Expression and Purification, 2011, 77, 86-91.	1.3	10
6	Structural basis for the binding of the membraneâ€proximal Câ€terminal region of chemokine receptor <scp>CCR</scp> 2 with the cytosolic regulator <scp>FROUNT</scp> . FEBS Journal, 2014, 281, 5552-5566.	4.7	9
7	Identification and Preparation of a Novel Chemokine Receptor-Binding Domain in the Cytoplasmic Regulator FROUNT. Molecular Biotechnology, 2017, 59, 141-150.	2.4	5
8	1H, 13C and 15N resonance assignments for a chemokine receptor-binding domain of FROUNT, a cytoplasmic regulator of chemotaxis. Biomolecular NMR Assignments, 2018, 12, 259-262.	0.8	3
9	Proton Relaxation Time in Water-soluble Metal Complex Nanoparticles. Chemistry Letters, 2018, 47, 598-600.	1.3	2
10	Efficient identification of compounds suppressing protein precipitation via solvent screening using serial deletion mutants of the target protein. Genes To Cells, 2018, 23, 70-79.	1.2	2
11	Expression and purification of mouse peptide ESP4 in Escherichia coli. Protein Expression and Purification, 2014, 96, 20-25.	1.3	0
12	Rational Design of Monodispersed Mutants of Proteins by Identifying Aggregation Contact Sites Using Solubilizing Agents. Biochemistry, 2020, 59, 3639-3649.	2.5	0