Christopher R Pudney

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/579592/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	On the Temperature Dependence of Enzyme-Catalyzed Rates. Biochemistry, 2016, 55, 1681-1688.	2.5	233
2	Biocatalysis with Thermostable Enzymes: Structure and Properties of a Thermophilic â€~ene'â€Reductase related to Old Yellow Enzyme. ChemBioChem, 2010, 11, 197-207.	2.6	110
3	Fast Protein Motions Are Coupled to Enzyme H-Transfer Reactions. Journal of the American Chemical Society, 2013, 135, 2512-2517.	13.7	83
4	Evidence To Support the Hypothesis That Promoting Vibrations Enhance the Rate of an Enzyme Catalyzed H-Tunneling Reaction. Journal of the American Chemical Society, 2009, 131, 17072-17073.	13.7	79
5	Direct Analysis of Donorâ^'Acceptor Distance and Relationship to Isotope Effects and the Force Constant for Barrier Compression in Enzymatic H-Tunneling Reactions. Journal of the American Chemical Society, 2010, 132, 11329-11335.	13.7	74
6	α-Secondary Isotope Effects as Probes of "Tunneling-Ready―Configurations in Enzymatic H-Tunneling:Â Insight from Environmentally Coupled Tunneling Models. Journal of the American Chemical Society, 2006, 128, 14053-14058.	13.7	66
7	Mutagenesis of Morphinone Reductase Induces Multiple Reactive Configurations and Identifies Potential Ambiguity in Kinetic Analysis of Enzyme Tunneling Mechanisms. Journal of the American Chemical Society, 2007, 129, 13949-13956.	13.7	55
8	Enzyme evolution and the temperature dependence of enzyme catalysis. Current Opinion in Structural Biology, 2020, 65, 96-101.	5.7	54
9	Barrier Compression Enhances an Enzymatic Hydrogenâ€īransfer Reaction. Angewandte Chemie - International Edition, 2009, 48, 1452-1454.	13.8	52
10	Coupled Motions Direct Electrons along Human Microsomal P450 Chains. PLoS Biology, 2011, 9, e1001222.	5.6	48
11	The red edge excitation shift phenomenon can be used to unmask protein structural ensembles: implications for NEMO–ubiquitin interactions. FEBS Journal, 2016, 283, 2272-2284.	4.7	44
12	A complete thermodynamic analysis of enzyme turnover links the free energy landscape to enzyme catalysis. FEBS Journal, 2017, 284, 2829-2842.	4.7	39
13	Are Environmentally Coupled Enzymatic Hydrogen Tunneling Reactions Influenced by Changes in Solution Viscosity?. Angewandte Chemie - International Edition, 2008, 47, 537-540.	13.8	34
14	Gating mechanisms for biological electron transfer: Integrating structure with biophysics reveals the nature of redox control in cytochrome P450 reductase and copperâ€dependent nitrite reductase. FEBS Letters, 2012, 586, 578-584.	2.8	31
15	Structural and mechanistic aspects of flavoproteins: probes of hydrogen tunnelling. FEBS Journal, 2009, 276, 3930-3941.	4.7	27
16	Bipartite recognition and conformational sampling mechanisms for hydride transfer from nicotinamide coenzyme to FMN in pentaerythritol tetranitrate reductase. FEBS Journal, 2009, 276, 4780-4789.	4.7	24
17	Polyubiquitin Drives the Molecular Interactions of the NF-κB Essential Modulator (NEMO) by Allosteric Regulation. Journal of Biological Chemistry, 2015, 290, 14130-14139.	3.4	23
18	Parallel Pathways and Freeâ€Energy Landscapes for Enzymatic Hydride Transfer Probed by Hydrostatic Pressure. ChemBioChem, 2009, 10, 1379-1384.	2.6	22

