# Alessandra Corazza

## List of Publications by Year in Descending Order



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| 85 papers | 2,117 citations | 25 h-index | 43 g-index |
| 86 ext. papers | 2,349 ext. citations | 4.6 avg, IF | 4.1 L-index |

| # | Paper | IF | Citations |
|---|---|---|---|
| 85 | Amyloid Formation by Globular Proteins: The Need to Narrow the Gap Between and Mechanisms.. *Frontiers in Molecular Biosciences*, **2022**, 9, 830006 | 5.6 | 2 |
| 84 | Insights on peptide topology in the computational design of protein ligands: the example of lysozyme binding peptides. *Physical Chemistry Chemical Physics*, **2021**, 23, 23158-23172 | 3.6 | 0 |
| 83 | Clinical ApoA-IV amyloid is associated with fibrillogenic signal sequence. *Journal of Pathology*, **2021**, 255, 311-318 | 9.4 | 1 |
| 82 | Comparative study of the stabilities of synthetic and natural transthyretin amyloid fibrils. *Journal of Biological Chemistry*, **2020**, 295, 11379-11387 | 5.4 | 8 |
| 81 | Binding of Monovalent and Bivalent Ligands by Transthyretin Causes Different Short- and Long-Distance Conformational Changes. *Journal of Medicinal Chemistry*, **2019**, 62, 8274-8283 | 8.3 | 17 |
| 80 | Citrate stabilized gold nanoparticles interfere with amyloid fibril formation: D76N and N6 -microglobulin variants. *Nanoscale*, **2018**, 10, 4793-4806 | 7.7 | 25 |
| 79 | Interference of citrate-stabilized gold nanoparticles with -microglobulin oligomeric association. *Chemical Communications*, **2018**, 54, 5422-5425 | 5.8 | 11 |
| 78 | H, C and N backbone resonance assignments of the lactamase BlaP from Bacillus licheniformis 749/C and two mutational variants. *Biomolecular NMR Assignments*, **2018**, 12, 69-77 | 0.7 | |
| 77 | Free Energy, Enthalpy and Entropy from Implicit Solvent End-Point Simulations. *Frontiers in Molecular Biosciences*, **2018**, 5, 11 | 5.6 | 18 |
| 76 | Plasminogen activation triggers transthyretin amyloidogenesis. *Journal of Biological Chemistry*, **2018**, 293, 14192-14199 | 5.4 | 42 |
| 75 | PDB2ENTROPY and PDB2TRENT: Conformational and Translational-Rotational Entropy from Molecular Ensembles. *Journal of Chemical Information and Modeling*, **2018**, 58, 1319-1324 | 6.1 | 12 |
| 74 | Dynamics and Thermodynamics of Transthyretin Association from Molecular Dynamics Simulations. *BioMed Research International*, **2018**, 2018, 7480749 | 3 | 5 |
| 73 | Computational design of cyclic peptides for the customized oriented immobilization of globular proteins. *Physical Chemistry Chemical Physics*, **2017**, 19, 2740-2748 | 3.6 | 12 |
| 72 | Citrate-stabilized gold nanoparticles hinder fibrillogenesis of a pathological variant of microglobulin. *Nanoscale*, **2017**, 9, 3941-3951 | 7.7 | 22 |
| 71 | Automation of peak-tracking analysis of stepwise perturbed NMR spectra. *Journal of Biomolecular NMR*, **2017**, 67, 121-134 | 3 | 3 |
| 70 | A specific nanobody prevents amyloidogenesis of D76N microglobulin in vitro and modifies its tissue distribution in vivo. *Scientific Reports*, **2017**, 7, 46711 | 4.9 | 12 |
| 69 | Inhibition of the mechano-enzymatic amyloidogenesis of transthyretin: role of ligand affinity, binding cooperativity and occupancy of the inner channel. *Scientific Reports*, **2017**, 7, 182 | 4.9 | 25 |

