## Chao Wang

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Structural Insights into the Redox-Regulated Dynamic Conformations of Human Protein Disulfide Isomerase. Antioxidants and Redox Signaling, 2013, 19, 36-45.	5.4	183
2	Human Protein-disulfide Isomerase Is a Redox-regulated Chaperone Activated by Oxidation of Domain a′. Journal of Biological Chemistry, 2012, 287, 1139-1149.	3.4	110
3	Plasticity of Human Protein Disulfide Isomerase. Journal of Biological Chemistry, 2010, 285, 26788-26797.	3.4	61
4	Bridging Genomics to Phenomics at Atomic Resolution through Variation Spatial Profiling. Cell Reports, 2018, 24, 2013-2028.e6.	6.4	33
5	Compact Conformations of Human Protein Disulfide Isomerase. PLoS ONE, 2014, 9, e103472.	2.5	26
6	Quantitating the epigenetic transformation contributing to cholesterol homeostasis using Gaussian process. Nature Communications, 2019, 10, 5052.	12.8	18
7	Individualized management of genetic diversity in Niemann-Pick C1 through modulation of the Hsp70 chaperone system. Human Molecular Genetics, 2020, 29, 1-19.	2.9	18
8	Spatial covariance analysis reveals the residue-by-residue thermodynamic contribution of variation to the CFTR fold. Communications Biology, 2022, 5, 356.	4.4	10
9	Measuring the Effect of Histone Deacetylase Inhibitors (HDACi) on the Secretion and Activity of Alpha-1 Antitrypsin. Methods in Molecular Biology, 2017, 1639, 185-193.	0.9	8
10	Leveraging Population Genomics for Individualized Correction of the Hallmarks of Alpha-1 Antitrypsin Deficiency. Chronic Obstructive Pulmonary Diseases (Miami, Fla ), 2020, 7, 224-246.	0.7	4
11	Managing the Adaptive Proteostatic Landscape: Restoring Resilience in Alpha-1 Antitrypsin Deficiency. Respiratory Medicine, 2016. , 53-83.	0.1	1