

Sun-gu Lee

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3990684/publications.pdf>

Version: 2024-02-01

72
papers

1,220
citations

471061

17
h-index

395343

33
g-index

75
all docs

75
docs citations

75
times ranked

1938
citing authors

#	ARTICLE	IF	CITATIONS
1	A Study on the Effect of Surface Lysine to Arginine Mutagenesis on Protein Stability and Structure Using Green Fluorescent Protein. <i>PLoS ONE</i> , 2012, 7, e40410.	1.1	198
2	Design of artificial cell-cell communication using gene and metabolic networks. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 2299-2304.	3.3	151
3	Development of a Selective, Sensitive, and Reversible Biosensor by the Genetic Incorporation of a Metal-Binding Site into Green Fluorescent Protein. <i>Angewandte Chemie - International Edition</i> , 2011, 50, 6534-6537.	7.2	55
4	Production of cytidine 5'-monophosphate-N-acetylneuraminic acid using recombinant <i>Escherichia coli</i> as a biocatalyst. <i>Biotechnology and Bioengineering</i> , 2002, 80, 516-524.	1.7	50
5	Functional expression of single-chain variable fragment antibody against c-Met in the cytoplasm of <i>Escherichia coli</i> . <i>Protein Expression and Purification</i> , 2006, 47, 203-209.	0.6	48
6	Overexpression of α -type cytochrome, CymA in <i>Shewanella oneidensis</i> for enhanced bioelectricity generation and cell growth in a microbial fuel cell. <i>Journal of Chemical Technology and Biotechnology</i> , 2019, 94, 2115-2122.	1.6	44
7	Engineering Protein Sequence Composition for Folding Robustness Renders Efficient Noncanonical Amino acid Incorporations. <i>ChemBioChem</i> , 2010, 11, 2521-2524.	1.3	33
8	Modulation of protein stability and aggregation properties by surface charge engineering. <i>Molecular BioSystems</i> , 2013, 9, 2379.	2.9	32
9	In Silico Characterization of the Binding Modes of Surfactants with Bovine Serum Albumin. <i>Scientific Reports</i> , 2019, 9, 10643.	1.6	32
10	Synthesis of Well-Defined (Nitrilotriacetic Acid)-End-Functionalized Polystyrenes and Their Bioconjugation with Histidine-Tagged Green Fluorescent Proteins. <i>Macromolecules</i> , 2011, 44, 4672-4680.	2.2	30
11	Simultaneously Enhancing the Stability and Catalytic Activity of Multimeric Lysine Decarboxylase CadA by Engineering Interface Regions for Enzymatic Production of Cadaverine at High Concentration of Lysine. <i>Biotechnology Journal</i> , 2017, 12, 1700278.	1.8	30
12	Exploring the differences and similarities between urea and thermally driven denaturation of bovine serum albumin: intermolecular forces and solvation preferences. <i>Journal of Molecular Modeling</i> , 2018, 24, 75.	0.8	27
13	Enhancing the thermal stability of a single-chain Fv fragment by in vivo global fluorination of the proline residues. <i>Molecular BioSystems</i> , 2011, 7, 258-265.	2.9	26
14	Improving the productivity of single-chain Fv antibody against c-Met by rearranging the order of its variable domains. <i>Journal of Microbiology and Biotechnology</i> , 2008, 18, 1186-90.	0.9	22
15	Importance of expression system in the production of unnatural recombinant proteins in <i>Escherichia coli</i> . <i>Biotechnology and Bioprocess Engineering</i> , 2009, 14, 257-265.	1.4	20
16	Engineering an aldehyde dehydrogenase toward its substrates, 3-hydroxypropanal and NAD ⁺ , for enhancing the production of 3-hydroxypropionic acid. <i>Scientific Reports</i> , 2017, 7, 17155.	1.6	19
17	Effect of ligand torsion number on the AutoDock mediated prediction of protein-ligand binding affinity. <i>Journal of Industrial and Engineering Chemistry</i> , 2020, 83, 359-365.	2.9	19
18	Redesigning of anti-c-Met single chain Fv antibody for the cytoplasmic folding and its structural analysis. <i>Biotechnology and Bioengineering</i> , 2010, 106, 367-375.	1.7	18

