

# Kwnag-Im Oh

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3977804/publications.pdf>

Version: 2024-02-01

27  
papers

1,021  
citations

471509

17  
h-index

552781

26  
g-index

28  
all docs

28  
docs citations

28  
times ranked

974  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Systematic Approach to Find the Global Minimum of Relaxation Dispersion Data for Protein-Induced Bâ€“Z Transition of DNA. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3517.          | 4.1  | 2         |
| 2  | Non-Canonical Helical Structure of Nucleic Acids Containing Base-Modified Nucleotides. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9552.   | 4.1  | 9         |
| 3  | Conformational exchange of the ZÎ± domain of human RNA editing enzyme ADAR1 studied by NMR spectroscopy. <i>Biochemical and Biophysical Research Communications</i> , 2021, 580, 63-66.                 | 2.1  | 2         |
| 4  | Protein-induced B-Z transition of DNA duplex containing a 2â€“OMe guanosine. <i>Biochemical and Biophysical Research Communications</i> , 2020, 533, 417-423.   | 2.1  | 2         |
| 5  | Molecular heterogeneity in aqueous cosolvent systems. <i>Journal of Chemical Physics</i> , 2020, 152, 190901.   | 3.0  | 17        |
| 6  | Liquidâ€“Liquid Phase Separation Produces Fast H-Bond Dynamics in DMSOâ€“Water Mixtures. <i>Journal of Physical Chemistry Letters</i> , 2020, 11, 1903-1908.  | 4.6  | 28        |
| 7  | Dynamics Studies of DNA with Non-canonical Structure Using NMR Spectroscopy. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2673.   | 4.1  | 12        |
| 8  | Empirical S=O stretch vibrational frequency map. <i>Journal of Chemical Physics</i> , 2019, 151, 234107.  | 3.0  | 16        |
| 9  | Crowding Stabilizes DMSOâ€“Water Hydrogen-Bonding Interactions. <i>Journal of Physical Chemistry B</i> , 2018, 122, 5984-5990.  | 2.6  | 37        |
| 10 | Quantifying Hydrogenâ€“Bond Populations in Dimethyl Sulfoxide/Water Mixtures. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 11375-11379.   | 13.8 | 94        |
| 11 | Quantifying Hydrogenâ€“Bond Populations in Dimethyl Sulfoxide/Water Mixtures. <i>Angewandte Chemie</i> , 2017, 129, 11533-11537.  | 2.0  | 25        |
| 12 | Titelbild: Quantifying Hydrogenâ€“Bond Populations in Dimethyl Sulfoxide/Water Mixtures (Angew.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5  | 2.0  | 0         |
| 13 | How Sensitive is the Amideâ€“I Vibration of the Polypeptide Backbone to Electric Fields?. <i>ChemPhysChem</i> , 2015, 16, 3595-3598.  | 2.1  | 16        |
| 14 | Kinetics of peptide folding in lipid membranes. <i>Biopolymers</i> , 2015, 104, 281-290.  | 2.4  | 5         |
| 15 | Neighboring Residue Effects in Terminally Blocked Dipeptides: Implications for Residual Secondary Structures in Intrinsically Unfolded/Disordered Proteins. <i>Chirality</i> , 2014, 26, 443-452.       | 2.6  | 11        |
| 16 | A comprehensive library of blocked dipeptides reveals intrinsic backbone conformational propensities of unfolded proteins. <i>Proteins: Structure, Function and Bioinformatics</i> , 2012, 80, 977-990. | 2.6  | 30        |
| 17 | Conformational distributions of denatured and unstructured proteins are similar to those of 20Å—20 blocked dipeptides. <i>Journal of Biomolecular NMR</i> , 2012, 53, 25-41.                            | 2.8  | 22        |
| 18 | Circular dichroism eigenspectra of polyproline II and Î±-strand conformers of trialanine in water: Singular value decomposition analysis. <i>Chirality</i> , 2010, 22, E186-201.                        | 2.6  | 39        |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Azido Gauche Effect on the Backbone Conformation of $\hat{\text{I}}^2$ -Azidoalanine Peptides. <i>Journal of Physical Chemistry B</i> , 2010, 114, 13021-13029.   | 2.6 | 27        |
| 20 | Classical and quantum mechanical/molecular mechanical molecular dynamics simulations of alanine dipeptide in water: Comparisons with IR and vibrational circular dichroism spectra. <i>Journal of Chemical Physics</i> , 2008, 128, 105106. | 3.0 | 78        |
| 21 | $\hat{\text{I}}^2$ -Azidoalanine as an IR Probe: Application to Amyloid A $\hat{\text{I}}^2$ (16-22) Aggregation. <i>Journal of Physical Chemistry B</i> , 2008, 112, 10352-10357.  | 2.6 | 108       |
| 22 | Azido-derivatized compounds as IR probes of local electrostatic environment: Theoretical studies. <i>Journal of Chemical Physics</i> , 2008, 129, 174512.   | 3.0 | 72        |
| 23 | Nitrile and thiocyanate IR probes: Quantum chemistry calculation studies and multivariate least-square fitting analysis. <i>Journal of Chemical Physics</i> , 2008, 128, 134506.  | 3.0 | 168       |
| 24 | Nitrile and thiocyanate IR probes: Molecular dynamics simulation studies. <i>Journal of Chemical Physics</i> , 2008, 128, 154504.   | 3.0 | 124       |
| 25 | Dipeptide Structure Determination by Vibrational Circular Dichroism Combined with Quantum Chemistry Calculations. <i>ChemPhysChem</i> , 2007, 8, 2218-2226.   | 2.1 | 19        |
| 26 | Structure of N-Acetylproline Amide in Liquid Water: Experimentally Measured and Numerically Simulated Infrared and Vibrational Circular Dichroism Spectra. <i>Journal of Physical Chemistry B</i> , 2006, 110, 18834-18843.                 | 2.6 | 38        |
| 27 | Site-Specific Hydrogen-Bonding Interaction between N-Acetylproline Amide and Protic Solvent Molecules: Comparisons of IR and VCD Measurements with MD Simulations. <i>Journal of Physical Chemistry A</i> , 2006, 110, 13355-13365.         | 2.5 | 20        |