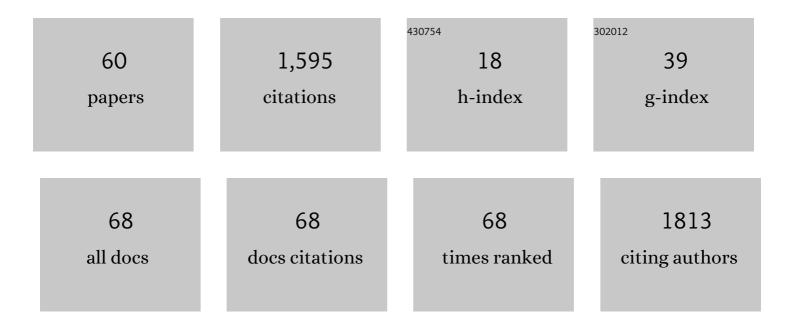
List of Publications by Year in descending order

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| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Designing artificial enzymes by intuition and computation. Nature Chemistry, 2010, 2, 15-24.  | 6.6  | 232       |
| 2  | Design and engineering of an O2 transport protein. Nature, 2009, 458, 305-309.  | 13.7 | 228       |
| 3  | Structures of Nitroreductase in Three States. Journal of Biological Chemistry, 2002, 277, 11513-11520.  | 1.6  | 130       |
| 4  | Steady-state kinetic mechanism, stereospecificity, substrate and inhibitor specificity of Enterobacter cloacae nitroreductase. BBA - Proteins and Proteomics, 1998, 1387, 395-405.  | 2.1  | 116       |
| 5  | Flavin Thermodynamics Explain the Oxygen Insensitivity of Enteric Nitroreductasesâ€. Biochemistry,<br>2002, 41, 14197-14205.  | 1.2  | 78        |
| 6  | The HP-1 maquette: From an apoprotein structure to a structured hemoprotein designed to promote redox-coupled proton exchange. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 5536-5541. | 3.3  | 70        |
| 7  | Intelligent design: the de novo engineering of proteins with specified functions. Dalton Transactions, 2006, , 3045.  | 1.6  | 65        |
| 8  | Mutational and spectroscopic studies of the significance of the active site glutamine to metal ion specificity in superoxide dismutase. Journal of Inorganic Biochemistry, 2000, 80, 247-256.   | 1.5  | 60        |
| 9  | Quantitative Structure–Activity Relationships in Two-Electron Reduction of Nitroaromatic<br>Compounds by Enterobacter cloacae NAD(P)H:Nitroreductase. Archives of Biochemistry and<br>Biophysics, 2001, 385, 170-178.                 | 1.4  | 56        |
| 10 | Hydrophilic to amphiphilic design in redox protein maquettes. Current Opinion in Chemical Biology, 2003, 7, 741-748.  | 2.8  | 48        |
| 11 | Nativelike Structure in Designed Four α-Helix Bundles Driven by Buried Polar Interactions. Journal of<br>the American Chemical Society, 2006, 128, 14450-14451.   | 6.6  | 43        |
| 12 | Computational Design of Thermostabilizing <scp>d</scp> -Amino Acid Substitutions. Journal of the American Chemical Society, 2011, 133, 18750-18759.   | 6.6  | 38        |
| 13 | Probing Charge Transport through Peptide Bonds. Journal of Physical Chemistry Letters, 2018, 9,<br>763-767.   | 2.1  | 38        |
| 14 | <i>De Novo</i> Self-Assembling Collagen Heterotrimers Using Explicit Positive and Negative Design.<br>Biochemistry, 2010, 49, 2307-2316.  | 1.2  | 34        |
| 15 | Two-electron reduction of quinones by Enterobacter cloacae NAD(P)H:nitroreductase: quantitative structure-activity relationships. Archives of Biochemistry and Biophysics, 2002, 403, 249-258.  | 1.4  | 32        |
| 16 | Geometric constraints for porphyrin binding in helical protein binding sites. Proteins: Structure,<br>Function and Bioinformatics, 2009, 74, 400-416.   | 1.5  | 25        |
| 17 | Manipulating Cofactor Binding Thermodynamics in an Artificial Oxygen Transport Protein.<br>Biochemistry, 2011, 50, 10254-10261.   | 1.2  | 21        |
| 18 | <sup>15</sup> N Solid-State NMR as a Probe of Flavin H-Bonding. Journal of Physical Chemistry B, 2011, 115, 7788-7798.  | 1.2  | 20        |

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|----|---|------|-----------|
| 19 | Design principles for chlorophyllâ€binding sites in helical proteins. Proteins: Structure, Function and<br>Bioinformatics, 2011, 79, 463-476.   | 1.5  | 20        |
