

# Anna Rita Bizzarri

## List of Publications by Year in descending order

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88  
papers

2,915  
citations

201674

27  
h-index

182427

51  
g-index

88  
all docs

88  
docs citations

88  
times ranked

3165  
citing authors

#	ARTICLE	IF	CITATIONS
1	Interaction between miR4749 and Human Serum Albumin as Revealed by Fluorescence, FRET, Atomic Force Spectroscopy and Computational Modelling. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1291.	4.1	4
2	A Competitive O-Acetylserine Sulphydrylase Inhibitor Modulates the Formation of Cysteine Synthase Complex. <i>Catalysts</i> , 2021, 11, 700.	3.5	4
3	Solution structure of the anticancer p28 peptide in biomimetic medium. <i>Journal of Peptide Science</i> , 2021, 27, e3357.	1.4	3
4	Temperature Modulation of the DBDp53 Structure as Monitored by Static and Time-Resolved Fluorescence Combined with Molecular Dynamics Simulations. <i>Journal of Physical Chemistry B</i> , 2021, 125, 10166-10173.	2.6	2
5	The use of a commercial ESI Z-spray source for ambient ion soft landing and microdroplet reactivity experiments. <i>International Journal of Mass Spectrometry</i> , 2021, 468, 116658.	1.5	9
6	Direct Interaction of miRNA and circRNA with the Oncosuppressor p53: An Intriguing Perspective in Cancer Research. <i>Cancers</i> , 2021, 13, 6108.	3.7	7
7	A Reliable BioFET Immunosensor for Detection of p53 Tumour Suppressor in Physiological-Like Environment. <i>Sensors</i> , 2020, 20, 6364.	3.8	18
8	Time-Resolved Fluorescence and Essential Dynamics Study on the Structural Heterogeneity of p53DBD Bound to the Anticancer p28 Peptide. <i>Journal of Physical Chemistry B</i> , 2020, 124, 9820-9828.	2.6	3
9	Toward Cancer Diagnostics of the Tumor Suppressor p53 by Surface Enhanced Raman Spectroscopy. <i>Sensors</i> , 2020, 20, 7153.	3.8	3
10	Nanogap Sensors Decorated with SnO <sub>2</sub> Nanoparticles Enable Low-Temperature Detection of Volatile Organic Compounds. <i>ACS Applied Nano Materials</i> , 2020, 3, 3337-3346.	5.0	13
11	Investigation of a Direct Interaction between miR4749 and the Tumor Suppressor p53 by Fluorescence, FRET and Molecular Modeling. <i>Biomolecules</i> , 2020, 10, 346.	4.0	8
12	Portable Immunosensor Based on Extended Gate <sup>+</sup> Field Effect Transistor for Rapid, Sensitive Detection of Cancer Markers. <i>Proceedings (mdpi)</i> , 2019, 15, .	0.2	1
13	Raman Evidence of p53-DBD Disorder Decrease upon Interaction with the Anticancer Protein Azurin. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3078.	4.1	13
14	Probing direct interaction of oncomiR-21-3p with the tumor suppressor p53 by fluorescence, FRET and atomic force spectroscopy. <i>Archives of Biochemistry and Biophysics</i> , 2019, 671, 35-41.	3.0	16
15	Interaction Force Fluctuations in Antigen <sup>+</sup> Antibody Biorecognition Studied by Atomic Force Spectroscopy. <i>ACS Omega</i> , 2019, 4, 3627-3634.	3.5	3
16	Interaction of human hemoglobin and semi-hemoglobins with the <i>Staphylococcus aureus</i> hemophore IsdB: a kinetic and mechanistic insight. <i>Scientific Reports</i> , 2019, 9, 18629.	3.3	21
17	Interaction of the anticancer p28 peptide with p53-DBD as studied by fluorescence, FRET, docking and MD simulations. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2019, 1863, 342-350.	2.4	20
18	Surface enhanced Raman spectroscopy based immunosensor for ultrasensitive and selective detection of wild type p53 and mutant p53R175H. <i>Analytica Chimica Acta</i> , 2018, 1029, 86-96.	5.4	29

