## Brent L Nannenga

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/889741/publications.pdf

Version: 2024-02-01

41 papers 2,675 citations

430874 18 h-index 302126 39 g-index

48 all docs

48 docs citations

48 times ranked

2953 citing authors

| #  | Article  | IF                 | CITATIONS           |
|----|--|--------------------|---------------------|
| 1  | Protein–Nanoparticle Complex Structure Determination by Cryo-Electron Microscopy. ACS Applied Bio Materials, 2022, 5, 4696-4700.   | 4.6                | 3                   |
| 2  | Efficient Free Triplet Generation Follows Singlet Fission in Diketopyrrolopyrrole Polymorphs with Goldilocks Coupling. Journal of Physical Chemistry C, 2021, 125, 12207-12213.                                      | 3.1                | 14                  |
| 3  | MicroED structure of the human adenosine receptor determined from a single nanocrystal in LCP.<br>Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .                      | 7.1                | 36                  |
| 4  | Structural insights into the function of the catalytically active human Taspase 1. Structure, 2021, 29, 873-885.e5.  | 3.3                | 4                   |
| 5  | Structureâ€guided identification of a peptide for bioâ€enabled gold nanoparticle synthesis.<br>Biotechnology and Bioengineering, 2021, 118, 4867-4873.   | 3.3                | 1                   |
| 6  | Recent Developments Toward Integrated Metabolomics Technologies (UHPLC-MS-SPE-NMR and) Tj ETQq0 0 0 rg<br>Biosciences, 2021, 8, 720955.  | gBT /Overlo<br>3.5 | ock 10 Tf 50 5<br>9 |
| 7  | MicroED Sample Preparation and Data Collection For Protein Crystals. Methods in Molecular Biology, 2021, 2215, 287-297.  | 0.9                | 7                   |
| 8  | Heterologous expression and purification of the bicarbonate transporter BicA from Synechocystis sp. PCC 6803. Protein Expression and Purification, 2020, 175, 105716.  | 1.3                | 1                   |
| 9  | Structure Determination from Lipidic Cubic Phase Embedded Microcrystals by MicroED. Structure, 2020, 28, 1149-1159.e4.   | 3.3                | 21                  |
| 10 | Beam-sensitive metal-organic framework structure determination by microcrystal electron diffraction. Ultramicroscopy, 2020, 216, 113048.   | 1.9                | 18                  |
| 11 | Crystal structure and orientation of organic semiconductor thin films by microcrystal electron diffraction and grazing-incidence wide-angle X-ray scattering. Chemical Communications, 2020, 56, 4204-4207.          | 4.1                | 27                  |
| 12 | MicroED methodology and development. Structural Dynamics, 2020, 7, 014304.   | 2.3                | 34                  |
| 13 | Rapid Structural Analysis of a Synthetic Non-canonical Amino Acid by Microcrystal Electron Diffraction. Frontiers in Molecular Biosciences, 2020, 7, 609999.   | <b>3.</b> 5        | 6                   |
| 14 | Tetragonal crystal form of the cyanobacterial bicarbonate-transporter regulator SbtB from <i>Synechocystis</i> sp. PCC 6803. Acta Crystallographica Section F, Structural Biology Communications, 2020, 76, 438-443. | 0.8                | 0                   |
| 15 | The complementarity of serial femtosecond crystallography and MicroED for structure determination from microcrystals. Current Opinion in Structural Biology, 2019, 58, 286-293.                                      | 5.7                | 18                  |
| 16 | Proteinâ€facilitated gold nanoparticle formation as indicators of ionizing radiation. Biotechnology and Bioengineering, 2019, 116, 3160-3167.  | 3.3                | 5                   |
| 17 | Electrophoretic exclusion microscale sample preparation for cryo-EM structural determination of proteins. Biomicrofluidics, 2019, 13, 054112.  | 2.4                | 5                   |
| 18 | The cryo-EM method microcrystal electron diffraction (MicroED). Nature Methods, 2019, 16, 369-379.   | 19.0               | 170                 |

