David G Norman

List of Publications by Year
in descending order

11	547	933447 10	1281871 11
papers	citations	h-index	g-index

13	13	13	573
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	EPR distance measurements in deuterated proteins. Journal of Magnetic Resonance, 2010, 207, 164-167.	2.1	134
2	Activity, disulphate mapping and structural modelling of the fifth domain of human Î²2 -glycoprotein I. FEBS Letters, 1992, 313, 193-197.	2.8	89
3	Structure of the chromatin remodelling enzyme Chd1 bound to a ubiquitinylated nucleosome. ELife, 2018, 7, .	6.0	72
4	Subâ€Micromolar Pulse Dipolar EPR Spectroscopy Reveals Increasing Cu^{II}â€labelling of Doubleâ€Histidine Motifs with Lower Temperature. Angewandte Chemie - International Edition, 2019, 58, 11681-11685.	13.8	61
5	Structural reorganization of the chromatin remodeling enzyme Chd1 upon engagement with nucleosomes. ELife, 2017, 6, .	6.0	51
6	A Gadolinium Spin Label with Both a Narrow Central Transition and Short Tether for Use in Double Electron Electron Resonance Distance Measurements. Inorganic Chemistry, 2019, 58, 3015-3025.	4.0	39
7	The histone chaperone Vps75 forms multiple oligomeric assemblies capable of mediating exchange between histone H3â€“H4 tetramers and Asf1â€“H3â€“H4 complexes. Nucleic Acids Research, 2016, 44, 6157-6172.	14.5	30
8	Conserved structure and domain organization among bacterial Slc26 transporters. Biochemical Journal, 2014, 463, 297-307.	3.7	25
9	Subâ€Micromolar Pulse Dipolar EPR Spectroscopy Reveals Increasing Cu^{II}â€labelling of Doubleâ€Histidine Motifs with Lower Temperature. Angewandte Chemie, 2019, 131, 11807-11811.	2.0	21
10	A general model to optimise Cu^{II}labelling efficiency of double-histidine motifs for pulse dipolar EPR applications. Physical Chemistry Chemical Physics, 2021, 23, 3810-3819.	2.8	21
11	Analysis of the Intrinsically Disordered N-Terminus of the DNA Junction-Resolving Enzyme T7 Endonuclease I: Identification of Structure Formed upon DNA Binding. Biochemistry, 2016, 55, 4166-4172.	2.5	3