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19	Imaging and kinetics of the bimolecular complex formed by the tumor suppressor p53 with ubiquitin ligase COP1 as studied by atomic force microscopy and surface plasmon resonance. <i>International Journal of Nanomedicine</i> , 2018, Volume 13, 251-259.	6.7	16
20	Structural Characterization of the Intrinsically Disordered Protein p53 Using Raman Spectroscopy. <i>Applied Spectroscopy</i> , 2017, 71, 823-832.	2.2	26
21	Binding of Amphipathic Cell Penetrating Peptide p28 to Wild Type and Mutated p53 as studied by Raman, Atomic Force and Surface Plasmon Resonance spectroscopies. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017, 1861, 910-921.	2.4	20
22	Binding kinetics of mutant p53R175H with wild type p53 and p63: A Surface Plasmon Resonance and Atomic Force Spectroscopy study. <i>Biophysical Chemistry</i> , 2017, 228, 55-61.	2.8	5
23	Structure, Dynamics, and Electron Transfer of Azurin Bound to a Gold Electrode. <i>Langmuir</i> , 2017, 33, 9190-9200.	3.5	5
24	Vibrational Changes Induced by Electron Transfer in Surface Bound Azurin Metalloprotein Studied by Tip-Enhanced Raman Spectroscopy and Scanning Tunneling Microscopy. <i>ACS Nano</i> , 2017, 11, 12824-12831.	14.6	25
25	Surface Plasmon Resonance Sensing of Biorecognition Interactions within the Tumor Suppressor p53 Network. <i>Sensors</i> , 2017, 17, 2680.	3.8	22
26	MDM2&ndash;MDM4 molecular interaction investigated by atomic force spectroscopy and surface plasmon resonance. <i>International Journal of Nanomedicine</i> , 2016, Volume 11, 4221-4229.	6.7	11
27	Electron transfer, conduction and biorecognition properties of the redox metalloprotein Azurin assembled onto inorganic substrates. <i>European Polymer Journal</i> , 2016, 83, 407-427.	5.4	32
28	Kinetics and binding geometries of the complex between $\hat{2}$ -microglobulin and its antibody: An AFM and SPR study. <i>Biophysical Chemistry</i> , 2016, 211, 19-27.	2.8	18
29	Energy landscape investigation by wavelet transform analysis of atomic force spectroscopy data in a biorecognition experiment. <i>Journal of Biological Physics</i> , 2016, 42, 167-176.	1.5	1
30	Chirality Switching within an Anionic Cell-Penetrating Peptide Inhibits Translocation without Affecting Preferential Entry. <i>Molecular Pharmaceutics</i> , 2015, 12, 140-149.	4.6	31
31	Calcium Ions Modulate the Mechanics of Tomato Bushy Stunt Virus. <i>Biophysical Journal</i> , 2015, 109, 390-397.	0.5	25
32	Electron tunnelling through single azurin molecules can be on/off switched by voltage pulses. <i>Applied Physics Letters</i> , 2015, 106, 183701.	3.3	15
33	A nanotechnological, molecular-modeling, and immunological approach to study the interaction of the anti-tumorigenic peptide p28 with the p53 family of proteins. <i>International Journal of Nanomedicine</i> , 2014, 9, 1799.	6.7	11
34	Binding of azurin to cytochrome <i>c</i> 551 as investigated by surface plasmon resonance and fluorescence. <i>Journal of Molecular Recognition</i> , 2014, 27, 124-130.	2.1	14
35	Antigen&quot;antibody biorecognition events as discriminated by noise analysis of force spectroscopy curves. <i>Nanotechnology</i> , 2014, 25, 335102.	2.6	17
36	Interaction of mutant p53 with p73: A Surface Plasmon Resonance and Atomic Force Spectroscopy study. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 1958-1964.	2.4	15

