

Nina Hakulinen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9322442/publications.pdf>

Version: 2024-02-01

42
papers

1,143
citations

430754

18
h-index

377752

34
g-index

42
all docs

42
docs citations

42
times ranked

1572
citing authors

#	ARTICLE	IF	CITATIONS
1	Three-dimensional structures of thermophilic beta-1,4-xylanases from <i>Chaetomium thermophilum</i> and <i>Nonomuraea flexuosa</i> . Comparison of twelve xylanases in relation to their thermal stability. <i>FEBS Journal</i> , 2003, 270, 1399-1412.	0.2	188
2	Crystal structure of a laccase from <i>Melanocarpus albomyces</i> with an intact trinuclear copper site. <i>Nature Structural Biology</i> , 2002, 9, 601-5.	9.7	151
3	Essential role of the C-terminus in <i>Melanocarpus albomyces</i> laccase for enzyme production, catalytic properties and structure. <i>FEBS Journal</i> , 2009, 276, 6285-6300.	2.2	73
4	Crystal structure of an ascomycete fungal laccase from <i>Thielavia farenaria</i> – common structural features of ascomycete laccases. <i>FEBS Journal</i> , 2011, 278, 2283-2295.	2.2	71
5	Three-Dimensional Structure of the Catalytic Core of Acetylxylan Esterase from <i>Trichoderma reesei</i> : Insights into the Deacetylation Mechanism. <i>Journal of Structural Biology</i> , 2000, 132, 180-190.	1.3	66
6	Structural analysis, enzymatic characterization, and catalytic mechanisms of β -galactosidase from <i>Bacillus circulans</i> sp. <i>alkalophilus</i> . <i>FEBS Journal</i> , 2012, 279, 1788-1798.	2.2	59
7	A crystallographic and spectroscopic study on the effect of X-ray radiation on the crystal structure of <i>Melanocarpus albomyces</i> laccase. <i>Biochemical and Biophysical Research Communications</i> , 2006, 350, 929-934.	1.0	49
8	Crystal structures of <i>Trichoderma reesei</i> β -galactosidase reveal conformational changes in the active site. <i>Journal of Structural Biology</i> , 2011, 174, 156-163.	1.3	47
9	The Crystal Structure of β -Glucosidase from <i>Bacillus circulans</i> sp. <i>alkalophilus</i> : Ability to Form Long Polymeric Assemblies. <i>Journal of Structural Biology</i> , 2000, 129, 69-79.	1.3	43
10	The crystal structure of an extracellular catechol oxidase from the ascomycete fungus <i>Aspergillus oryzae</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2013, 18, 917-929.	1.1	40
11	Insights into the roles of non-catalytic residues in the active site of a GH10 xylanase with activity on cellulose. <i>Journal of Biological Chemistry</i> , 2017, 292, 19315-19327.	1.6	35
12	Determination of thioxylo-oligosaccharide binding to family 11 xylanases using electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry and X-ray crystallography. <i>FEBS Journal</i> , 2005, 272, 2317-2333.	2.2	28
13	Characterization and mutagenesis of two novel iron-sulphur cluster pentonate dehydratases. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 7549-7563.	1.7	27
14	The Crystal Structure of a Bacterial l-Arabinonate Dehydratase Contains a [2Fe-2S] Cluster. <i>ACS Chemical Biology</i> , 2017, 12, 1919-1927.	1.6	25
15	The Contribution of Polystyrene Nanospheres towards the Crystallization of Proteins. <i>PLoS ONE</i> , 2009, 4, e4198.	1.1	24
16	Improving the catalytic performance of Proteinase K from <i>Parabryodonta album</i> for use in feather degradation. <i>International Journal of Biological Macromolecules</i> , 2020, 154, 1586-1595.	3.6	24
17	The crystal structure of D-xylonate dehydratase reveals functional features of enzymes from the lly/ED dehydratase family. <i>Scientific Reports</i> , 2018, 8, 865.	1.6	21
18	Substrate specificity of 2-deoxy-D-ribose 5-phosphate aldolase (DERA) assessed by different protein engineering and machine learning methods. <i>Applied Microbiology and Biotechnology</i> , 2020, 104, 10515-10529.	1.7	21