#	Article	IF	CITATIONS
19	Atomistic insight into the origin of the temperature-dependence of kinetic isotope effects and H-tunnelling in enzyme systems is revealed through combined experimental studies and biomolecular simulation. Biochemical Society Transactions, 2008, 36, 16-21.	3.4	21
20	Enzymatic Single-Molecule Kinetic Isotope Effects. Journal of the American Chemical Society, 2013, 135, 3855-3864.	13.7	21
21	Steady-State Kinetics of α-Synuclein Ferrireductase Activity Identifies the Catalytically Competent Species. Biochemistry, 2017, 56, 2497-2505.	2.5	21
22	Conformation control of the histidine kinase BceS of <i>Bacillus subtilis</i> by its cognate ABCâ€ŧransporter facilitates needâ€based activation of antibiotic resistance. Molecular Microbiology, 2021, 115, 157-174.	2.5	20
23	Sensing Enzyme Activation Heat Capacity at the Single-Molecule Level Using Gold-Nanorod-Based Optical Whispering Gallery Modes. ACS Applied Nano Materials, 2021, 4, 4576-4583.	5.0	20
24	Change in heat capacity accurately predicts vibrational coupling in enzyme catalyzed reactions. FEBS Letters, 2015, 589, 2200-2206.	2.8	19
25	Kinetic and spectroscopic probes of motions and catalysis in the cytochrome P450 reductase family of enzymes. FEBS Journal, 2012, 279, 1534-1544.	4.7	18
26	Excited State Dynamics Can Be Used to Probe Donor-Acceptor Distances for H-Tunneling Reactions Catalyzed by Flavoproteins. Biophysical Journal, 2013, 105, 2549-2558.	0.5	17
27	Solvent as a Probe of Active Site Motion and Chemistry during the Hydrogen Tunnelling Reaction in Morphinone Reductase. ChemPhysChem, 2008, 9, 1875-1881.	2.1	16
28	Probing active site geometry using high pressure and secondary isotope effects in an enzymeâ€eatalysed â€~deep' Hâ€ŧunnelling reaction. Journal of Physical Organic Chemistry, 2010, 23, 696-701.	1.9	16
29	Rigidifying a <i>De Novo</i> Enzyme Increases Activity and Induces a Negative Activation Heat Capacity. ACS Catalysis, 2021, 11, 11532-11541.	11.2	15
30	Uncovering the Relationship between the Change in Heat Capacity for Enzyme Catalysis and Vibrational Frequency through Isotope Effect Studies. ACS Catalysis, 2018, 8, 5340-5349.	11.2	13
31	Synthetic Cannabinoid Receptor Agonists Detection Using Fluorescence Spectral Fingerprinting. Analytical Chemistry, 2019, 91, 12971-12979.	6.5	13
32	Monoclonal antibody stability can be usefully monitored using the excitation-energy-dependent fluorescence edge-shift. Biochemical Journal, 2020, 477, 3599-3612.	3.7	13
33	Exposing the Interplay Between Enzyme Turnover, Protein Dynamics, and the Membrane Environment in Monoamine Oxidase B. Biochemistry, 2019, 58, 2362-2372.	2.5	12
34	Analysis of synthetic cannabinoid agonists and their degradation products after combustion in a smoking simulator. Analytical Methods, 2019, 11, 3101-3107.	2.7	10
35	Structure and <i>in silico</i> simulations of a cold-active esterase reveals its prime cold-adaptation mechanism. Open Biology, 2021, 11, 210182.	3.6	10
36	Correction of Pre-Steady-State KIEs for Isotopic Impurities and the Consequences of Kinetic Isotope Fractionation. Journal of Physical Chemistry A, 2008, 112, 13109-13115.	2.5	9

#	Article	IF	CITATIONS
37	Molecular Rules Underpinning Enhanced Affinity Binding of Human T Cell Receptors Engineered for Immunotherapy. Molecular Therapy - Oncolytics, 2020, 18, 443-456.	4.4	9
38	Excitation-Energy-Dependent Molecular Beacon Detects Early Stage Neurotoxic AÎ ² Aggregates in the Presence of Cortical Neurons. ACS Chemical Neuroscience, 2019, 10, 1240-1250.	3.5	8
39	Reliable <i>In Silico</i> Ranking of Engineered Therapeutic TCR Binding Affinities with MMPB/GBSA. Journal of Chemical Information and Modeling, 2022, 62, 577-590.	5.4	8
40	Modelling flavoenzymatic charge transfer events: development of catalytic indole deuteration strategies. Organic and Biomolecular Chemistry, 2016, 14, 3787-3792.	2.8	6
41	Peptide cargo tunes a network of correlated motions in human leucocyte antigens. FEBS Journal, 2020, 287, 3777-3793.	4.7	6
42	Practical Aspects on the Use of Kinetic Isotope Effects as Probes of Flavoprotein Enzyme Mechanisms. Methods in Molecular Biology, 2014, 1146, 161-175.	0.9	6
43	Chemical Mapping Exposes the Importance of Active Site Interactions in Governing the Temperature Dependence of Enzyme Turnover. ACS Catalysis, 2021, 11, 14854-14863.	11.2	6
44	Switching protein metalloporphyrin binding specificity by design from iron to fluorogenic zinc. Chemical Communications, 2020, 56, 4308-4311.	4.1	4
45	Flavin mimetics: Synthesis and photophysical properties. Tetrahedron, 2021, 82, 131925.	1.9	3