#	ARTICLE	IF	CITATIONS
19	Conjugation of Proteins by Installing BIO-Orthogonally Reactive Groups at Their N-Termini. PLoS ONE, 2012, 7, e46741.	1.1	18
20	Production of (S)-styrene oxide using styrene oxide isomerase negative mutant of Pseudomonas putida SN1. Enzyme and Microbial Technology, 2006, 39, 1264-1269.	1.6	17
21	Biosynthetic substitution of tyrosine in green fluorescent protein with its surrogate fluorotyrosine in Escherichia coli. Biotechnology Letters, 2011, 33, 2201-2207.	1.1	17
22	Encapsulation of Nanoparticles Using Nitrilotriacetic Acid End-Functionalized Polystyrenes and Their Application for the Separation of Proteins. Advanced Functional Materials, 2012, 22, 4032-4037.	7.8	17
23	In silico study on the effect of surface lysines and arginines on the electrostatic interactions and protein stability. Biotechnology and Bioprocess Engineering, 2013, 18, 18-26.	1.4	17
24	Comparison of P aprE , P amyE , and P P43 promoter strength for Î²-galactosidase and staphylokinase expression in Bacillus subtilis. Biotechnology and Bioprocess Engineering, 2008, 13, 313-318.	1.4	16
25	In situ formation of polymer-protein hybrid spherical aggregates from (nitrilotriacetic) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 9 5	1.9	16
26	Structural and sequence features of two residue turns in beta-hairpins. Proteins: Structure, Function and Bioinformatics, 2014, 82, 1721-1733.	1.5	14
27	Development of recombinant Pseudomonas putida containing homologous styrene monooxygenase genes for the production of (S)-styrene oxide. Biotechnology and Bioprocess Engineering, 2006, 11, 530-537.	1.4	13
28	One-step immobilization and purification of his-tagged enzyme using poly(2-acetamidoacrylic acid) hydrogel. Macromolecular Research, 2013, 21, 5-9.	1.0	12
29	A variant of green fluorescent protein exclusively deposited to active intracellular inclusion bodies. Microbial Cell Factories, 2014, 13, 68.	1.9	12
30	Stable isotopic labeling-based quantitative targeted glycomics (iâ€‹scp>QT</scp>a<scp>G</scp>). Biotechnology Progress, 2015, 31, 840-848.	1.3	12
31	In vivo Protein Evolution, Next Generation Protein Engineering Strategy: from Random Approach to Target-specific Approach. Biotechnology and Bioprocess Engineering, 2019, 24, 85-94.	1.4	12
32	Construction and characterization of a recombinant whole-cell biocatalyst of Escherichia coli expressing styrene monooxygenase under the control of arabinose promoter. Biotechnology and Bioprocess Engineering, 2008, 13, 69-76.	1.4	11
33	Assessment of Computational Modeling of Fc-Fc Receptor Binding Through Protein-protein Docking Tool. Biotechnology and Bioprocess Engineering, 2020, 25, 734-741.	1.4	11
34	A comparative study on the stability and structure of two different green fluorescent proteins in organic co-solvent systems. Biotechnology and Bioprocess Engineering, 2013, 18, 342-349.	1.4	10
35	Title is missing!. Biotechnology Letters, 2000, 22, 819-823.	1.1	8
36	Deletional Protein Engineering Based on Stable Fold. PLoS ONE, 2012, 7, e51510.	1.1	8

#	ARTICLE	IF	CITATIONS
37	Site-specific reversible immobilization and purification of His-tagged protein on poly(ϵ -acetamidoacrylic) Tj ETQg1 1 0.784314 rgb	1.6	7
38	Effect of molecular properties of the protein-ligand complex on the prediction accuracy of AutoDock. Journal of Molecular Graphics and Modelling, 2021, 106, 107921.	1.3	7
39	Improving the growth rate of Escherichia coli DH5 α at low temperature through engineering of GroEL/S chaperone system. Biotechnology and Bioengineering, 2008, 99, 515-520.	1.7	6
40	NADH-dependent lactate dehydrogenase from Alcaligenes eutrophus H16 reduces 2-oxoadipate to 2-hydroxyadipate. Biotechnology and Bioprocess Engineering, 2014, 19, 1048-1057.	1.4	6
41	Identification of an Ideal-like Fingerprint for a Protein Fold using Overlapped Conserved Residues based Approach. Scientific Reports, 2015, 4, 5643.	1.6	6
42	RiSLnet: Rapid identification of smart mutant libraries using protein structure network. Application to thermal stability enhancement. Biotechnology and Bioengineering, 2019, 116, 250-259.	1.7	6
43	In vivo Production of Functional Single-Chain Fv Fragment with an N-Terminal-Specific Bio-orthogonal Reactive Group. ChemBioChem, 2010, 11, 498-501.	1.3	5
44	Soft Immobilization of Proteins onto Single-Walled Carbon Nanotubes through Nickel Complexed Nitrilotriacetic Acid-End Functionalized Polystyrenes. Israel Journal of Chemistry, 2012, 52, 359-363.	1.0	5
45	Sequence and Structural Features of Subsite Residues in GH10 and GH11 Xylanases. Biotechnology and Bioprocess Engineering, 2018, 23, 311-318.	1.4	5
46	Validation on the molecular docking efficiency of lipocalin family of proteins. Journal of Industrial and Engineering Chemistry, 2018, 67, 293-300.	2.9	5
47	Identification of Novel Cupredoxin Homologs Using Overlapped Conserved Residues Based Approach. Journal of Microbiology and Biotechnology, 2015, 25, 127-136.	0.9	5
48	The effect of the cspA 5'-untranslated region on recombinant protein production at low temperature. Biotechnology and Bioprocess Engineering, 2008, 13, 366-371.	1.4	4
49	Identification of novel cytochrome P450 homologs using overlapped conserved residues based approach. Biotechnology and Bioprocess Engineering, 2015, 20, 431-438.	1.4	4
50	Combinatorial Effect of Ligand and Ligand-Binding Site Hydrophobicities on Binding Affinity. Journal of Chemical Information and Modeling, 2020, 60, 1678-1684.	2.5	4
51	Production of sialyltrisaccharides using β -galactosidase and trans-sialidase in one pot. Biotechnology and Bioprocess Engineering, 2000, 5, 215-218.	1.4	3
52	Deciphering the factors responsible for the stability of a GFP variant resistant to alkaline pH using molecular dynamics simulations. Biotechnology and Bioprocess Engineering, 2013, 18, 858-867.	1.4	3
53	Redesigning the type II' β -turn in green fluorescent protein to type I': Implications for folding kinetics and stability. Proteins: Structure, Function and Bioinformatics, 2014, 82, 2812-2822.	1.5	3
54	Joint-based description of protein structure: its application to the geometric characterization of membrane proteins. Scientific Reports, 2017, 7, 1056.	1.6	3

