List of Publications by Year in descending order

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YONG-DOO PARK

#	Article	IF	CITATIONS
1	Effect of hesperetin on tyrosinase: Inhibition kinetics integrated computational simulation study. International Journal of Biological Macromolecules, 2012, 50, 257-262.	3.6	90
2	Tyrosinase inhibition by isophthalic acid: Kinetics and computational simulation. International Journal of Biological Macromolecules, 2011, 48, 700-704.	3.6	68
3	Inhibitory effect of hesperetin on α-glucosidase: Molecular dynamics simulation integrating inhibition kinetics. International Journal of Biological Macromolecules, 2017, 101, 32-39.	3.6	51
4	Towards a proteomic analysis of atopic dermatitis: A two-dimensional-polyacrylamide gel electrophoresis/mass spectrometric analysis of cultured patient-derived fibroblasts. Proteomics, 2004, 4, 3446-3455.	1.3	45
5	Metabolic responses and arginine kinase expression of juvenile cuttlefish (Sepia pharaonis) under salinity stress. International Journal of Biological Macromolecules, 2018, 113, 881-888.	3.6	33
6	Mixed-Type Inhibition of Tyrosinase from Agaricus bisporus by Terephthalic Acid: Computational Simulations and Kinetics. Protein Journal, 2011, 30, 273-280.	0.7	31
7	Inhibitory effect of raspberry ketone on α-glucosidase: Docking simulation integrating inhibition kinetics. International Journal of Biological Macromolecules, 2018, 113, 212-218.	3.6	30
8	The inhibitory effect of pyrogallol on tyrosinase activity and structure: Integration study of inhibition kinetics with molecular dynamics simulation. International Journal of Biological Macromolecules, 2019, 121, 463-471.	3.6	27
9	A new type of uncompetitive inhibition of tyrosinase induced by Cl– binding. Biochimie, 2005, 87, 931-937.	1.3	25
10	Profiling the dysregulated genes of keratinocytes in atopic dermatitis patients: cDNA microarray and interactomic analyses. Journal of Dermatological Science, 2009, 54, 126-129.	1.0	24
11	Two-dimensional electrophoretic profiling of atopic dermatitis in primary cultured keratinocytes from patients. Proteomics, 2006, 6, 1362-1370.	1.3	22
12	Inhibitory effect of pyrogallol on α-glucosidase: Integrating docking simulations with inhibition kinetics. International Journal of Biological Macromolecules, 2018, 112, 686-693.	3.6	21
13	Detection of down-regulated acetaldehyde dehydrogenase 1 in atopic dermatitis patients by two-dimensional electrophoresis. Experimental Dermatology, 2007, 16, 130-134.	1.4	19
14	Effects of osmolytes on arginine kinase from Euphausia superba: A study on thermal denaturation and aggregation. Process Biochemistry, 2014, 49, 936-947.	1.8	19
15	A folding study of Antarctic krill (Euphausia superba) alkaline phosphatase using denaturants. International Journal of Biological Macromolecules, 2014, 70, 266-274.	3.6	19
16	Two-dimensional electrophoresis analyses of atopic dermatitis and the chances to detect new candidate proteins by the variations in immobilized pH gradient strips. Journal of Dermatological Science, 2007, 47, 9-17.	1.0	17
17	Effects of osmolytes on Pelodiscus sinensis creatine kinase: A study on thermal denaturation and aggregation. International Journal of Biological Macromolecules, 2013, 60, 277-287.	3.6	17
18	Inhibitory effect of phloroglucinol on α-glucosidase: Kinetics and molecular dynamics simulation integration study. International Journal of Biological Macromolecules, 2019, 124, 771-779.	3.6	17