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| 68 | Molecular dynamics simulations of β-microglobulin interaction with hydrophobic surfaces. *Molecular BioSystems*, **2017**, 13, 2625-2637 | | 4 |
| 67 | Short-Chain Alkanethiol Coating for Small-Size Gold Nanoparticles Supporting Protein Stability. *Magnetochemistry*, **2017**, 3, 40 | 3.1 | 3 |
| 66 | Accurate Estimation of the Entropy of Rotation-Translation Probability Distributions. *Journal of Chemical Theory and Computation*, **2016**, 12, 1-8 | 6.4 | 14 |
| 65 | Rational design of mutations that change the aggregation rate of a protein while maintaining its native structure and stability. *Scientific Reports*, **2016**, 6, 25559 | 4.9 | 41 |
| 64 | Accuracy assessment of the linear Poisson-Boltzmann equation and reparametrization of the OBC generalized Born model for nucleic acids and nucleic acid-protein complexes. *Journal of Computational Chemistry*, **2015**, 36, 585-96 | 3.5 | 5 |
| 63 | Probing the influence of citrate-capped gold nanoparticles on an amyloidogenic protein. *ACS Nano*, **2015**, 9, 2600-13 | 16.7 | 68 |
| 62 | Distance-Based Configurational Entropy of Proteins from Molecular Dynamics Simulations. *PLoS ONE*, **2015**, 10, e0132356 | 3.7 | 24 |
| 61 | The Accuracy of Generalized Born Forces **2015**, 143-155 | | |
| 60 | Oligomeric states along the folding pathways of β-microglobulin: kinetics, thermodynamics, and structure. *Journal of Molecular Biology*, **2013**, 425, 2722-36 | 6.5 | 27 |
| 59 | Structure, folding dynamics, and amyloidogenesis of D76N β-microglobulin: roles of shear flow, hydrophobic surfaces, and α-crystallin. *Journal of Biological Chemistry*, **2013**, 288, 30917-30 | 5.4 | 63 |
| 58 | Carnosine inhibits Aβ(1-42) aggregation by perturbing the H-bond network in and around the central hydrophobic cluster. *ChemBioChem*, **2013**, 14, 583-92 | 3.8 | 61 |
| 57 | A differential equation for the Generalized Born radii. *Physical Chemistry Chemical Physics*, **2013**, 15, 9783-91 | 3.9 | 1 |
| 56 | Generalized Born forces: surface integral formulation. *Journal of Chemical Physics*, **2013**, 138, 054112 | 3.9 | 4 |
| 55 | Reduction of conformational mobility and aggregation in W60G β-microglobulin: assessment by 15N NMR relaxation. *Magnetic Resonance in Chemistry*, **2013**, 51, 795-807 | 2.1 | 9 |
| 54 | Monitoring the interaction between β-microglobulin and the molecular chaperone B-crystallin by NMR and mass spectrometry: B-crystallin dissociates β-microglobulin oligomers. *Journal of Biological Chemistry*, **2013**, 288, 17844-58 | 5.4 | 29 |
| 53 | Protein Aggregation Measurement through Electrical Impedance Spectroscopy. *Journal of Physics: Conference Series*, **2013**, 459, 012049 | 0.3 | 2 |
| 52 | Absolute Quantification of Choline-Related Biomarkers in Breast Cancer Biopsies by Liquid Chromatography Electrospray Ionization Mass Spectrometry. *Analytical Cellular Pathology*, **2013**, 36, 71-83 | 3.4 | 16 |
| 51 | Absolute quantification of choline-related biomarkers in breast cancer biopsies by liquid chromatography electrospray ionization mass spectrometry. *Analytical Cellular Pathology*, **2013**, 36, 71-83 | 3.4 | 12 |

