

Jesus Mendieta

List of Publications by Year in descending order

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53
papers

1,278
citations

279487

23
h-index

377514

34
g-index

55
all docs

55
docs citations

55
times ranked

1518
citing authors

#	ARTICLE	IF	CITATIONS
1	Proton Transfer in Guanine-Cytosine Base Pairs in B-DNA. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 6984-6991.	2.3	21
2	Amino acid residues in HIV-2 reverse transcriptase that restrict the development of nucleoside analogue resistance through the excision pathway. <i>Journal of Biological Chemistry</i> , 2018, 293, 2247-2259.	1.6	9
3	Effect of water-DNA interactions on elastic properties of DNA self-assembled monolayers. <i>Scientific Reports</i> , 2017, 7, 536.	1.6	33
4	Two-step ATP-driven opening of cohesin head. <i>Scientific Reports</i> , 2017, 7, 3266.	1.6	19
5	Quantum Mechanics/Molecular Mechanics Free Energy Maps and Nonadiabatic Simulations for a Photochemical Reaction in DNA: Cyclobutane Thymine Dimer. <i>Journal of Physical Chemistry Letters</i> , 2016, 7, 4391-4397.	2.1	20
6	A Practical Quantum Mechanics Molecular Mechanics Method for the Dynamical Study of Reactions in Biomolecules. <i>Advances in Protein Chemistry and Structural Biology</i> , 2015, 100, 67-88.	1.0	5
7	MEPSA: minimum energy pathway analysis for energy landscapes. <i>Bioinformatics</i> , 2015, 31, 3853-3855.	1.8	84
8	Torsion and curvature of FtsZ filaments. <i>Soft Matter</i> , 2014, 10, 1977.	1.2	27
9	<scp>fireball</scp>/<scp>amber</scp>: An Efficient Local-Orbital DFT QM/MM Method for Biomolecular Systems. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 2185-2193.	2.3	42
10	Molecular basis of the association of H208Y and thymidine analogue resistance mutations M41L, L210W and T215Y in the HIV-1 reverse transcriptase of treated patients. <i>Antiviral Research</i> , 2014, 106, 42-52.	1.9	3
11	Simulation of Catalytic Water Activation in Mitochondrial F ₁ -ATPase Using a Hybrid Quantum Mechanics/Molecular Mechanics Approach: An Alternative Role for Î ² -Glu 188. <i>Biochemistry</i> , 2013, 52, 959-966.	1.2	6
12	Functional Specificity of a Protein-DNA Complex Mediated by Two Arginines Bound to the Minor Groove. <i>Journal of Bacteriology</i> , 2012, 194, 4727-4735.	1.0	4
13	Molecular dynamics analysis of conformational change of paramyxovirus F protein during the initial steps of membrane fusion. <i>Biochemical and Biophysical Research Communications</i> , 2012, 420, 42-47.	1.0	2
14	The Role of Gln61 in HRas GTP Hydrolysis: A Quantum Mechanics/Molecular Mechanics Study. <i>Biophysical Journal</i> , 2012, 102, 152-157.	0.2	48
15	Molecular dynamics simulation of GTPase activity in polymers of the cell division protein FtsZ. <i>FEBS Letters</i> , 2012, 586, 1236-1239.	1.3	9
16	Functional phosphoproteomics for current immunology research. <i>Journal of Integrated OMICS</i> , 2011, 1, .	0.5	3
17	Technical phosphoproteomic and bioinformatic tools useful in cancer research. <i>Journal of Clinical Bioinformatics</i> , 2011, 1, 26.	1.2	16
18	Nanomechanics of the Cadherin Ectodomain. <i>Journal of Biological Chemistry</i> , 2011, 286, 9405-9418.	1.6	45