| 20 | Overexpression, Isotopic Labeling, and Spectral Characterization ofEnterobacter cloacaeNitroreductase. Protein Expression and Purification, 1998, 13, 53-60.  | 0.6  | 18        |
| 21 | Controlling complexity and water penetration in functional <i>de novo</i> protein design.<br>Biochemical Society Transactions, 2008, 36, 1106-1111.   | 1.6  | 18        |
| 22 | Reversible proton coupled electron transfer in a peptide-incorporated naphthoquinoneamino acid.<br>Chemical Communications, 2009, , 168-170.  | 2.2  | 18        |
| 23 | Mechanism-Informed Refinement Reveals Altered Substrate-Binding Mode for Catalytically Competent<br>Nitroreductase. Structure, 2017, 25, 978-987.e4.  | 1.6  | 18        |
| 24 | A three-dimensional printed cell for rapid, low-volume spectroelectrochemistry. Analytical<br>Biochemistry, 2013, 439, 1-3.   | 1.1  | 17        |
| 25 | 15N Solid-State NMR Provides a Sensitive Probe of Oxidized Flavin Reactive Sites. Journal of the<br>American Chemical Society, 2006, 128, 15200-15208.  | 6.6  | 15        |
| 26 | Rational design of a zinc phthalocyanine binding protein. Journal of Structural Biology, 2014, 185,<br>178-185.   | 1.3  | 15        |
| 27 | Order, Disorder, and Temperature-Driven Compaction in a Designed Elastin Protein. Journal of<br>Physical Chemistry B, 2018, 122, 2725-2736.   | 1.2  | 15        |
| 28 | Dynamic Factors Affecting Gaseous Ligand Binding in an Artificial Oxygen Transport Protein.<br>Biochemistry, 2013, 52, 447-455.   | 1.2  | 14        |
| 29 | A flavin analogue with improved solubility in organic solvents. Tetrahedron Letters, 2007, 48, 5517-5520.   | 0.7  | 12        |
| 30 | Hydrogen bond-free flavin redox properties: managing flavins in extreme aprotic solvents. Organic<br>and Biomolecular Chemistry, 2008, 6, 2204.   | 1.5  | 12        |
| 31 | Fast, cheap and out of control — Insights into thermodynamic and informatic constraints on natural<br>protein sequences from de novo protein design. Biochimica Et Biophysica Acta - Bioenergetics, 2016,<br>1857, 485-492.                   | 0.5  | 10        |
| 32 | Designing heterotropically activated allosteric conformational switches using supercharging.<br>Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 5291-5297.  | 3.3  | 10        |
| 33 | Manipulating reduction potentials in an artificial safranin cofactor. Tetrahedron Letters, 2012, 53, 1201-1203.   | 0.7  | 9         |
| 34 | Observation of persistent αâ€helical content and discrete types of backbone disorder during a molten<br>globule to ordered peptide transition via deepâ€UV resonance Raman spectroscopy. Journal of Raman<br>Spectroscopy, 2013, 44, 957-962. | 1.2  | 8         |
| 35 | Thermalization of Fluorescent Protein Exciton–Polaritons at Room Temperature. Advanced Materials,<br>2022, 34, e2109107.  | 11.1 | 7         |
| 36 | An extended scope synthesis of an artificial safranine cofactor. Tetrahedron Letters, 2014, 55, 2487-2491.  | 0.7  | 5         |

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|----|---|-----|-----------|
| 37 | Retro-Nitroreductase, a Putative Evolutionary Precursor toEnterobacter cloacaeStrain 96-3<br>Nitroreductase. Antioxidants and Redox Signaling, 2001, 3, 747-755.                  | 2.5 | 4         |
| 38 | The design features cells use to build their transmembrane proton gradient. Physical Biology, 2017, 14, 013001.   | 0.8 | 4         |
