

Annette Eva Langkilde

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8362351/publications.pdf>

Version: 2024-02-01

25
papers

773
citations

623188

14
h-index

642321

23
g-index

27
all docs

27
docs citations

27
times ranked

1409
citing authors

#	ARTICLE	IF	CITATIONS
1	Insight into calcium-binding motifs of intrinsically disordered proteins. <i>Biophysical Journal</i> , 2022, 121, 300a.	0.2	0
2	Recommendations for addressing the translational gap between experimental and clinical research on amyloid diseases. <i>Journal of Translational Medicine</i> , 2022, 20, 213.	1.8	0
3	Structural insights into protein folding, stability and activity using <i>in vivo</i> perdeuteration of hen egg-white lysozyme. <i>IUCr</i> , 2021, 8, 372-386.	1.0	4
4	Refinement of $\hat{\mu}$ -Synuclein Ensembles Against SAXS Data: Comparison of Force Fields and Methods. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 654333.	1.6	51
5	$\hat{\mu}$ -Synuclein Responses in the Laterodorsal Tegmentum, the Pedunculo-pontine Tegmentum, and the Substantia Nigra: Implications for Early Appearance of Sleep Disorders in Parkinson's Disease. <i>Journal of Parkinson's Disease</i> , 2021, 11, 1-18.	1.5	3
6	Insight into Calcium-Binding Motifs of Intrinsically Disordered Proteins. <i>Biomolecules</i> , 2021, 11, 1173.	1.8	16
7	Structure and thermodynamics of transient protein-protein complexes by chemometric decomposition of SAXS datasets. <i>Structure</i> , 2021, 29, 1074-1090.e4.	1.6	7
8	The impact of folding modes and deuteration on the atomic resolution structure of hen egg-white lysozyme. <i>Acta Crystallographica Section D: Structural Biology</i> , 2021, 77, 1579-1590.	1.1	3
9	Developing Inhibitors of the p47phox-p22phox Protein-Protein Interaction by Fragment-Based Drug Discovery. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 1156-1177.	2.9	25
10	The Non-Fibrillating N-Terminal of $\hat{\mu}$ -Synuclein Binds and Co-Fibrillates with Heparin. <i>Biomolecules</i> , 2020, 10, 1192.	1.8	6
11	The structural basis of fungal glucuronoyl esterase activity on natural substrates. <i>Nature Communications</i> , 2020, 11, 1026.	5.8	16
12	Size-Selective Phagocytic Clearance of Fibrillar $\hat{\mu}$ -Synuclein through Conformational Activation of Complement Receptor 4. <i>Journal of Immunology</i> , 2020, 204, 1345-1361.	0.4	23
13	Avidity within the N-terminal anchor drives $\hat{\mu}$ -synuclein membrane interaction and insertion. <i>FASEB Journal</i> , 2020, 34, 7462-7482.	0.2	28
14	Distinct $\hat{\mu}$ -Synuclein:Lipid Co-Structure Complexes Affect Amyloid Nucleation through Fibril Mimetic Behavior. <i>Biochemistry</i> , 2019, 58, 5052-5065.	1.2	12
15	Femtosecond X-ray coherent diffraction of aligned amyloid fibrils on low background graphene. <i>Nature Communications</i> , 2018, 9, 1836.	5.8	34
16	Noninvasive Structural Analysis of Intermediate Species During Fibrillation: An Application of Small-Angle X-Ray Scattering. <i>Methods in Molecular Biology</i> , 2018, 1779, 209-239.	0.4	5
17	Monoclonal Antibodies Follow Distinct Aggregation Pathways During Production-Relevant Acidic Incubation and Neutralization. <i>Pharmaceutical Research</i> , 2016, 33, 716-728.	1.7	45
18	In-depth analysis of subclass-specific conformational preferences of IgG antibodies. <i>IUCr</i> , 2015, 2, 9-18.	1.0	59

#	ARTICLE	IF	CITATIONS
19	The architecture of amyloid-like peptide fibrils revealed by X-ray scattering, diffraction and electron microscopy. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 882-895.	2.5	50
20	Small-Angle X-ray Scattering Screening Complements Conventional Biophysical Analysis: Comparative Structural and Biophysical Analysis of Monoclonal Antibodies IgG1, IgG2, and IgG4. <i>Journal of Pharmaceutical Sciences</i> , 2014, 103, 1701-1710.	1.6	54
21	Protein/Lipid Coaggregates are Formed During $\hat{1}\pm$ -Synuclein-Induced Disruption of Lipid Bilayers. <i>Biomacromolecules</i> , 2014, 15, 3643-3654.	2.6	51
22	Wildtype and A30P Mutant Alpha-Synuclein Form Different Fibril Structures. <i>PLoS ONE</i> , 2013, 8, e67713.	1.1	48
23	Structural and Functional Insight into How the Plasmodium falciparum VAR2CSA Protein Mediates Binding to Chondroitin Sulfate A in Placental Malaria. <i>Journal of Biological Chemistry</i> , 2012, 287, 23332-23345.	1.6	154
24	Structural Characterization of Prefibrillar Intermediates and Amyloid Fibrils by Small-Angle X-Ray Scattering. <i>Methods in Molecular Biology</i> , 2012, 849, 137-155.	0.4	10
25	Methods for structural characterization of prefibrillar intermediates and amyloid fibrils. <i>FEBS Letters</i> , 2009, 583, 2600-2609.	1.3	63