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19	Proteomic approaches to the analysis of atopic dermatitis and new insights from interactomics. Proteomics - Clinical Applications, 2008, 2, 290-300.	0.8	15
20	Purification, characterization, and unfolding studies of arginine kinase from Antarctic krill. International Journal of Biological Macromolecules, 2014, 67, 426-432.	3.6	15
21	TXM13 human melanoma cells: a novel source for the inhibition kinetics of human tyrosinase and for screening whitening agents. Biochemistry and Cell Biology, 2006, 84, 112-116.	0.9	14
22	The effect of Zn2+ on Euphausia superba arginine kinase: Unfolding and aggregation studies. Process Biochemistry, 2014, 49, 821-829.	1.8	14
23	Dysregulated genes of extrinsic type of atopic dermatitis: 34K microarray and interactomic analyses. Journal of Dermatological Science, 2009, 53, 146-150.	1.0	13
24	DNA microarray analyses and interactomic predictions for atopic dermatitis. Journal of Dermatological Science, 2009, 55, 123-125.	1.0	10
25	Thermal stable characteristics of acid- and pepsin-soluble collagens from the carapace tissue of Chinese soft-shelled turtle (Pelodiscus sinensis). Tissue and Cell, 2020, 67, 101424.	1.0	10
26	Combination of free-flow electrophoresis and interactomics to analyze the dysregulated proteins of fibroblasts from atopic dermatitis patients. Journal of Dermatological Science, 2011, 61, 148-150.	1.0	9
27	Inhibition of α-glucosidase by 2-thiobarbituric acid: Molecular dynamics simulation integrating parabolic noncompetitive inhibition kinetics. Process Biochemistry, 2018, 65, 62-70.	1.8	9
28	Kinetics for Cu 2+ induced Sepia pharaonis arginine kinase inactivation and aggregation. International Journal of Biological Macromolecules, 2016, 91, 926-933.	3.6	8
29	Effect of Cd2+ on tyrosinase: Integration of inhibition kinetics with computational simulation. International Journal of Biological Macromolecules, 2017, 94, 836-844.	3.6	8
30	Proteomic analyses for profiling regulated proteins/enzymes by Fucus vesiculosus fucoidan in B16 melanoma cells: A combination of enzyme kinetics functional study. International Journal of Biological Macromolecules, 2018, 112, 667-674.	3.6	8
31	Analysis of the peptides detected in atopic dermatitis and various inflammatory diseases patients-derived sera. International Journal of Biological Macromolecules, 2018, 106, 1052-1061.	3.6	8
32	Effects of Cu2+ on alkaline phosphatase from Macrobrachium rosenbergii. International Journal of Biological Macromolecules, 2018, 117, 116-123.	3.6	8
33	Comparative studies of the expression of creatine kinase isoforms under immune stress in Pelodiscus sinensis. International Journal of Biological Macromolecules, 2020, 162, 11-23.	3.6	8
34	The Inhibitory Effects of Cu2+ on Exopalaemon carinicauda Arginine Kinase via Inhibition Kinetics and Molecular Dynamics Simulations. Applied Biochemistry and Biotechnology, 2015, 176, 1217-1236.	1.4	7
35	Tyrosinase-mediated melanogenesis in melanoma cells: Array comparative genome hybridization integrating proteomics and bioinformatics studies. International Journal of Biological Macromolecules, 2021, 170, 150-163.	3.6	7
36	Effect of Cysteine Modification on Creatine Kinase Aggregation. Applied Biochemistry and Biotechnology, 2009, 152, 15-28.	1.4	6