| 39 | Dynamics in natural and designed elastins and their relation to elastic fiber structure and recoil.<br>Biophysical Journal, 2021, 120, 4623-4634.                                 | 0.2 | 4         |
| 40 | Fundamental Limits on Wavelength, Efficiency and Yield of the Charge Separation Triad. PLoS ONE, 2012, 7, e36065.   | 1.1 | 3         |
| 41 | Handheld chem/biosensor using extreme conformational changes in designed binding proteins to enhance surface plasmon resonance (SPR). Proceedings of SPIE, 2016, , .              | 0.8 | 2         |
| 42 | An Artificial Safranine Enzyme which Activates Chemotherapeutic Prodrugs. Biophysical Journal, 2013, 104, 205a.   | 0.2 | 1         |
| 43 | Hemoprotein Design using Minimal Sequence Information. Biophysical Journal, 2013, 104, 661a.  | 0.2 | 1         |
| 44 | Dynamics in Natural and Designed Elastins and their Relation to Elastic Fiber Structure and Recoil.<br>Biophysical Journal, 2020, 118, 536a-537a.                                 | 0.2 | 1         |
| 45 | Persistent α-Helical Content and Local Helical Structural Fluctuations from a Molten Globule to<br>Ordered Peptide Transition. Biophysical Journal, 2012, 102, 444a.              | 0.2 | Ο         |
| 46 | Rational Design of a Zinc Phthalocyanine Binding Protein. Biophysical Journal, 2013, 104, 685a.   | 0.2 | 0         |
| 47 | Photosynthesis in a Single Protein. Biophysical Journal, 2015, 108, 605a.   | 0.2 | 0         |
| 48 | Designed Enzymes and the Driving Forces Behind Interdomain Electron Transfer. Biophysical Journal, 2017, 112, 66a.  | 0.2 | 0         |
| 49 | Optimizing Protein Dynamics in Metalloenzyme Design. Biophysical Journal, 2017, 112, 193a.  | 0.2 | Ο         |
| 50 | Design of Supercharged Proteins to Impart Allosteric Behavior and their Use in Biosensing.<br>Biophysical Journal, 2017, 112, 510a.   | 0.2 | 0         |
| 51 | Engendering Methane Monooxygenase and Hydrogen Peroxide Oxidase Activity into a Designed Dimetal<br>Protein by Increasing Protein Dynamics. Biophysical Journal, 2018, 114, 411a. | 0.2 | Ο         |
| 52 | Designed Enzymes: Creating a more Efficient Nitric Oxide Dioxygenase. Biophysical Journal, 2018, 114,<br>35a.   | 0.2 | 0         |
| 53 | NMR Studies of Secondary Structure and Compaction of Minielastin. Biophysical Journal, 2018, 114, 365a.   | 0.2 | 0         |
| 54 | Supercharging as a General Strategy for Making Proteins into Conformational Switches and their Use<br>in Biosensing. Biophysical Journal, 2018, 114, 588a.                        | 0.2 | 0         |

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|----|---|-----|-----------|
| 55 | A General Method to Design Allosteric Conformational Switches. Biophysical Journal, 2019, 116, 164a.  | 0.2 | Ο         |
| 56 | Engendering Catalytic Activity by Increasing Dynamics in a Designed Enzyme. Biophysical Journal, 2019,<br>116, 68a.                           | 0.2 | 0         |
| 57 | Non-Covalent Coatings on Carbon Nanotubes Mediate Photosensitizer Interactions. ACS Applied<br>Materials & Interfaces, 2021, 13, 51343-51350. | 4.0 | Ο         |
| 58 | Enhanced Transverse Photo-Induced Voltage by Slow Light. , 2015, , .  |     | 0         |
| 59 | Handheld highly selective plasmonic chem/biosensor using engineered binding proteins for extreme conformational changes. , 2017, , .          |     | Ο         |
| 60 | Oxidation-reduction and photophysical properties of isomeric forms of Safranin. PLoS ONE, 2022, 17, e0265105.                                 | 1.1 | 0         |