#	ARTICLE	IF	CITATIONS
55	In silico Study on Binding Specificities of Cellular Retinol Binding Protein and Its Q108R Mutant. <i>Biotechnology and Bioprocess Engineering</i> , 2022, 27, 126-134.	1.4	3
56	Structural Study on the Impact of S239D/I332E Mutations in the Binding of Fc and Fc γ R1IIa. <i>Biotechnology and Bioprocess Engineering</i> , 2021, 26, 985-992.	1.4	3
57	Colorimetric monitoring of the activity of recombinant <i>Escherichia coli</i> expressing styrene monooxygenase. <i>Journal of Industrial and Engineering Chemistry</i> , 2009, 15, 520-523.	2.9	2
58	Multivalent (Nitrilotriacetic Acid)-End-Functionalized Polystyrenes by ATRP and Their Self-Assembly. <i>Macromolecular Chemistry and Physics</i> , 2013, 214, 2027-2035.	1.1	2
59	Generation of efficient fingerprint for GFP-like fold and computational identification of potential GFP-like homologs. <i>Biotechnology and Bioprocess Engineering</i> , 2016, 21, 712-719.	1.4	2
60	Computational screening of potential non-immunoglobulin scaffolds using overlapped conserved residues (OCR)-based fingerprints. <i>Korean Journal of Chemical Engineering</i> , 2018, 35, 717-724.	1.2	2
61	Comparative Analysis of TM and Cytoplasmic β -barrel Conformations Using Joint Descriptor. <i>Scientific Reports</i> , 2018, 8, 14185.	1.6	2
62	In Silico Study on Retinoid-binding Modes in Human RBP and ApoD Lipocalins. <i>Biotechnology and Bioprocess Engineering</i> , 2018, 23, 158-167.	1.4	2
63	Control of acetate production rate in <i>Escherichia coli</i> by regulating expression of single-copy pta using lacI(Q) in multicopy plasmid. <i>Journal of Microbiology and Biotechnology</i> , 2008, 18, 334-7.	0.9	2
64	Biological synthesis of alkyne-terminated telechelic recombinant protein. <i>Macromolecular Research</i> , 2009, 17, 424-429.	1.0	1
65	Modulation of intracellular protein activity at level of protein folding by beta-turn engineering. <i>Biotechnology and Bioprocess Engineering</i> , 2014, 19, 433-441.	1.4	1
66	Engineering a beta-turn in green fluorescent protein to a foreign loop. <i>Journal of Industrial and Engineering Chemistry</i> , 2016, 33, 330-335.	2.9	1
67	Measuring the Conformational Distance of GPCR-related Proteins Using a Joint-based Descriptor. <i>Scientific Reports</i> , 2017, 7, 15205.	1.6	1
68	Identification of common and distinct features of ligand-binding sites in kernel and outlier lipocalins. <i>Journal of Industrial and Engineering Chemistry</i> , 2019, 78, 344-351.	2.9	1
69	Generation of anti-c-met single domain antibody fragment based on human stable frameworks. <i>Biotechnology and Bioprocess Engineering</i> , 2012, 17, 1120-1127.	1.4	0
70	(Nitrilotriacetic Acid)-End-Functionalized Polystyrenes Synthesized by ATRP. <i>ACS Symposium Series</i> , 2012, , 303-314.	0.5	0
71	Separation efficiency of free-solution conjugated electrophoresis with drag-tags incorporating a synthetic amino acid. <i>Electrophoresis</i> , 2016, 37, 818-825.	1.3	0
72	Characterization on the aggregation of self-aggregating green fluorescent protein variant. <i>Journal of Industrial and Engineering Chemistry</i> , 2017, 46, 337-341.	2.9	0