

# Sik Lok Lam

## List of Publications by Year in descending order

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48  
papers

644  
citations

567281

15  
h-index

642732

23  
g-index

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48  
docs citations

48  
times ranked

572  
citing authors

#	ARTICLE	IF	CITATIONS
1	NMR identification of left-handed polyproline type II helices. <i>Biopolymers</i> , 2003, 69, 270-281.	2.4	62
2	Effect of 1-methyladenine on double-helical DNA structures. <i>FEBS Letters</i> , 2008, 582, 1629-1633.	2.8	45
3	Use of chemical shifts for structural studies of nucleic acids. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2010, 56, 289-310.	7.5	45
4	Structural roles of CTG repeats in slippage expansion during DNA replication. <i>Nucleic Acids Research</i> , 2005, 33, 1604-1617.	14.5	41
5	Minidumbbell: A New Form of Native DNA Structure. <i>Journal of the American Chemical Society</i> , 2016, 138, 12534-12540.	13.7	28
6	The origin of genetic instability in CCTG repeats. <i>Nucleic Acids Research</i> , 2011, 39, 6260-6268.	14.5	26
7	Effect of 1-methyladenine on thermodynamic stabilities of double-helical DNA structures. <i>FEBS Letters</i> , 2009, 583, 1548-1553.	2.8	23
8	DSHIFT: a web server for predicting DNA chemical shifts. <i>Nucleic Acids Research</i> , 2007, 35, W713-W717.	14.5	22
9	Sequence-specific local structural variations in solution structures of d(CGXX <sup>2</sup> CG) <sub>2</sub> and d(CAXX <sup>2</sup> TC) <sub>2</sub> self-complementary deoxyribonucleic acids. <i>Journal of Molecular Biology</i> , 1997, 266, 745-760.	4.2	19
10	Unusual structures of TTTA repeats in <i>icaC</i> gene of <i>Staphylococcus aureus</i> . <i>FEBS Letters</i> , 2015, 589, 1296-1300.	2.8	19
11	Preferential base pairing modes of T-A mismatches. <i>FEBS Letters</i> , 2011, 585, 3953-3958.	2.8	17
12	New insights into the genetic instability in CCTG repeats. <i>FEBS Letters</i> , 2015, 589, 3058-3063.	2.8	17
13	CAG RNAs induce DNA damage and apoptosis by silencing <i>NUDT16</i> expression in polyglutamine degeneration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	17
14	Random coil proton chemical shifts of deoxyribonucleic acids. <i>Journal of Biomolecular NMR</i> , 2002, 24, 329-337.	2.8	16
15	Crosslinking reactions of 4-amino-6-oxo-2-vinylpyrimidine with guanine derivatives and structural analysis of the adducts. <i>Nucleic Acids Research</i> , 2015, 43, 7717-7730.	14.5	16
16	Random coil phosphorus chemical shift of deoxyribonucleic acids. <i>Journal of Magnetic Resonance</i> , 2004, 171, 193-200.	2.1	15
17	NMR investigation of DNA primer-template models: Structural insights into dislocation mutagenesis in DNA replication. <i>FEBS Letters</i> , 2006, 580, 6496-6500.	2.8	15
18	Nuclear Magnetic Resonance Investigation of Primer-Template Models: Formation of a Pyrimidine Bulge upon Misincorporation. <i>Biochemistry</i> , 2008, 47, 4469-4476.	2.5	15