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19	Thymidine Analogue Excision and Discrimination Modulated by Mutational Complexes Including Single Amino Acid Deletions of Asp-67 or Thr-69 in HIV-1 Reverse Transcriptase. <i>Journal of Biological Chemistry</i> , 2011, 286, 20615-20624.	1.6	14
20	Structural and Functional Model for Ionic (K ⁺ /Na ⁺) and pH Dependence of GTPase Activity and Polymerization of FtsZ, the Prokaryotic Ortholog of Tubulin. <i>Journal of Molecular Biology</i> , 2009, 390, 17-25.	2.0	38
21	A Mg ²⁺ -induced conformational switch rendering a competent DNA polymerase catalytic complex. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 71, 565-574.	1.5	28
22	Mechanistic Basis of Zidovudine Hypersusceptibility and Lamivudine Resistance Conferred by the Deletion of Codon 69 in the HIV-1 Reverse Transcriptase Coding Region. <i>Journal of Molecular Biology</i> , 2008, 382, 327-341.	2.0	18
23	Residues in human respiratory syncytial virus P protein that are essential for its activity on RNA viral synthesis. <i>Virus Research</i> , 2008, 132, 160-173.	1.1	20
24	Use of a Dominant rpsL Allele Conferring Streptomycin Dependence for Positive and Negative Selection in <i>Thermus thermophilus</i> . <i>Applied and Environmental Microbiology</i> , 2007, 73, 5138-5145.	1.4	14
25	DNA sequence-specific recognition by a transcriptional regulator requires indirect readout of A-tracts. <i>Nucleic Acids Research</i> , 2007, 35, 3252-3261.	6.5	10
26	Binding of 5'-GMP to the GluR2 AMPA Receptor: An Insight from Targeted Molecular Dynamics Simulations. <i>Biochemistry</i> , 2005, 44, 14470-14476.	1.2	28
27	Phosphorylation modulates the alpha-helical structure and polymerization of a peptide from the third tau microtubule-binding repeat. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2005, 1721, 16-26.	1.1	22
28	In silico activation of Src tyrosine kinase reveals the molecular basis for intramolecular autophosphorylation. <i>Journal of Molecular Graphics and Modelling</i> , 2004, 23, 189-198.	1.3	21
29	Role of Histidine-85 in the Catalytic Mechanism of Thymidine Phosphorylase As Assessed by Targeted Molecular Dynamics Simulations and Quantum Mechanical Calculations. <i>Biochemistry</i> , 2004, 43, 405-414.	1.2	34
30	Towards New Thymidine Phosphorylase/PD-ECGF Inhibitors Based on the Transition State of the Enzyme Reaction. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2003, 22, 951-953.	0.4	3
31	A 3'-ET743-DNA Complex That Both Resembles an RNA-DNA Hybrid and Mimicks Zinc Finger-Induced DNA Structural Distortions. <i>Journal of Medicinal Chemistry</i> , 2002, 45, 871-880.	2.9	30
32	Combined Use of Differential Pulse Polarography and Multivariate Curve Resolution: As Applied to the Study of Metal Mixed Complexes of the Metallothionein Related Hexapeptide. <i>Electroanalysis</i> , 2002, 14, 50-56.	1.5	10
33	Molecular dynamics simulations of the conformational changes of the glutamate receptor ligand-binding core in the presence of glutamate and kainate. <i>Proteins: Structure, Function and Bioinformatics</i> , 2001, 44, 460-469.	1.5	37
34	Soft- and Hard-Modeling Approaches for the Determination of Stability Constants of Metal-Peptide Systems by Voltammetry. <i>Analytical Biochemistry</i> , 2000, 279, 189-201.	1.1	41
35	Zinc-binding properties of the C-terminal hexapeptide Lys-Cys-Thr-Cys-Cys-Ala from mouse metallothionein: analysis by differential pulse polarography and multivariate curve resolution. <i>Analytica Chimica Acta</i> , 1999, 385, 353-363.	2.6	17
36	Complexation of cadmium by the C-terminal hexapeptide Lys-Cys-Thr-Cys-Cys-Ala from mouse metallothionein: study by differential pulse polarography and circular dichroism spectroscopy with multivariate curve resolution analysis. <i>Analytica Chimica Acta</i> , 1999, 390, 15-25.	2.6	27

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37	Multivariate Curve Resolution of Cyclic Voltammetric Data: Application to the Study of the Cadmium-Binding Properties of Glutathione. <i>Analytical Chemistry</i> , 1999, 71, 4629-4636.	3.2	53
38	Two-Phase Induction of the Nonnative $\hat{1}\pm$ -Helical Form of $\hat{1}^2$ -Lactoglobulin in the Presence of Trifluoroethanol. <i>Biophysical Journal</i> , 1999, 76, 451-457.	0.2	38
39	Study of the zinc-binding properties of glutathione by differential pulse polarography and multivariate curve resolution. <i>Journal of Inorganic Biochemistry</i> , 1998, 70, 91-98.	1.5	58
40	Multivariate Curve Resolution: A Possible Tool in the Detection of Intermediate Structures in Protein Folding. <i>Biophysical Journal</i> , 1998, 74, 2876-2888.	0.2	53
41	Cadmium-binding properties of glutathione: A chemometrical analysis of voltammetric data. <i>Journal of Inorganic Biochemistry</i> , 1997, 66, 29-36.	1.5	77
42	Electrochemical study of the binding properties of a metallothionein I related peptide with cadmium or/and zinc. <i>Electroanalysis</i> , 1996, 8, 473-479.	1.5	25
43	Application of Multivariate Curve Resolution to Voltammetric Data. <i>Analytical Biochemistry</i> , 1996, 240, 134-141.	1.1	74
44	The electrochemical behaviour of Cd,Zn thioneins depending on the solution pH using differential pulse polarography. <i>Analytica Chimica Acta</i> , 1995, 305, 285-294.	2.6	32
45	Electrochemical behavior of metallothioneins and related molecules. Part I: Lys-Cys-Thr-Cys-Cys-Ala thionein fragment [56-61] MT I. <i>Electroanalysis</i> , 1995, 7, 663-669.	1.5	33
46	Transferrin-Binding Capacity by Rat Bone Marrow Populations Containing Different Proportions of Erythroid Cells. <i>Biological Chemistry Hoppe-Seyler</i> , 1994, 375, 135-140.	1.4	0
47	Transferrin binding capacity as a marker of differentiation and maturation of rat erythroid cells fractionated by counter current distribution in aqueous polymer two-phase systems. <i>Bioscience Reports</i> , 1994, 14, 119-130.	1.1	0
48	[33] Charge-directed affinity partitioning of cells. <i>Methods in Enzymology</i> , 1994, 228, 363-368.	0.4	0
49	Fractionation of erythroblasts with affinity-mediated modifications of their electrical properties using counter-current distribution. <i>Molecular and Cellular Biochemistry</i> , 1993, 121, 93-98.	1.4	0
50	Rat bone marrow erythroid cell fractionation by counter current distribution in non-charge-sensitive two-phase systems. <i>Bioscience Reports</i> , 1992, 12, 77-85.	1.1	1
51	Ligand-receptor interactions in affinity cell partitioning. <i>Journal of Chromatography A</i> , 1992, 594, 97-103.	1.8	11
52	Affinity-mediated modification of electrical charge on a cell surface: A new approach to the affinity partitioning of biological particles. <i>Analytical Biochemistry</i> , 1992, 200, 280-285.	1.1	5
53	Analysis by partitioning in aqueous two-phase systems of the loss of transferrin-binding capacity during maturation of rat reticulocytes. <i>Bioscience Reports</i> , 1989, 9, 541-548.	1.1	10