

Liqi Li

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

409
citations

840776

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1199594

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13
all docs

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docs citations

13
times ranked

602
citing authors

#	ARTICLE	IF	CITATIONS
1	Prediction of bacterial protein subcellular localization by incorporating various features into Chou's PseAAC and a backward feature selection approach. <i>Biochimie</i> , 2014, 104, 100-107.	2.6	63
2	Lipidome in colorectal cancer. <i>Oncotarget</i> , 2016, 7, 33429-33439.	1.8	61
3	An Ensemble Classifier for Eukaryotic Protein Subcellular Location Prediction Using Gene Ontology Categories and Amino Acid Hydrophobicity. <i>PLoS ONE</i> , 2012, 7, e31057.	2.5	55
4	A Transferrin-Conjugated Hollow Nanoplatform for Redox-Controlled and Targeted Chemotherapy of Tumor with Reduced Inflammatory Reactions. <i>Theranostics</i> , 2018, 8, 518-532.	10.0	48
5	YTHDF1 upregulation mediates hypoxia-dependent breast cancer growth and metastasis through regulating PKM2 to affect glycolysis. <i>Cell Death and Disease</i> , 2022, 13, 258.	6.3	35
6	Sustained raloxifene release from hyaluronan-alendronate-functionalized titanium nanotube arrays capable of enhancing osseointegration in osteoporotic rabbits. <i>Materials Science and Engineering C</i> , 2018, 82, 345-353.	7.3	32
7	Enhancement of local bone remodeling in osteoporotic rabbits by biomimic multilayered structures on Ti6Al4V implants. <i>Journal of Biomedical Materials Research - Part A</i> , 2016, 104, 1437-1451.	4.0	30
8	PSSP-RFE: Accurate Prediction of Protein Structural Class by Recursive Feature Extraction from PSI-BLAST Profile, Physical-Chemical Property and Functional Annotations. <i>PLoS ONE</i> , 2014, 9, e92863.	2.5	24
9	An ROS-Activatable Nanoassembly Remodulates Tumor Cell Metabolism for Enhanced Ferroptosis Therapy. <i>Advanced Healthcare Materials</i> , 2022, 11, e2101702.	7.6	22
10	Autophagy-Dependent Ferroptosis as a Therapeutic Target in Cancer. <i>ChemMedChem</i> , 2021, 16, 2942-2950.	3.2	21
11	Protein submitochondrial localization from integrated sequence representation and SVM-based backward feature extraction. <i>Molecular BioSystems</i> , 2015, 11, 170-177.	2.9	13
12	A machine-learning approach for predicting palmitoylation sites from integrated sequence-based features. <i>Journal of Bioinformatics and Computational Biology</i> , 2017, 15, 1650025.	0.8	5