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37	Folding Studies of Arginine Kinase from Euphausia superba Using Denaturants. Applied Biochemistry and Biotechnology, 2014, 172, 3888-3901.	1.4	6
38	The effect of Cu2+ on arginine kinase from Euphausia superba: A computational simulation integrating unfolding and aggregation studies. Process Biochemistry, 2015, 50, 395-405.	1.8	6
39	Functional study of 14-3-3 protein epsilon (YWHAE) in keratinocytes: microarray integrating bioinformatics approaches. Journal of Biomolecular Structure and Dynamics, 2020, 38, 2633-2649.	2.0	6
40	Computational prediction for the protein interactions of tyrosinase: Protein experimental interactome MAP. Process Biochemistry, 2013, 48, 638-648.	1.8	5
41	Serum proteomic analyses for probing C3 fragment protein. Process Biochemistry, 2016, 51, 981-988.	1.8	5
42	Hydrogen peroxide (H2O2) irreversibly inactivates creatine kinase from Pelodiscus sinensis by targeting the active site cysteine. International Journal of Biological Macromolecules, 2017, 105, 1595-1601.	3.6	5
43	An OMICS-based study of the role of C3dg in keratinocytes: RNA sequencing, antibody-chip array, and bioinformatics approaches. International Journal of Biological Macromolecules, 2019, 133, 391-411.	3.6	5
44	Towards creatine kinase aggregation due to the cysteine modification at the flexible active site and refolding pathway. International Journal of Biological Macromolecules, 2007, 41, 439-446.	3.6	4
45	An integrated method for the detection of basic proteins in serum-derived proteomes. Process Biochemistry, 2016, 51, 973-980.	1.8	3
46	The omics based study for the role of superoxide dismutase 2 (SOD2) in keratinocytes: RNA sequencing, antibody-chip array and bioinformatics approaches. Journal of Biomolecular Structure and Dynamics, 2020, 38, 2884-2897.	2.0	3
47	Trifluoroethanol-Induced Changes in Activity and Conformation of Manganese-Containing Superoxide Dismutase. Applied Biochemistry and Biotechnology, 2012, 166, 276-288.	1.4	2
48	Inhibitory effect of α-ketoglutaric acid on α-glucosidase: integrating molecular dynamics simulation and inhibition kinetics. Journal of Biomolecular Structure and Dynamics, 2020, 38, 3496-3503.	2.0	2
49	Functional study of acetaldehyde dehydrogenase 1 (ALDH1) in keratinocytes: microarray integrating bioinformatics approaches. Journal of Biomolecular Structure and Dynamics, 2021, 39, 2133-2151.	2.0	2
50	An RNA interference based study for the role of ALDH1 in keratinocytes: DNA microarray, antibody–chip array and bioinformatics approaches. Process Biochemistry, 2014, 49, 1612-1621.	1.8	1
51	Are Podoplanin Gene Polymorphisms Associated with Atopic Dermatitis in Koreans?. Annals of Dermatology, 2015, 27, 275.	0.3	1
52	Seasonal expression of cytoplasmic creatine kinase in the epididymal epithelium of <i>Pelodiscus sinensis</i> . Biotechnic and Histochemistry, 2022, 97, 21-29.	0.7	1
53	A strategy to improve the efficiency of spermatogenesis during the breeding seasons of <i>Pelodiscus sinensis</i> . Journal of the World Aquaculture Society, 0, , .	1.2	1
54	Characterization and tissue expression analysis of mitochondrial creatine kinases (types I and II) from <i>Pelodiscus sinensis</i> . Journal of Biomolecular Structure and Dynamics, 2023, 41, 1388-1402.	2.0	1

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55	A computational integrating kinetic study on the flexible active site of human acetaldehyde dehydrogenase 1. Process Biochemistry, 2016, 51, 725-733.	1.8	0
56	Integration of Inhibition Kinetics and Molecular Dynamics Simulations: A Urea-Mediated Folding Study on Acetaldehyde Dehydrogenase 1. Applied Biochemistry and Biotechnology, 2016, 179, 1101-1114.	1.4	0
57	Effects of osmolytes on the refolding of recombinant Pelodiscus sinensis brain-type creatine kinase. Process Biochemistry, 2018, 68, 83-92.	1.8	0
58	Screening and analysis of agouti signaling protein interaction partners in Pelodiscus sinensis suggests a role in lipid metabolism. International Journal of Biological Macromolecules, 2020, 157, 695-705.	3.6	0
59	A Knock-Down Cell-Based Study for the Functional Analysis of Chloride Intracellular Channel 1 (CLIC1): Integrated Proteomics and Microarray Study. Protein and Peptide Letters, 2021, 28, 84-100.	0.4	0