

Irina Kuznetsova

List of Publications by Year in descending order

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176
papers

6,371
citations

76196

40
h-index

79541

73
g-index

196
all docs

196
docs citations

196
times ranked

7034
citing authors

#	ARTICLE	IF	CITATIONS
1	What Macromolecular Crowding Can Do to a Protein. <i>International Journal of Molecular Sciences</i> , 2014, 15, 23090-23140.	1.8	425
2	Thioflavin T as a Molecular Rotor: Fluorescent Properties of Thioflavin T in Solvents with Different Viscosity. <i>Journal of Physical Chemistry B</i> , 2008, 112, 15893-15902.	1.2	300
3	Computational Study of Thioflavin T Torsional Relaxation in the Excited State. <i>Journal of Physical Chemistry A</i> , 2007, 111, 4829-4835.	1.1	206
4	Intrinsically disordered proteins as crucial constituents of cellular aqueous two phase systems and coacervates. <i>FEBS Letters</i> , 2015, 589, 15-22.	1.3	203
5	Spectral Properties of Thioflavin T and Its Complexes with Amyloid Fibrils. <i>Journal of Applied Spectroscopy</i> , 2003, 70, 868-874.	0.3	199
6	Spectral Properties of Thioflavin T in Solvents with Different Dielectric Properties and in a Fibril-Incorporated Form. <i>Journal of Proteome Research</i> , 2007, 6, 1392-1401.	1.8	187
7	Fluorescence of Dyes in Solutions with High Absorbance. Inner Filter Effect Correction. <i>PLoS ONE</i> , 2014, 9, e103878.	1.1	182
8	The protein kingdom extended: Ordered and intrinsically disordered proteins, their folding, supramolecular complex formation, and aggregation. <i>Progress in Biophysics and Molecular Biology</i> , 2010, 102, 73-84.	1.4	181
9	Transcriptomic and proteomic landscape of mitochondrial dysfunction reveals secondary coenzyme Q deficiency in mammals. <i>ELife</i> , 2017, 6, .	2.8	169
10	Beyond the Excluded Volume Effects: Mechanistic Complexity of the Crowded Milieu. <i>Molecules</i> , 2015, 20, 1377-1409.	1.7	157
11	Fluorescence Quantum Yield of Thioflavin T in Rigid Isotropic Solution and Incorporated into the Amyloid Fibrils. <i>PLoS ONE</i> , 2010, 5, e15385.	1.1	152
12	Modern fluorescent proteins: from chromophore formation to novel intracellular applications. <i>BioTechniques</i> , 2011, 51, 313-327.	0.8	137
13	Fluorescent Proteins as Biomarkers and Biosensors: Throwing Color Lights on Molecular and Cellular Processes. <i>Current Protein and Peptide Science</i> , 2008, 9, 338-369.	0.7	136
14	Use of the Phase Diagram Method to Analyze the Protein Unfolding-Refolding Reactions: Fishing Out the "Invisible" Intermediates. <i>Journal of Proteome Research</i> , 2004, 3, 485-494.	1.8	130
15	Partially Folded Conformations in the Folding Pathway of Bovine Carbonic Anhydrase II: A Fluorescence Spectroscopic Analysis. <i>ChemBioChem</i> , 2001, 2, 813.	1.3	121
16	Hierarchical RNA Processing Is Required for Mitochondrial Ribosome Assembly. <i>Cell Reports</i> , 2016, 16, 1874-1890.	2.9	116
17	Polarization-dependent optical 2D Fourier transform spectroscopy of semiconductors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 14227-14232.	3.3	110
18	High Stability of Discosoma DsRed As Compared to Aequorea EGFP. <i>Biochemistry</i> , 2003, 42, 7879-7884.	1.2	108

