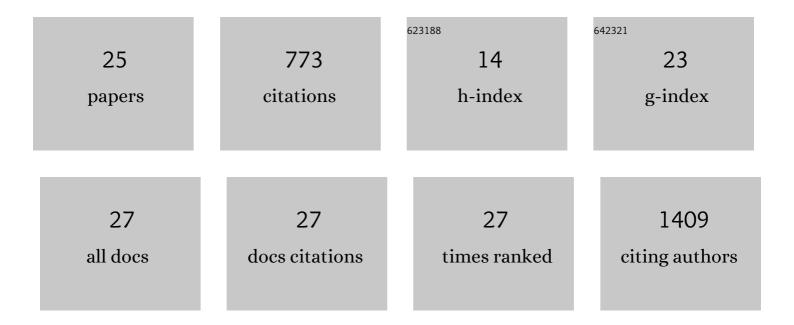
Annette Eva Langkilde

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Structural and Functional Insight into How the Plasmodium falciparum VAR2CSA Protein Mediates Binding to Chondroitin Sulfate A in Placental Malaria. Journal of Biological Chemistry, 2012, 287, 23332-23345. | 1.6 | 154 |
| 2 | Methods for structural characterization of prefibrillar intermediates and amyloid fibrils. FEBS Letters, 2009, 583, 2600-2609. | 1.3 | 63 |
| 3 | In-depth analysis of subclass-specific conformational preferences of IgG antibodies. IUCrJ, 2015, 2, 9-18. | 1.0 | 59 |
| 4 | Small-Angle X-ray Scattering Screening Complements Conventional Biophysical Analysis: Comparative Structural and Biophysical Analysis of Monoclonal Antibodies IgG1, IgG2, and IgG4. Journal of Pharmaceutical Sciences, 2014, 103, 1701-1710. | 1.6 | 54 |
| 5 | Protein/Lipid Coaggregates are Formed During $\hat{I}\pm$ -Synuclein-Induced Disruption of Lipid Bilayers. Biomacromolecules, 2014, 15, 3643-3654. | 2.6 | 51 |
| 6 | Refinement of α-Synuclein Ensembles Against SAXS Data: Comparison of Force Fields and Methods. Frontiers in Molecular Biosciences, 2021, 8, 654333. | 1.6 | 51 |
| 7 | The architecture of amyloid-like peptide fibrils revealed by X-ray scattering, diffraction and electron microscopy. Acta Crystallographica Section D: Biological Crystallography, 2015, 71, 882-895. | 2.5 | 50 |
| 8 | Wildtype and A30P Mutant Alpha-Synuclein Form Different Fibril Structures. PLoS ONE, 2013, 8, e67713. | 1.1 | 48 |
| 9 | Monoclonal Antibodies Follow Distinct Aggregation Pathways During Production-Relevant Acidic Incubation and Neutralization. Pharmaceutical Research, 2016, 33, 716-728. | 1.7 | 45 |
| 10 | Femtosecond X-ray coherent diffraction of aligned amyloid fibrils on low background graphene. Nature Communications, 2018, 9, 1836. | 5.8 | 34 |
| 11 | Avidity within the Nâ€ŧerminal anchor drives αâ€synuclein membrane interaction and insertion. FASEB Journal, 2020, 34, 7462-7482. | 0.2 | 28 |
| 12 | Developing Inhibitors of the p47phox–p22phox Protein–Protein Interaction by Fragment-Based Drug Discovery. Journal of Medicinal Chemistry, 2020, 63, 1156-1177. | 2.9 | 25 |
| 13 | Size-Selective Phagocytic Clearance of Fibrillar α-Synuclein through Conformational Activation of Complement Receptor 4. Journal of Immunology, 2020, 204, 1345-1361. | 0.4 | 23 |
| 14 | The structural basis of fungal glucuronoyl esterase activity on natural substrates. Nature Communications, 2020, 11, 1026. | 5.8 | 16 |
| 15 | Insight into Calcium-Binding Motifs of Intrinsically Disordered Proteins. Biomolecules, 2021, 11, 1173. | 1.8 | 16 |
| 16 | Distinct α-Synuclein:Lipid Co-Structure Complexes Affect Amyloid Nucleation through Fibril Mimetic Behavior. Biochemistry, 2019, 58, 5052-5065. | 1.2 | 12 |
| 17 | Structural Characterization of Prefibrillar Intermediates and Amyloid Fibrils by Small-Angle X-Ray Scattering. Methods in Molecular Biology, 2012, 849, 137-155. | 0.4 | 10 |
| 18 | Structure and thermodynamics of transient protein-protein complexes by chemometric decomposition of SAXS datasets. Structure, 2021, 29, 1074-1090.e4. | 1.6 | 7 |

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | The Non-Fibrillating N-Terminal of α-Synuclein Binds and Co-Fibrillates with Heparin. Biomolecules, 2020, 10, 1192. | 1.8 | 6 |
| 20 | Noninvasive Structural Analysis of Intermediate Species During Fibrillation: An Application of Small-Angle X-Ray Scattering. Methods in Molecular Biology, 2018, 1779, 209-239. | 0.4 | 5 |
| 21 | Structural insights into protein folding, stability and activity using <i>in vivo</i> perdeuteration of hen egg-white lysozyme. IUCrJ, 2021, 8, 372-386. | 1.0 | 4 |
| 22 | α-Synuclein Responses in the Laterodorsal Tegmentum, the Pedunculopontine Tegmentum, and the Substantia Nigra: Implications for Early Appearance of Sleep Disorders in Parkinson's Disease. Journal of Parkinson's Disease, 2021, 11, 1-18. | 1.5 | 3 |
| 23 | The impact of folding modes and deuteration on the atomic resolution structure of hen egg-white lysozyme. Acta Crystallographica Section D: Structural Biology, 2021, 77, 1579-1590. | 1.1 | 3 |
| 24 | Insight into calcium-binding motifs of intrinsically disordered proteins. Biophysical Journal, 2022, 121, 300a. | 0.2 | 0 |
| 25 | Recommendations for addressing the translational gap between experimental and clinical research on amyloid diseases. Journal of Translational Medicine, 2022, 20, 213. | 1.8 | Ο |