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37	Excitation of the ligand-to-metal charge transfer band induces electron tunnelling in azurin. <i>Applied Physics Letters</i> , 2014, 104, 093702.	3.3	10
38	Conductive atomic force microscopy study of single molecule electron transport through the Azurin-gold nanoparticle system. <i>Applied Physics Letters</i> , 2013, 102, 203704.	3.3	14
39	$1/f$ Noise in the Dynamic Force Spectroscopy Curves Signals the Occurrence of Biorecognition. <i>Physical Review Letters</i> , 2013, 110, 048104.	7.8	21
40	Ultrafast Pump-Probe Study of the Excited-State Charge-Transfer Dynamics in Blue Copper Rusticyanin. <i>Journal of Physical Chemistry B</i> , 2012, 116, 4192-4198.	2.6	15
41	Surface-enhanced Raman scattering detection of wild-type and mutant p53 proteins at very low concentration in human serum. <i>Analytical Biochemistry</i> , 2012, 421, 9-15.	2.4	70
42	Free energy evaluation of the p53-Mdm2 complex from unbinding work measured by dynamic force spectroscopy. <i>Physical Chemistry Chemical Physics</i> , 2011, 13, 2738-2743.	2.8	14
43	Steered Molecular Dynamics Simulations of the Electron Transfer Complex between Azurin and Cytochrome <i>c</i> <sub>551</sub> . <i>Journal of Physical Chemistry B</i> , 2011, 115, 1211-1219.	2.6	9
44	Interaction of an anticancer peptide fragment of azurin with p53 and its isolated domains studied by atomic force spectroscopy. <i>International Journal of Nanomedicine</i> , 2011, 6, 3011.	6.7	50
45	SERS-based nanobiosensing for ultrasensitive detection of the p53 tumor suppressor. <i>International Journal of Nanomedicine</i> , 2011, 6, 2033.	6.7	34
46	Azurin modulates the association of Mdm2 with p53: SPR evidence from interaction of the full-length proteins. <i>Journal of Molecular Recognition</i> , 2011, 24, 707-714.	2.1	26
47	Modelling the interaction between the p53 DNA-binding domain and the p28 peptide fragment of Azurin. <i>Journal of Molecular Recognition</i> , 2011, 24, 1043-1055.	2.1	25
48	Interaction of p53 with Mdm2 and azurin as studied by atomic force spectroscopy. <i>Journal of Molecular Recognition</i> , 2010, 23, 343-351.	2.1	25
49	The application of atomic force spectroscopy to the study of biological complexes undergoing a biorecognition process. <i>Chemical Society Reviews</i> , 2010, 39, 734-749.	38.1	120
50	Modeling the interaction between the N-terminal domain of the tumor suppressor p53 and azurin. <i>Journal of Molecular Recognition</i> , 2009, 22, 215-222.	2.1	23
51	A combined atomic force microscopy imaging and docking study to investigate the complex between p53 DNA binding domain and Azurin. <i>Journal of Molecular Recognition</i> , 2009, 22, 506-515.	2.1	13
52	Surface-enhanced Raman spectroscopy combined with atomic force microscopy for ultrasensitive detection of thrombin. <i>Analytical Biochemistry</i> , 2009, 393, 149-154.	2.4	23
53	Atomic Force Spectroscopy in Biological Complex Formation: Strategies and Perspectives. <i>Journal of Physical Chemistry B</i> , 2009, 113, 16449-16464.	2.6	44
54	Probing the interaction between p53 and the bacterial protein azurin by single molecule force spectroscopy. <i>Journal of Molecular Recognition</i> , 2008, 21, 63-70.	2.1	59

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55	Statistical analysis of intensity fluctuations in single molecule SERS spectra. <i>Physical Chemistry Chemical Physics</i> , 2007, 9, 5315.	2.8	23
56	Docking and molecular dynamics simulation of the Azurin-Cytochrome c551 electron transfer complex. <i>Journal of Molecular Recognition</i> , 2007, 20, 122-131.	2.1	24
57	Docking study and free energy simulation of the complex between p53 DNA-binding domain and azurin. <i>Journal of Molecular Recognition</i> , 2007, 20, 215-226.	2.1	54
58	SERS detection of thrombin by protein recognition using functionalized gold nanoparticles. <i>Nanomedicine: Nanotechnology, Biology, and Medicine</i> , 2007, 3, 306-310.	3.3	58
59	Quenching and Blinking of Fluorescence of a Single Dye Molecule Bound to Gold Nanoparticles. <i>Journal of Physical Chemistry B</i> , 2006, 110, 16491-16498.	2.6	85
60	Time-dependent study of single-molecule SERS signal from yeast cytochrome c. <i>Chemical Physics</i> , 2006, 326, 356-362.	1.9	26
61	Topological and dynamical properties of Azurin anchored to a gold substrate as investigated by molecular dynamics simulation. <i>Biophysical Chemistry</i> , 2006, 122, 206-214.	2.8	21
62	Electron tunneling in a metal-protein-metal junction investigated by scanning tunneling and conductive atomic force spectroscopies. <i>Applied Physics Letters</i> , 2006, 89, 183125.	3.3	23
63	Temporal Fluctuations in Single-Molecule SERS Spectra. , 2006, , 279-296.		7
64	Temporal Fluctuations in Single-Molecule SERS Spectra. , 2006, , 279-296.		0
65	Single-molecule detection of yeast cytochrome c by Surface-Enhanced Raman Spectroscopy. <i>Biophysical Chemistry</i> , 2005, 113, 41-51.	2.8	89
66	LÃ©vy Statistics of Vibrational Mode Fluctuations of Single Molecules from Surface-Enhanced Raman Scattering. <i>Physical Review Letters</i> , 2005, 94, 068303.	7.8	51
67	SERS and Tunneling Spectroscopy Investigation of Iron-Protoporphyrin IX Adsorbed on a Silver Tip. <i>Journal of Physical Chemistry B</i> , 2005, 109, 16571-16574.	2.6	15
68	Optical Spectroscopic Investigation of the Alkaline Transition in Ume cyanin from Horseradish Root. <i>Biochemistry</i> , 2005, 44, 16090-16097.	2.5	14
69	Neutron scattering and molecular dynamics simulation: a conjugate approach to investigate the dynamics of electron transfer proteins. <i>Journal of Physics Condensed Matter</i> , 2004, 16, R83-R110.	1.8	13
70	Evidence of electron-transfer in the SERS spectra of a single iron-protoporphyrin IX molecule. <i>Chemical Physics Letters</i> , 2004, 395, 222-226.	2.6	26
71	Topological and Electron-Transfer Properties of Yeast Cytochrome c Adsorbed on Bare Gold Electrodes. <i>ChemPhysChem</i> , 2003, 4, 1183-1188.	2.1	49
72	A Combined Atomic Force Microscopy and Molecular Dynamics Simulation Study on a Plastocyanin Mutant Chemisorbed on a Gold Surface. <i>ChemPhysChem</i> , 2003, 4, 1189-1195.	2.1	22