#	ARTICLE	IF	CITATIONS
19	NMR Investigation of Primer-Template Models: Structural Effect of Sequence Downstream of a Thymine Template on Mutagenesis in DNA Replication. <i>Biochemistry</i> , 2007, 46, 9292-9300.	2.5	12
20	Low Temperature Solution Structures and Base Pair Stacking of Double Helical d(CGTACG) <sub>2</sub> . <i>Journal of Biomolecular Structure and Dynamics</i> , 2002, 19, 907-917.	3.5	11
21	Minidumbbell structures formed by ATTCT pentanucleotide repeats in spinocerebellar ataxia type 10. <i>Nucleic Acids Research</i> , 2020, 48, 7557-7568.	14.5	11
22	NMR Investigation of DNA Primer-Template Models: Guanine Templates Are Less Prone to Strand Slippage upon Misincorporation. <i>Biochemistry</i> , 2009, 48, 11478-11486.	2.5	10
23	The competing mini-dumbbell mechanism: new insights into CCTG repeat expansion. <i>Signal Transduction and Targeted Therapy</i> , 2016, 1, 16028.	17.1	10
24	Unusual structures of CCTG repeats and their participation in repeat expansion. <i>Biomolecular Concepts</i> , 2016, 7, 331-340.	2.2	10
25	Achilles Tendon Xanthomas: Fat-Water Separation at Baseline and after Treatment. <i>Radiology</i> , 2017, 285, 876-884.	7.3	10
26	An Extraordinarily Stable DNA Minidumbbell. <i>Journal of Physical Chemistry Letters</i> , 2017, 8, 3478-3481.	4.6	9
27	Random coil carbon chemical shifts of deoxyribonucleic acids. <i>Journal of Magnetic Resonance</i> , 2004, 166, 11-18.	2.1	8
28	Proton chemical shift prediction of A•A mismatches in B-DNA duplexes. <i>Journal of Magnetic Resonance</i> , 2007, 187, 105-111.	2.1	8
29	Sequence Context Effect on Strand Slippage in Natural DNA Primer-Template. <i>Journal of Physical Chemistry B</i> , 2012, 116, 1999-2007.	2.6	8
30	Rational design of a reversible Mg <sup>2+</sup> /EDTA-controlled molecular switch based on a DNA minidumbbell. <i>Chemical Communications</i> , 2020, 56, 10127-10130.	4.1	8
31	Effect of hyperoxidized guanine on DNA primer-template structures: Spiroiminodihydroantoin leads to strand slippage. <i>FEBS Letters</i> , 2008, 582, 4169-4175.	2.8	7
32	Effects of Adenine Methylation on the Structure and Thermodynamic Stability of a DNA Minidumbbell. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3633.	4.1	7
33	NMR proton chemical shift prediction of T•T mismatches in B-DNA duplexes. <i>Journal of Magnetic Resonance</i> , 2013, 234, 184-189.	2.1	6
34	Formation of a DNA Mini-Dumbbell with a Quasi-Type II Loop. <i>Journal of Physical Chemistry B</i> , 2017, 121, 2554-2560.	2.6	6
35	Effect of an Abasic Site on Strand Slippage in DNA Primer-Template. <i>Journal of Physical Chemistry B</i> , 2012, 116, 14781-14787.	2.6	5
36	NMR proton chemical shift prediction of C•C mismatches in B-DNA. <i>Journal of Magnetic Resonance</i> , 2015, 252, 87-93.	2.1	5

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37	Unprecedented hydrophobic stabilizations from a reverse wobble T <sup>+</sup> A mispair in DNA minidumbbell. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 1946-1953.	3.5	5
38	High-Resolution Structures of DNA Minidumbbells Comprising Type II Tetraloops with a Purine Minor Groove Residue. <i>Journal of Physical Chemistry B</i> , 2020, 124, 5131-5138.	2.6	4
39	NMR solution structures of d(GGCCTG) <sub>n</sub> repeats associated with spinocerebellar ataxia type 36. <i>International Journal of Biological Macromolecules</i> , 2022, 201, 607-615.	7.5	4
40	Determination of Backbone Torsion Angle $\gamma$ of DNA Duplexes in Solution from <sup>3</sup> J <sub>C4</sub> and <sup>3</sup> J <sub>H3</sub> , Using Heteronuclear Single-Quantum-Coherence Spectroscopy. <i>Journal of Magnetic Resonance Series B</i> , 1996, 113, 59-64.	1.6	3
41	The J-coupling Restrained Molecular Mechanics (JrMM) Protocol - An Efficient Alternative for Deriving DNA Endocyclic Torsion Angle Constraints Part II: Experimental Application of the JrMM Protocol. <i>Journal of Biomolecular Structure and Dynamics</i> , 1996, 13, 815-825.	3.5	3
42	Cobalt(II) amido complexes derived from a monodentate arylamido ligand featuring a highly electron-withdrawing C <sub>6</sub> F <sub>5</sub> substituent. <i>Dalton Transactions</i> , 2015, 44, 17950-17959.	3.3	3
43	Sequence Effect on the Formation of DNA Minidumbbells. <i>Journal of Physical Chemistry B</i> , 2017, 121, 10338-10343.	2.6	3
44	The J-coupling Restrained Molecular Mechanics (JrMM) Protocol - An Efficient Alternative for Deriving DNA Endocyclic Torsion Angle Constraints Part I: Correlation of Endocyclic Torsion Angles and Vicinal Torsion Angle $\gamma$ . <i>Journal of Biomolecular Structure and Dynamics</i> , 1996, 13, 803-814.	3.5	2
45	5-Methylcytosine Substantially Enhances the Thermal Stability of DNA Minidumbbells. <i>Chemistry - A European Journal</i> , 2021, 27, 6740-6747.	3.3	2
46	Conformational flexibility in the RNA stem-loop structures formed by CAG repeats. <i>FEBS Letters</i> , 2017, 591, 1752-1760.	2.8	2
47	A pH and Mg <sup>2+</sup> -Responsive Molecular Switch Based on a Stable DNA Minidumbbell Bearing 5' and 3'-Overhangs. <i>ACS Omega</i> , 2021, 6, 28263-28269.	3.5	2
48	A purine and a backbone discontinuous site alter the structure and thermal stability of DNA minidumbbells containing two pentaloops. <i>FEBS Letters</i> , 2022, 596, 826-840.	2.8	2