

# Clay Bracken

## List of Publications by Year in descending order

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28  
papers

2,164  
citations

394421

19  
h-index

580821

25  
g-index

28  
all docs

28  
docs citations

28  
times ranked

3527  
citing authors

#	ARTICLE	IF	CITATIONS
1	A method for efficient isotopic labeling of recombinant proteins. <i>Journal of Biomolecular NMR</i> , 2001, 20, 71-75.	2.8	676
2	Temperature dependence of intramolecular dynamics of the basic leucine zipper of GCN4: implications for the entropy of association with DNA 1 Edited by P. E. Wright. <i>Journal of Molecular Biology</i> , 1999, 285, 2133-2146.	4.2	212
3	Val66Met polymorphism of BDNF alters prodomain structure to induce neuronal growth cone retraction. <i>Nature Communications</i> , 2013, 4, 2490.	12.8	185
4	Structural Architecture of the CARMA1/Bcl10/MALT1 Signalosome: Nucleation-Induced Filamentous Assembly. <i>Molecular Cell</i> , 2013, 51, 766-779.	9.7	163
5	An Effective Method for the Discrimination of Motional Anisotropy and Chemical Exchange. <i>Journal of the American Chemical Society</i> , 2002, 124, 1852-1853.	13.7	162
6	Combining prediction, computation and experiment for the characterization of protein disorder. <i>Current Opinion in Structural Biology</i> , 2004, 14, 570-576.	5.7	125
7	Synthesis and Nuclear Magnetic Resonance Structure Determination of an $\alpha$ -Helical, Bicyclic, Lactam-Bridged Hexapeptide. <i>Journal of the American Chemical Society</i> , 1994, 116, 6431-6432.	13.7	96
8	Molecular Motions and Protein Folding: Characterization of the Backbone Dynamics and Folding Equilibrium of $\beta$ -2D Using $^{13}\text{C}$ NMR Spin Relaxation. <i>Journal of the American Chemical Society</i> , 2000, 122, 11610-11619.	13.7	73
9	The BDNF Val66Met Prodomain Disassembles Dendritic Spines Altering Fear Extinction Circuitry and Behavior. <i>Neuron</i> , 2018, 99, 163-178.e6.	8.1	53
10	The volatile anesthetic isoflurane perturbs conformational activation of integrin LFA $\beta$ 1 by binding to the allosteric regulatory cavity. <i>FASEB Journal</i> , 2008, 22, 4109-4116.	0.5	50
11	NMR spin relaxation methods for characterization of disorder and folding in proteins. <i>Journal of Molecular Graphics and Modelling</i> , 2001, 19, 3-12.	2.4	47
12	Structural Model of the Extracellular Assembly of the TCR-CD3 Complex. <i>Cell Reports</i> , 2016, 14, 2833-2845.	6.4	46
13	(H)N(COCA)NH and HN(COCA)NH experiments for $^1\text{H}$ - $^{15}\text{N}$ backbone assignments in $^{13}\text{C}/^{15}\text{N}$ -labeled proteins. <i>Journal of Biomolecular NMR</i> , 1997, 9, 94-100.	2.8	41
14	Characterization of millisecond time-scale dynamics in the molten globule state of $\beta$ -lactalbumin by NMR. <i>Journal of Molecular Biology</i> , 1999, 294, 551-560.	4.2	36
15	Buried Polar Interactions and Conformational Stability in the Simian Immunodeficiency Virus (SIV) gp41 Core. <i>Biochemistry</i> , 2000, 39, 676-685.	2.5	34
16	Assembling Ligands In Situ Using Bioorthogonal Boronate Ester Synthesis. <i>Chemistry and Biology</i> , 2010, 17, 1171-1176.	6.0	34
17	Use of paramagnetic $^{19}\text{F}$ NMR to monitor domain movement in a glutamate transporter homolog. <i>Nature Chemical Biology</i> , 2020, 16, 1006-1012.	8.0	31
18	The DNA-binding domain in the <i>Bacillus subtilis</i> transition-state regulator AbrB employs significant motion for promiscuous DNA recognition. <i>Journal of Molecular Biology</i> , 2001, 305, 429-439.	4.2	24

#	ARTICLE	IF	CITATIONS
19	Helix formation and the unfolded state of a 52-residue helical protein. <i>Protein Science</i> , 2004, 13, 177-189.	7.6	19
20	pH Dependence of Amide Chemical Shifts in Natively Disordered Polypeptides Detects Medium-Range Interactions with Ionizable Residues. <i>Biophysical Journal</i> , 2005, 89, 3293-3302.	0.5	18
21	Post-translational modifications within tau paired helical filament nucleating motifs perturb microtubule interactions and oligomer formation. <i>Journal of Biological Chemistry</i> , 2022, 298, 101442.	3.4	16
22	Zinc induced structural changes in the intrinsically disordered BDNF Met prodomain confer synaptic elimination. <i>Metallomics</i> , 2020, 12, 1208-1219.	2.4	6
23	Spin Relaxation Methods for Characterizing Picosecond-Nanosecond and Microsecond-Millisecond Motions in Proteins. , 1999, , 171-190.		6
24	Determination of amide exchange rates by measurement of 2D NMR line-broadening. <i>Journal of the American Chemical Society</i> , 1993, 115, 6346-6348.	13.7	4
25	CO <sub>2</sub> H(N)CACB experiments for assigning backbone resonances in <sup>13</sup> C/ <sup>15</sup> N-labeled proteins. <i>Journal of Biomolecular NMR</i> , 1998, 11, 451-456.	2.8	4
26	NMR backbone resonance assignments of the prodomain variants of BDNF in the urea denatured state. <i>Biomolecular NMR Assignments</i> , 2018, 12, 43-45.	0.8	3
27	Elasticity measurements predict no membrane toxicity for the cardiolipin-targeted mitochondrial therapeutic SSâ€³1 (1054.3). <i>FASEB Journal</i> , 2014, 28, 1054.3.	0.5	0
28	Targeting the [CL/cyt c] complex to protect electron transport and prevent ROS formation prevents mitochondrial IR injury without ROS scavenging (LB169). <i>FASEB Journal</i> , 2014, 28, LB169.	0.5	0