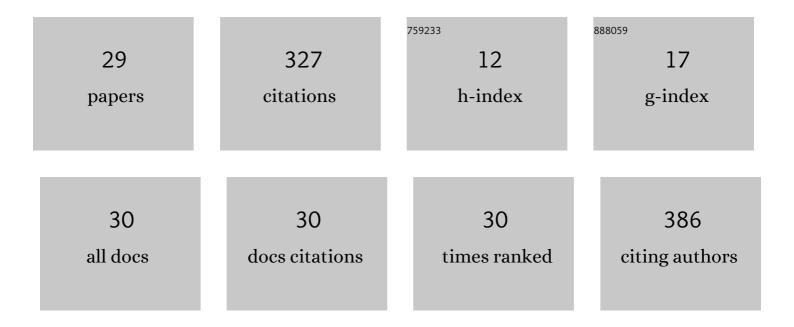
Mojtaba Mortazavi

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | 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. | 5.3 | 2 |
| 2 | 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. | 2.9 | 27 |
| 3 | Isolation, Identification and In Silico Study of Native Cellulase Producing Bacteria. Current Proteomics, 2021, 18, 3-11. | 0.3 | 4 |
| 4 | Identification of Lepidium draba Δ1-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. | 1.0 | 6 |
| 5 | Immobilization of Lepidium draba peroxidase on a novel Zn-MOF nanostructure. International Journal of Biological Macromolecules, 2021, 173, 366-378. | 7.5 | 13 |
| 6 | Enhancement of Thermostability of Aspergillus flavus Urate Oxidase by Immobilization on the Ni-Based Magnetic Metal–Organic Framework. Nanomaterials, 2021, 11, 1759. | 4.1 | 22 |
| 7 | In Silico Analysis of Relative Rareness, Codon Usage, and Enzymesubstrate Docking of Lampyroidea Maculata luciferase. Current Proteomics, 2021, 18, 424-434. | 0.3 | 0 |
| 8 | Assessment of E. Coli Expression System for Overexpression of Active Recombinant Ocriplasmin. Advanced Pharmaceutical Bulletin, 2021, 11, 564-569. | 1.4 | 1 |
| 9 | Engineering of Ocriplasmin Variants by Bioinformatics Methods for the Reduction of Proteolytic and Autolytic Activities. Iranian Journal of Medical Sciences, 2021, 46, 454-467. | 0.4 | 0 |
| 10 | In silico analysis of codon usage and rare codon clusters in the halophilic bacteria L-asparaginase. Biologia (Poland), 2020, 75, 151-160. | 1.5 | 5 |
| 11 | Mutational Analysis of Ocriplasmin to Reduce Proteolytic and Autolytic Activity in Pichia pastoris. Biological Procedures Online, 2020, 22, 25. | 2.9 | 5 |
| 12 | Design, synthesis and biological assessment of acridine derivatives containing 1,3,4-thiadiazole moiety as novel selective acetylcholinesterase inhibitors. Bioorganic Chemistry, 2020, 105, 104457. | 4.1 | 16 |
| 13 | 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. | 1.0 | 1 |
| 14 | Evaluation of Luciferase Thermal Stability by Arginine Saturation in the Flexible Loops. Current Proteomics, 2020, 17, 30-39. | 0.3 | 3 |
| 15 | <i>In Silico</i> Study of 1, 4 Alpha Glucan Branching Enzyme and Substrate Docking Studies. Current Proteomics, 2020, 17, 40-50. | 0.3 | 0 |
| 16 | <i>In silico</i> Evaluation of Substrate Binding Site and Rare Codons in the Structure of CYP152A1. Current Proteomics, 2020, 17, 10-22. | 0.3 | 0 |
| 17 | Genomic and protein structure analysis of the luciferase from the Iranian bioluminescent beetle, Luciola sp International Journal of Biological Macromolecules, 2019, 124, 689-698. | 7.5 | 9 |
| 18 | New Developments in Pichia pastoris Expression System, Review and Update. Current Pharmaceutical Biotechnology, 2018, 19, 451-467. | 1.6 | 35 |

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|----|--|-----|-----------|
| 19 | In Silico Analysis of L1/L2 Sequences of Human Papillomaviruses: Implication for Universal Vaccine Design. Viral Immunology, 2017, 30, 210-223. | 1.3 | 12 |
| 20 | Bifunctional role of leucine 300 of firefly luciferase in structural rigidity. International Journal of Biological Macromolecules, 2017, 101, 67-74. | 7.5 | 14 |
| 21 | 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. | 1.0 | 5 |
| 22 | Surface charge modification increases firefly luciferase rigidity without alteration in bioluminescence spectra. Enzyme and Microbial Technology, 2017, 96, 47-59. | 3.2 | 16 |
| 23 | 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. | 0.2 | 12 |
| 24 | 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. | 1.2 | 9 |
| 25 | 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. | 0.4 | 8 |
| 26 | Effects of 940 MHz EMF on luciferase solution: Structure, function, and dielectric studies. Bioelectromagnetics, 2013, 34, 489-498. | 1.6 | 15 |
| 27 | Design of thermostable luciferases through arginine saturation in solvent-exposed loops. Protein Engineering, Design and Selection, 2011, 24, 893-903. | 2.1 | 40 |
| 28 | Spectroscopic and functional characterization of <italic>Lampyris turkestanicus</italic> luciferase: a comparative study. Acta Biochimica Et Biophysica Sinica, 2008, 40, 365-374. | 2.0 | 22 |
| 29 | cDNA Cloning, Expression and Homology Modeling of a Luciferase from the Firefly Lampyroidea maculata. BMB Reports, 2006, 39, 578-585. | 2.4 | 18 |