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| 50 | Determining the energy landscape of proteins by a fast isotope exchange NMR approach. *Journal of the American Chemical Society*, **2012**, 134, 4457-60 | 16.4 | 9 |
| 49 | Bluues server: electrostatic properties of wild-type and mutated protein structures. *Bioinformatics*, **2012**, 28, 2189-90 | 7.2 | 54 |
| 48 | Single-shot NMR measurement of protein unfolding landscapes. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2012**, 1824, 842-9 | 4 | 7 |
| 47 | Bluues: a program for the analysis of the electrostatic properties of proteins based on generalized Born radii. *BMC Bioinformatics*, **2012**, 13 Suppl 4, S18 | 3.6 | 43 |
| 46 | Pathological self-aggregation of (2)-microglobulin: a challenge for protein biophysics. *Sub-Cellular Biochemistry*, **2012**, 65, 165-83 | 5.5 | 6 |
| 45 | Fast structure similarity searches among protein models: efficient clustering of protein fragments. *Algorithms for Molecular Biology*, **2012**, 7, 16 | 1.8 | 4 |
| 44 | Rapid oligomer formation of human muscle acylphosphatase induced by heparan sulfate. *Nature Structural and Molecular Biology*, **2012**, 19, 547-54, S1-2 | 17.6 | 25 |
| 43 | Studying interactions by molecular dynamics simulations at high concentration. *Journal of Biomedicine and Biotechnology*, **2012**, 2012, 303190 | | 12 |
| 42 | Molecular dynamics simulation of microglobulin in denaturing and stabilizing conditions. *Proteins: Structure, Function and Bioinformatics*, **2011**, 79, 986-1001 | 4.2 | 29 |
| 41 | Effect of tetracyclines on the dynamics of formation and destructuration of beta2-microglobulin amyloid fibrils. *Journal of Biological Chemistry*, **2011**, 286, 2121-31 | 5.4 | 77 |
| 40 | High-performance metabolic marker assessment in breast cancer tissue by mass spectrometry. *Clinical Chemistry and Laboratory Medicine*, **2011**, 49, 317-24 | 5.9 | 8 |
| 39 | Structural and dynamics characteristics of acylphosphatase from Sulfolobus solfataricus in the monomeric state and in the initial native-like aggregates. *Journal of Biological Chemistry*, **2010**, 285, 14689-700 | 5.4 | 20 |
| 38 | Native-unlike long-lived intermediates along the folding pathway of the amyloidogenic protein beta2-microglobulin revealed by real-time two-dimensional NMR. *Journal of Biological Chemistry*, **2010**, 285, 5827-35 | 5.4 | 53 |
| 37 | Folding and fibrillogenesis: clues from beta2-microglobulin. *Journal of Molecular Biology*, **2010**, 401, 286-97 | 6.5 | 33 |
| 36 | Molecular models for intrastrand DNA G-quadruplexes. *BMC Structural Biology*, **2009**, 9, 64 | 2.7 | 14 |
| 35 | Conformational stability of neuroglobin helix F--possible effects on the folding pathway within the globin family. *FEBS Journal*, **2009**, 276, 5177-90 | 5.7 | 8 |
| 34 | Equilibrium unfolding thermodynamics of beta2-microglobulin analyzed through native-state H/D exchange. *Biophysical Journal*, **2009**, 96, 169-79 | 2.9 | 18 |
| 33 | NMR-based homology model for the solution structure of the C-terminal globular domain of EMILIN1. *Journal of Biomolecular NMR*, **2009**, 43, 79-96 | 3 | 7 |

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| 32 | The controlling roles of Trp60 and Trp95 in beta2-microglobulin function, folding and amyloid aggregation properties. *Journal of Molecular Biology*, **2008**, 378, 887-97 | 6.5 | 76 |
| 31 | The solution structure of DNA-free Pax-8 paired box domain accounts for redox regulation of transcriptional activity in the pax protein family. *Journal of Biological Chemistry*, **2008**, 283, 33321-8 | 5.4 | 18 |
| 30 | The solution structure of EMILIN1 globular C1q domain reveals a disordered insertion necessary for interaction with the alpha4beta1 integrin. *Journal of Biological Chemistry*, **2008**, 283, 18947-56 | 5.4 | 25 |
| 29 | Helix mobility and recognition function of the rat thyroid transcription factor 1 homeodomain - hints from 15N-NMR relaxation studies. *FEBS Journal*, **2008**, 275, 435-48 | 5.7 | 7 |
| 28 | Molecular dynamics simulation suggests possible interaction patterns at early steps of beta2-microglobulin aggregation. *Biophysical Journal*, **2007**, 92, 1673-81 | 2.9 | 38 |
| 27 | Scoring predictive models using a reduced representation of proteins: model and energy definition. *BMC Structural Biology*, **2007**, 7, 15 | 2.7 | 20 |
| 26 | Estimation of 3JHN-Halpha and 3JHalpha-Hbeta coupling constants from heteronuclear TOCSY spectra. *Journal of Biomolecular NMR*, **2007**, 39, 213-22 | 3 | 2 |
| 25 | Collagen plays an active role in the aggregation of beta2-microglobulin under physiopathological conditions of dialysis-related amyloidosis. *Journal of Biological Chemistry*, **2006**, 281, 16521-9 | 5.4 | 122 |
| 24 | Variants of beta-microglobulin cleaved at lysine-58 retain the main conformational features of the native protein but are more conformationally heterogeneous and unstable at physiological temperature. *FEBS Journal*, **2006**, 273, 2461-74 | 5.7 | 19 |
| 23 | NMR solution structure of the acylphosphatase from Escherichia coli. *Journal of Biomolecular NMR*, **2006**, 36, 199-204 | 3 | 13 |
| 22 | Structure, conformational stability, and enzymatic properties of acylphosphatase from the hyperthermophile Sulfolobus solfataricus. *Proteins: Structure, Function and Bioinformatics*, **2006**, 62, 64-79 | 4.2 | 37 |
| 21 | Beta2-microglobulin isoforms display an heterogeneous affinity for type I collagen. *Protein Science*, **2005**, 14, 696-702 | 6.3 | 49 |
| 20 | Solution structure of beta(2)-microglobulin and insights into fibrillogenesis. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2005**, 1753, 76-84 | 4 | 22 |
| 19 | Properties of some variants of human beta2-microglobulin and amyloidogenesis. *Journal of Biological Chemistry*, **2004**, 279, 9176-89 | 5.4 | 59 |
| 18 | Sequence-specific backbone NMR assignments for the C-terminal globular domain of EMILIN-1. *Journal of Biomolecular NMR*, **2004**, 29, 91-2 | 3 | 4 |
| 17 | Interaction of copper with cysteine: stability of cuprous complexes and catalytic role of cupric ions in anaerobic thiol oxidation. *Journal of Inorganic Biochemistry*, **2004**, 98, 1495-501 | 4.2 | 148 |
| 16 | beta2-microglobulin H31Y variant 3D structure highlights the protein natural propensity towards intermolecular aggregation. *Journal of Molecular Biology*, **2004**, 335, 1051-64 | 6.5 | 35 |
| 15 | Structural and folding dynamic properties of the T70N variant of human lysozyme. *Journal of Biological Chemistry*, **2003**, 278, 25910-8 | 5.4 | 19 |