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73	Temporal fluctuations in the SERRS spectra of single iron protoporphyrin IX molecule. <i>Chemical Physics</i> , 2003, 290, 297-306.	1.9	39
74	MD simulation of a plastocyanin mutant adsorbed onto a gold surface. <i>Biophysical Chemistry</i> , 2003, 106, 111-123.	2.8	36
75	Excited state charge-transfer dynamics study of poplar plastocyanin by ultrafast pump-probe spectroscopy and molecular dynamics simulation. <i>Biophysical Chemistry</i> , 2003, 106, 221-231.	2.8	22
76	Surface-Enhanced Resonance Raman Spectroscopy Signals from Single Myoglobin Molecules. <i>Applied Spectroscopy</i> , 2002, 56, 1531-1537.	2.2	91
77	Molecular Dynamics of Water at the Protein-Solvent Interface. <i>Journal of Physical Chemistry B</i> , 2002, 106, 6617-6633.	2.6	484
78	Vibrational coherence in Azurin with impulsive excitation of the LMCT absorption band. <i>Chemical Physics Letters</i> , 2002, 362, 497-503.	2.6	31
79	Concerted motions in copper plastocyanin and azurin: an essential dynamics study. <i>Biophysical Chemistry</i> , 2001, 90, 45-56.	2.8	46
80	Molecular dynamics simulation and essential dynamics study of mutated plastocyanin: structural, dynamical and functional effects of a disulfide bridge insertion at the protein surface. <i>Biophysical Chemistry</i> , 2001, 92, 183-199.	2.8	26
81	Intensity fluctuations of the copper site resonant vibrational modes as observed by MD simulation in single plastocyanin molecule. <i>Chemical Physics Letters</i> , 2001, 349, 503-510.	2.6	12
82	Glasslike dynamical behavior of the plastocyanin hydration water. <i>Physical Review E</i> , 2000, 62, 3991-3999.	2.1	48
83	Long-term molecular dynamics simulation of copper azurin: structure, dynamics and functionality. <i>Biophysical Chemistry</i> , 1999, 78, 247-257.	2.8	33
84	Incoherent neutron scattering of copper azurin: a comparison with molecular dynamics simulation results. <i>European Biophysics Journal</i> , 1999, 28, 447-456.	2.2	44
85	Role of interfacial water in the molecular dynamics-simulated dynamical transition of plastocyanin. <i>Chemical Physics Letters</i> , 1998, 291, 7-14.	2.6	52
86	Water dynamical anomalies evidenced by molecular-dynamics simulations at the solvent-protein interface. <i>Physical Review E</i> , 1998, 57, 3315-3325.	2.1	166
87	Flickering noise in the potential energy fluctuations of proteins as investigated by MD simulation. <i>Physics Letters, Section A: General, Atomic and Solid State Physics</i> , 1997, 236, 596-601.	2.1	25
88	Molecular dynamics simulation evidence of anomalous diffusion of protein hydration water. <i>Physical Review E</i> , 1996, 53, R3040-R3043.	2.1	79