#	ARTICLE	IF	CITATIONS
19	Probing the Dioxygen Route in <i>Melanocarpus albomyces</i> Laccase with Pressurized Xenon Gas. <i>Biochemistry</i> , 2011, 50, 4396-4398.	1.2	19
20	Crystallization and preliminary X-ray characterization of <i>Trichoderma reesei</i> hydrophobin HFBII. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004, 60, 163-165.	2.5	16
21	Patulin Detoxification by Recombinant Manganese Peroxidase from <i>Moniliophthora roreri</i> Expressed by <i>Pichia pastoris</i> . <i>Toxins</i> , 2022, 14, 440.	1.5	13
22	Structure and function of <i>Caulobacter crescentus</i> aldose oxidoreductase. <i>Biochemical Journal</i> , 2015, 472, 297-307.	1.7	10
23	Polysaccharide utilization loci-driven enzyme discovery reveals BD-FAE: a bifunctional feruloyl and acetyl xylan esterase active on complex natural xylans. <i>Biotechnology for Biofuels</i> , 2021, 14, 127.	6.2	10
24	Structure and Function of a Decarboxylating <i>Agrobacterium tumefaciens</i> Keto-deoxy-D-galactarate Dehydratase. <i>Biochemistry</i> , 2014, 53, 8052-8060.	1.2	9
25	Unraveling Substrate Specificity and Catalytic Promiscuity of <i>Aspergillus oryzae</i> Catechol Oxidase. <i>ChemBioChem</i> , 2018, 19, 2348-2352.	1.3	9
26	A new crystal form of <i>Aspergillus oryzae</i> catechol oxidase and evaluation of copper site structures in coupled binuclear copper enzymes. <i>PLoS ONE</i> , 2018, 13, e0196691.	1.1	8
27	Cysteine Engineering of an Endo-polygalacturonase from <i>Talaromyces leycettanus</i> JCM 12802 to Improve Its Thermostability. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 6351-6359.	2.4	7
28	Current state of and need for enzyme engineering of 2-deoxy-D-ribose 5-phosphate aldolases and its impact. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 6215-6228.	1.7	7
29	Structural Insights into the Mechanisms Underlying the Kinetic Stability of GH28 Endo-Polygalacturonase. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 815-823.	2.4	7
30	Boosting enzymatic degradation of cellulose using a fungal expansin: Structural insight into the pretreatment mechanism. <i>Bioresource Technology</i> , 2022, 358, 127434.	4.8	6
31	Crystallization and preliminary X-ray analysis of <i>Aspergillus oryzae</i> catechol oxidase. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2011, 67, 672-674.	0.7	5
32	Crystallization and preliminary X-ray diffraction studies of the catalytic core of acetyl xylan esterase from <i>Trichoderma reesei</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1998, 54, 430-432.	2.5	4
33	Purification, crystallization and preliminary X-ray diffraction analysis of a novel keto-deoxy-D-galactarate (KDG) dehydratase from <i>Agrobacterium tumefaciens</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014, 70, 49-52.	0.4	4
34	Xylonolactonase from <i>Caulobacter crescentus</i> Is a Mononuclear Nonheme Iron Hydrolase. <i>Biochemistry</i> , 2021, 60, 3046-3049.	1.2	4
35	Three-dimensional structure of xylonolactonase from <i>Caulobacter crescentus</i> : A mononuclear iron enzyme of the Î²-bladed Î²-propeller hydrolase family. <i>Protein Science</i> , 2022, 31, 371-383.	3.1	4
36	Crystallization and preliminary diffraction analysis of a Î²-galactosidase from <i>Trichoderma reesei</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009, 65, 767-769.	0.7	3

#	ARTICLE	IF	CITATIONS
37	Crystallization and X-ray diffraction analysis of an <i>L</i> -arabinonate dehydratase from <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> and a <i>D</i> -xylonate dehydratase from <i>Caulobacter crescentus</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2016, 72, 604-608.	0.4	3
38	A Novel Laccase from the Ascomycete <i>Melanocarpus albomyces</i> . <i>ACS Symposium Series</i> , 2003, , 315-331.	0.5	1
39	Preliminary X-ray analysis of twinned crystals of sarcosine dimethylglycine methyltransferase from <i>Halorhodospira halochoris</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009, 65, 805-808.	0.7	1
40	Linear birefringence magnitude of artificial self-assembled DNA crystals. <i>Optical Materials Express</i> , 2011, 1, 936.	1.6	1
41	Crystallization of 3-hexulose-6-phosphate synthase. <i>European Journal of Chemistry</i> , 2021, 12, 299-303.	0.3	0
42	Crystal structure of galactarolactone cycloisomerase from <i>Agrobacterium tumefaciens</i> . <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2013, 69, s306-s307.	0.3	0