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| 14 | Electrostatic compared with hydrophobic interactions between bovine serum amine oxidase and its substrates. *Biochemical Journal*, **2003**, 371, 549-56 | 3.8 | 25 |
| 13 | Biphasic behavior of the kinetics of 31P-containing metabolites in ischemic porcine kidneys. *Transplantation Proceedings*, **2003**, 35, 3111-5 | 1.1 | 3 |
| 12 | The solution structure of human beta2-microglobulin reveals the prodromes of its amyloid transition. *Protein Science*, **2002**, 11, 487-99 | 6.3 | 132 |
| 11 | Binding of cations of group IA and IIA to bovine serum amine oxidase: effect on the activity. *Biophysical Journal*, **2002**, 83, 2231-9 | 2.9 | 10 |
| 10 | Deuterium nuclear magnetic resonance for evaluating the metabolic status of livers subjected to warm ischemia. *Transplantation*, **2001**, 71, 1515-7 | 1.8 | 3 |
| 9 | Improved method for pulse width calibration in indirectly detected experiments. *Magnetic Resonance in Chemistry*, **2001**, 39, 249-250 | 2.1 | 3 |
| 8 | Enzyme mimics complexing Cu(II) ion: structure-function relationships. *Chemical Biology and Drug Design*, **1999**, 54, 491-504 | | 7 |
| 7 | Assessment of pretransplantation warm ischemia time by phosphorus-31 magnetic resonance spectroscopy in pig kidneys. *Transplantation Proceedings*, **1997**, 29, 3415-6 | 1.1 | 5 |
| 6 | Platination of a GG site on single-stranded and double-stranded forms of a 14-base oligonucleotide with diaqua cisplatin followed by NMR and HPLC -- influence of the platinum ligands and base sequence on 5∀G versus 3∀G platination selectivity. *FEBS Journal*, **1997**, 249, 370-82 | | 55 |
| 5 | Structural transitions of a GG-platinated DNA duplex induced by pH, temperature and box A of high-mobility-group protein 1. *FEBS Journal*, **1997**, 243, 782-91 | | 14 |
| 4 | 1H,13C-NMR and X-ray absorption studies of copper(I) glutathione complexes. *FEBS Journal*, **1996**, 236, 697-705 | | 104 |
| 3 | Effect of polyphosphates on the activity of amine oxidases. *BBA - Proteins and Proteomics*, **1995**, 1247, 246-52 | | 5 |
| 2 | Interactions between polyamines and nucleotides studied by31P and1H NMR. *Applied Magnetic Resonance*, **1994**, 7, 89-94 | 0.8 | 1 |
| 1 | Effect of phosphate ion on the activity of bovine plasma amine oxidase. *Biochemical and Biophysical Research Communications*, **1992**, 189, 722-7 | 3.4 | 11 |