Sheng-Wei Lin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1330050/publications.pdf

Version: 2024-02-01

759233 642732 24 551 12 23 h-index citations g-index papers 24 24 24 983 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Structure and Mechanism of Helicobacter pylori Fucosyltransferase. Journal of Biological Chemistry, 2007, 282, 9973-9982.	3.4	113
2	Identification, synthesis and evaluation of SARS-CoV and MERS-CoV 3C-like protease inhibitors. Bioorganic and Medicinal Chemistry, 2016, 24, 3035-3042.	3.0	81
3	Carboxyl Terminus ofHelicobacter pyloriα1,3-Fucosyltransferase Determines the Structure and Stabilityâ€. Biochemistry, 2006, 45, 8108-8116.	2.5	62
4	Galectin-3 Binding Protein and Galectin-1 Interaction in Breast Cancer Cell Aggregation and Metastasis. Journal of the American Chemical Society, 2015, 137, 9685-9693.	13.7	58
5	Rad51 presynaptic filament stabilization function of the mouse Swi5–Sfr1 heterodimeric complex. Nucleic Acids Research, 2012, 40, 6558-6569.	14.5	34
6	Enhancement of ADP release from the RAD51 presynaptic filament by the SWI5-SFR1 complex. Nucleic Acids Research, 2014, 42, 349-358.	14.5	27
7	Functional attributes of the Saccharomyces cerevisiae meiotic recombinase Dmc1. DNA Repair, 2013, 12, 707-712.	2.8	21
8	Predominant structural configuration of natural antibody repertoires enables potent antibody responses against protein antigens. Scientific Reports, 2015, 5, 12411.	3.3	17
9	Role of N-Linked Glycans in the Interactions of Recombinant HCV Envelope Glycoproteins with Cellular Receptors. ACS Chemical Biology, 2014, 9, 1437-1443.	3.4	16
10	The differential expression of the blood group <i>P¹</i> â€ <i>A4GALT</i> and <i>P²</i> â€ <i>A4GALT</i> alleles is stimulated by the transcription factor early growth response 1. Transfusion, 2018, 58, 1054-1064.	1.6	16
11	The molecular genetic background leading to the formation of the human erythroid-specific Xga/CD99 blood groups. Blood Advances, 2018, 2, 1854-1864.	5.2	16
12	RoleÂof the RAD51–SWI5–SFR1 Ensemble in homologous recombination. Nucleic Acids Research, 2016, 44, 6242-6251.	14.5	14
13	Functional Relationship of ATP Hydrolysis, Presynaptic Filament Stability, and Homologous DNA Pairing Activity of the Human Meiotic Recombinase DMC1. Journal of Biological Chemistry, 2015, 290, 19863-19873.	3.4	12
14	Chemoenzymatic Synthesis of GDPâ€ <scp>L</scp> â€Fucose Derivatives as Potent and Selective αâ€1,3â€Fucosyltransferase Inhibitors. Advanced Synthesis and Catalysis, 2012, 354, 1750-1758.	4.3	11
15	Coll(Chromomycin)2 Complex Induces a Conformational Change of CCG Repeats from i-Motif to Base-Extruded DNA Duplex. International Journal of Molecular Sciences, 2018, 19, 2796.	4.1	10
16	A chemical probe inhibitor targeting STAT1 restricts cancer stem cell traits and angiogenesis in colorectal cancer. Journal of Biomedical Science, 2022, 29, 20.	7.0	10
17	Moiety-Linkage Map Reveals Selective Nonbisphosphonate Inhibitors of Human Geranylgeranyl Diphosphate Synthase. Journal of Chemical Information and Modeling, 2013, 53, 2299-2311.	5.4	7
18	TRIM28 Regulates Dlk1 Expression in Adipogenesis. International Journal of Molecular Sciences, 2020, 21, 7245.	4.1	7

#	Article	IF	CITATION
19	Functional characterization of the meiosis-specific DNA double-strand break inducing factor SPO-11 from C. elegans. Scientific Reports, 2017, 7, 2370.	3.3	6
20	Fluorescent Farnesyl Diphosphate Analogue: A Probe To Validate trans-Prenyltransferase Inhibitors. Biochemistry, 2016, 55, 4366-4374.	2.5	4
21	Control Activity of Yeast Geranylgeranyl Diphosphate Synthase from Dimer Interface through H-Bonds and Hydrophobic Interaction. Biochemistry, 2013, 52, 2783-2792.	2.5	3
22	The functional characterization of phosphorylation of tristetraprolin at C-terminal NOT1-binding domain. Journal of Inflammation, 2021, 18, 22.	3.4	2
23	Structural and bioinformatic analyses of Azemiops venom serine proteases reveal close phylogeographic relationships to pitvipers from eastern China and the New World. Toxicon, 2021, 198, 93-101.	1.6	2
24	Generation of TRIM28 Knockout K562 Cells by CRISPR/Cas9 Genome Editing and Characterization of TRIM28-Regulated Gene Expression in Cell Proliferation and Hemoglobin Beta Subunits. International Journal of Molecular Sciences, 2022, 23, 6839.	4.1	2