Mojtaba Mortazavi

List of Publications by Year in descending order

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29	327	12	17
papers	citations	h-index	g-index
30	30	30	386
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Design of thermostable luciferases through arginine saturation in solvent-exposed loops. Protein Engineering, Design and Selection, 2011, 24, 893-903.	1.0	40
2	New Developments in Pichia pastoris Expression System, Review and Update. Current Pharmaceutical Biotechnology, 2018, 19, 451-467.	0.9	35
3	Synthesis and Characterization of GO/ZIF-67 Nanocomposite: Investigation of Catalytic Activity for the Determination of Epinine in the Presence of Dobutamine. Micromachines, 2022, 13, 88.	1.4	27
4	Spectroscopic and functional characterization of <italic>Lampyris turkestanicus</italic> luciferase: a comparative study. Acta Biochimica Et Biophysica Sinica, 2008, 40, 365-374.	0.9	22
5	Enhancement of Thermostability of Aspergillus flavus Urate Oxidase by Immobilization on the Ni-Based Magnetic Metal–Organic Framework. Nanomaterials, 2021, 11, 1759.	1.9	22
6	cDNA Cloning, Expression and Homology Modeling of a Luciferase from the Firefly Lampyroidea maculata. BMB Reports, 2006, 39, 578-585.	1.1	18
7	Surface charge modification increases firefly luciferase rigidity without alteration in bioluminescence spectra. Enzyme and Microbial Technology, 2017, 96, 47-59.	1.6	16
8	Design, synthesis and biological assessment of acridine derivatives containing 1,3,4-thiadiazole moiety as novel selective acetylcholinesterase inhibitors. Bioorganic Chemistry, 2020, 105, 104457.	2.0	16
9	Effects of 940 MHz EMF on luciferase solution: Structure, function, and dielectric studies. Bioelectromagnetics, 2013, 34, 489-498.	0.9	15
10	Bifunctional role of leucine 300 of firefly luciferase in structural rigidity. International Journal of Biological Macromolecules, 2017, 101, 67-74.	3.6	14
11	Immobilization of Lepidium draba peroxidase on a novel Zn-MOF nanostructure. International Journal of Biological Macromolecules, 2021, 173, 366-378.	3.6	13
12	In Silico Analysis of L1/L2 Sequences of Human Papillomaviruses: Implication for Universal Vaccine Design. Viral Immunology, 2017, 30, 210-223.	0.6	12
13	Bioinformatic Identification of Rare Codon Clusters (RCCs) in HBV Genome and Evaluation of RCCs in Proteins Structure of Hepatitis B Virus. Hepatitis Monthly, 2016, 16, e39909.	0.1	12
14	Genomic and protein structure analysis of the luciferase from the Iranian bioluminescent beetle, Luciola sp International Journal of Biological Macromolecules, 2019, 124, 689-698.	3.6	9
15	Impact of RGD Peptide Tethering to IL24/mda-7 (Melanoma Differentiation Associated Gene-7) on Apoptosis Induction in Hepatocellular Carcinoma Cells. Asian Pacific Journal of Cancer Prevention, 2015, 16, 6073-6080.	0.5	9
16	The characteristics of rare codon clusters in the genome and proteins of hepatitis C virus; a bioinformatics look. Middle East Journal of Digestive Diseases, 2014, 6, 214-27.	0.2	8
17	Identification of Lepidium draba î"1-pyrroline-5-carboxylate Synthetase (P5CS) and Assessment of its Expression Under NaCl stress: P5CS Identification in L. draba plant. Proceedings of the National Academy of Sciences India Section B - Biological Sciences, 2021, 91, 195-203.	0.4	6
18	The Effect of RGD/NGR Peptide Modification of Melanoma Differentiation-Associated Gene-7/Interleukin-24 on Its Receptor Attachment, an <i>In Silico</i> Analysis. Cancer Biotherapy and Radiopharmaceuticals, 2017, 32, 205-214.	0.7	5

#	Article	IF	CITATIONS
19	In silico analysis of codon usage and rare codon clusters in the halophilic bacteria L-asparaginase. Biologia (Poland), 2020, 75, 151-160.	0.8	5
20	Mutational Analysis of Ocriplasmin to Reduce Proteolytic and Autolytic Activity in Pichia pastoris. Biological Procedures Online, 2020, 22, 25.	1.4	5
21	Isolation, Identification and In Silico Study of Native Cellulase Producing Bacteria. Current Proteomics, 2021, 18, 3-11.	0.1	4
22	Evaluation of Luciferase Thermal Stability by Arginine Saturation in the Flexible Loops. Current Proteomics, 2020, 17, 30-39.	0.1	3
23	Thermally stable and acidic pH tolerant mutant phytases with high catalytic efficiency from Yersinia intermedia for potential application in feed industries. Environmental Science and Pollution Research, 2022, 29, 33713-33724.	2.7	2
24	The Evaluation of tLyP-1-Bound Mda-7/IL-24 Killing Activity on a Liver Tumor Cell Line. Cancer Biotherapy and Radiopharmaceuticals, 2020, 36, 827-836.	0.7	1
25	Assessment of E. Coli Expression System for Overexpression of Active Recombinant Ocriplasmin. Advanced Pharmaceutical Bulletin, 2021, 11, 564-569.	0.6	1
26	In Silico Analysis of Relative Rareness, Codon Usage, and Enzymesubstrate Docking of Lampyroidea Maculata luciferase. Current Proteomics, 2021, 18, 424-434.	0.1	0
27	<i>In Silico</i> Study of 1, 4 Alpha Glucan Branching Enzyme and Substrate Docking Studies. Current Proteomics, 2020, 17, 40-50.	0.1	O
28	<i>In silico</i> Evaluation of Substrate Binding Site and Rare Codons in the Structure of CYP152A1. Current Proteomics, 2020, 17, 10-22.	0.1	0
29	Engineering of Ocriplasmin Variants by Bioinformatics Methods for the Reduction of Proteolytic and Autolytic Activities. Iranian Journal of Medical Sciences, 2021, 46, 454-467.	0.3	0