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|----|--|------|-----------|
| 19 | Microcrystal electron diffraction methodology and applications. MRS Bulletin, 2019, 44, 956-960.   | 3.5  | 2         |
| 20 | The Evolution and the Advantages of MicroED. Frontiers in Molecular Biosciences, 2018, 5, 114.   | 3.5  | 15        |
| 21 | MicroED: a versatile cryoEM method for structure determination. Emerging Topics in Life Sciences, 2018, 2, 1-8.  | 2.6  | 22        |
| 22 | Atomic structures of fibrillar segments of hIAPP suggest tightly mated $\hat{l}^2$ -sheets are important for cytotoxicity. ELife, 2017, 6, .   | 6.0  | 95        |
| 23 | The collection of MicroED data for macromolecular crystallography. Nature Protocols, 2016, 11, 895-904.  | 12.0 | 117       |
| 24 | MicroED opens a new era for biological structure determination. Current Opinion in Structural Biology, 2016, 40, 128-135.  | 5.7  | 46        |
| 25 | MicroED data collection and processing. Acta Crystallographica Section A: Foundations and Advances, 2015, 71, 353-360.   | 0.1  | 115       |
| 26 | EKylation: Addition of an Alternating-Charge Peptide Stabilizes Proteins. Biomacromolecules, 2015, 16, 3357-3361.  | 5.4  | 51        |
| 27 | Structure of the toxic core of α-synuclein from invisible crystals. Nature, 2015, 525, 486-490.  | 27.8 | 528       |
| 28 | Structure of catalase determined by MicroED. ELife, 2014, 3, e03600.   | 6.0  | 115       |
| 29 | High thermodynamic stability of parametrically designed helical bundles. Science, 2014, 346, 481-485.  | 12.6 | 264       |
| 30 | Protein structure determination by MicroED. Current Opinion in Structural Biology, 2014, 27, 24-31.  | 5.7  | 46        |
| 31 | High-resolution structure determination by continuous-rotation data collection in MicroED. Nature Methods, 2014, 11, 927-930.  | 19.0 | 340       |
| 32 | Overview of Electron Crystallography of Membrane Proteins: Crystallization and Screening Strategies Using Negative Stain Electron Microscopy. Current Protocols in Protein Science, 2013, 72, Unit17.15. | 2.8  | 25        |
| 33 | Three-dimensional electron crystallography of protein microcrystals. ELife, 2013, 2, e01345.   | 6.0  | 340       |
| 34 | Folding Engineering Strategies for Efficient Membrane Protein Production in E. coli. Methods in Molecular Biology, 2012, 899, 187-202.   | 0.9  | 6         |
| 35 | Enhanced expression of membrane proteins in E. coli with a PBAD promoter mutant: synergies with chaperone pathway engineering strategies. Microbial Cell Factories, 2011, 10, 105.                       | 4.0  | 16        |
| 36 | Reprogramming chaperone pathways to improve membrane protein expression in <i>Escherichia coli</i> li>. Protein Science, 2011, 20, 1411-1420.  | 7.6  | 47        |

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 37 | A story of thrift unfolds. Nature Chemical Biology, 2010, 6, 880-881.  | 8.0  | 10        |
| 38 | Enhancing the secretory yields of leech carboxypeptidase inhibitor in Escherichia coli: Influence of trigger factor and signal recognition particle. Protein Expression and Purification, 2010, 74, 122-128. | 1.3  | 13        |
| 39 | Conformational Targeting of Fibrillar Polyglutamine Proteins in Live Cells Escalates Aggregation and Cytotoxicity. PLoS ONE, 2009, 4, e5727.   | 2.5  | 51        |
| 40 | Antiâ€oligomeric single chain variable domain antibody differentially affects huntingtin and αâ€synuclein aggregates. FEBS Letters, 2008, 582, 517-522.  | 2.8  | 12        |
| 41 | MicroED for the study of protein–ligand interactions and the potential for drug discovery. Nature Reviews Chemistry, 0, , .  | 30.2 | 8         |