Hideaki Shimizu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7264223/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	PINK1 autophosphorylation upon membrane potential dissipation is essential for Parkin recruitment to damaged mitochondria. Nature Communications, 2012, 3, 1016.	5.8	465
2	Semiâ€rational engineering of a coral fluorescent protein into an efficient highlighter. EMBO Reports, 2005, 6, 233-238.	2.0	320
3	Crystal structure of nitric oxide reductase from denitrifying fungus Fusarium oxysporum. Nature Structural Biology, 1997, 4, 827-832.	9.7	172
4	Crystal Structure of an Active Form of BACE1, an Enzyme Responsible for Amyloid β Protein Production. Molecular and Cellular Biology, 2008, 28, 3663-3671.	1.1	167
5	High-Resolution Crystal Structures and Spectroscopy of Native and Compound I CytochromecPeroxidaseâ€. Biochemistry, 2003, 42, 5600-5608.	1.2	140
6	The Novel Binding Mode of N-Alkyl-Nâ€~-hydroxyguanidine to Neuronal Nitric Oxide Synthase Provides Mechanistic Insights into NO Biosynthesis. Biochemistry, 2002, 41, 13868-13875.	1.2	122
7	RNA-binding Protein TLS Is a Major Nuclear Aggregate-interacting Protein in Huntingtin Exon 1 with Expanded Polyglutamine-expressing Cells. Journal of Biological Chemistry, 2008, 283, 6489-6500.	1.6	109
8	Comparison of the Heme-free and -bound Crystal Structures of Human Heme Oxygenase-1. Journal of Biological Chemistry, 2003, 278, 7834-7843.	1.6	104
9	Proton Delivery in NO Reduction by Fungal Nitric-oxide Reductase. Journal of Biological Chemistry, 2000, 275, 4816-4826.	1.6	100
10	Crystal Structure ofNitrosomonas europaeaCytochromecPeroxidase and the Structural Basis for Ligand Switching in Bacterial Di-heme Peroxidasesâ€. Biochemistry, 2001, 40, 13483-13490.	1.2	83
11	Crystal Structures of Epothilone D-bound, Epothilone B-bound, and Substrate-free Forms of Cytochrome P450epoK. Journal of Biological Chemistry, 2003, 278, 44886-44893.	1.6	75
12	Membrane microdomain switching: a regulatory mechanism of amyloid precursor protein processing. Journal of Cell Biology, 2008, 183, 339-352.	2.3	61
13	Singular localization of sodium channel β4 subunit in unmyelinated fibres and its role in the striatum. Nature Communications, 2014, 5, 5525.	5.8	61
14	The E1 Mechanism in Photo-Induced β-Elimination Reactions for Green-to-Red Conversion of Fluorescent Proteins. Chemistry and Biology, 2009, 16, 1140-1147.	6.2	56
15	The RIKEN structural biology beamline II (BL44B2) at the SPring-8. Nuclear Instruments and Methods in Physics Research, Section A: Accelerators, Spectrometers, Detectors and Associated Equipment, 2001, 467-468, 711-714.	0.7	50
16	Discovery of a small molecule inhibitor targeting dengue virus NS5 RNA-dependent RNA polymerase. PLoS Neglected Tropical Diseases, 2019, 13, e0007894.	1.3	49
17	Crystal structures of cytochrome P450nor and its mutants (Ser286→Val, Thr) in the ferric resting state at cryogenic temperature: a comparative analysis with monooxygenase cytochrome P450s. Journal of Inorganic Biochemistry, 2000, 81, 191-205.	1.5	45
18	A Novel Heme and Peroxide-dependent Tryptophan–tyrosine Cross-link in a Mutant of Cytochrome c Peroxidase. Journal of Molecular Biology, 2003, 328, 157-166.	2.0	39

HIDEAKI SHIMIZU

#	Article	IF	CITATIONS
19	Structure of cytochromec6from the red algaPorphyra yezoensisat 1.57â€Ã resolution. Acta Crystallographica Section D: Biological Crystallography, 2000, 56, 1577-1582.	2.5	24
20	Mutation effects of a conserved threonine (Thr243) of cytochrome P450nor on its structure and function. Journal of Inorganic Biochemistry, 2000, 82, 103-111.	1.5	22
21	A molecular mechanism realizing sequence-specific recognition of nucleic acids by TDP-43. Scientific Reports, 2016, 6, 20576.	1.6	22
22	X-ray structure of nitric oxide reductase (cytochrome P450nor) at atomic resolution. Acta Crystallographica Section D: Biological Crystallography, 2002, 58, 81-89.	2.5	21
23	Comparative functional analysis of CYP71AV1 natural variants reveals an important residue for the successive oxidation of amorphaâ€4,11â€diene. FEBS Letters, 2013, 587, 278-284.	1.3	21
24	Crystallization, preliminary diffraction and electron paramagnetic resonance studies of a single crystal of cytochromeP450nor. FEBS Letters, 1997, 412, 346-350.	1.3	19
25	Parallel homodimer structures of the extracellular domains of the voltage-gated sodium channel β4 subunit explain its role in cell–cell adhesion. Journal of Biological Chemistry, 2017, 292, 13428-13440.	1.6	16
26	Structure-based site-directed photo-crosslinking analyses of multimeric cell-adhesive interactions of voltage-gated sodium channel β subunits. Scientific Reports, 2016, 6, 26618.	1.6	13
27	Acidic Chitinase-Chitin Complex Is Dissociated in a Competitive Manner by Acetic Acid: Purification of Natural Enzyme for Supplementation Purposes. International Journal of Molecular Sciences, 2018, 19, 362.	1.8	12
28	Aggregation mechanism of polyglutamine diseases revealed using quantum chemical calculations, fragment molecular orbital calculations, molecular dynamics simulations, and binding free energy calculations. Computational and Theoretical Chemistry, 2006, 778, 85-95.	1.5	9
29	Crystallization and preliminary X-ray diffraction analysis of a rat biliverdin reductase. Acta Crystallographica Section D: Biological Crystallography, 2000, 56, 1180-1182.	2.5	8
30	Molecular mechanism of nitric oxide reduction catalyzed by fungal nitric oxide reductase. International Congress Series, 2002, 1233, 59-62.	0.2	2