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19	Unraveling multistate unfolding of rabbit muscle creatine kinase. <i>BBA - Proteins and Proteomics</i> , 2002, 1596, 138-155.	2.1	96
20	Stochasticity of Biological Soft Matter: Emerging Concepts in Intrinsically Disordered Proteins and Biological Phase Separation. <i>Trends in Biochemical Sciences</i> , 2019, 44, 716-728.	3.7	94
21	Interaction of Thioflavin T with Amyloid Fibrils: Stoichiometry and Affinity of Dye Binding, Absorption Spectra of Bound Dye. <i>Journal of Physical Chemistry B</i> , 2011, 115, 11519-11524.	1.2	92
22	Comparative Studies on the Structure and Stability of Fluorescent Proteins EGFP, zFP506, mRFP1, ΔC -dimer2, and DsRed1. <i>Biochemistry</i> , 2004, 43, 14913-14923.	1.2	85
23	Interaction of Thioflavin T with Amyloid Fibrils: Fluorescence Quantum Yield of Bound Dye. <i>Journal of Physical Chemistry B</i> , 2012, 116, 2538-2544.	1.2	84
24	CirGO: an alternative circular way of visualising gene ontology terms. <i>BMC Bioinformatics</i> , 2019, 20, 84.	1.2	84
25	Differences in the Pathways of Proteins Unfolding Induced by Urea and Guanidine Hydrochloride: Molten Globule State and Aggregates. <i>PLoS ONE</i> , 2010, 5, e15035.	1.1	77
26	Beta-Barrel Scaffold of Fluorescent Proteins. <i>International Review of Cell and Molecular Biology</i> , 2013, 302, 221-278.	1.6	75
27	Reevaluation of ANS Binding to Human and Bovine Serum Albumins: Key Role of Equilibrium Microdialysis in Ligand K_d Receptor Binding Characterization. <i>PLoS ONE</i> , 2012, 7, e40845.	1.1	71
28	Intrinsically disordered proteins in crowded milieu: when chaos prevails within the cellular gumbo. <i>Cellular and Molecular Life Sciences</i> , 2018, 75, 3907-3929.	2.4	71
29	Thioflavin T fluoresces as excimer in highly concentrated aqueous solutions and as monomer being incorporated in amyloid fibrils. <i>Scientific Reports</i> , 2017, 7, 2146.	1.6	66
30	Analyzing Thioflavin T Binding to Amyloid Fibrils by an Equilibrium Microdialysis-Based Technique. <i>PLoS ONE</i> , 2012, 7, e30724.	1.1	63
31	Concerted regulation of mitochondrial and nuclear non-coding RNAs by a dual-targeted RNase Z. <i>EMBO Reports</i> , 2018, 19, .	2.0	60
32	Intrinsic Fluorescence of Actin. <i>Journal of Fluorescence</i> , 2003, 13, 41-57.	1.3	57
33	A New Trend in the Experimental Methodology for the Analysis of the Thioflavin T Binding to Amyloid Fibrils. <i>Molecular Neurobiology</i> , 2012, 45, 488-498.	1.9	56
34	Accumulation of storage proteins in plant seeds is mediated by amyloid formation. <i>PLoS Biology</i> , 2020, 18, e3000564.	2.6	53
35	PTCD1 Is Required for 16S rRNA Maturation Complex Stability and Mitochondrial Ribosome Assembly. <i>Cell Reports</i> , 2018, 23, 127-142.	2.9	51
36	Physico-chemical properties of actin cleaved with bacterial protease from <i>E. coli</i> A2 strain. <i>FEBS Letters</i> , 1991, 279, 49-51.	1.3	50

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37	Multi-functionality of proteins involved in GPCR and G protein signaling: making sense of structureâ€“function continuum with intrinsic disorder-based proteoforms. <i>Cellular and Molecular Life Sciences</i> , 2019, 76, 4461-4492.	2.4	47
38	Fidelity of translation initiation is required for coordinated respiratory complex assembly. <i>Science Advances</i> , 2019, 5, eaay2118.	4.7	47
39	Effect of the fluorescent probes ThT and ANS on the mature amyloid fibrils. <i>Prion</i> , 2020, 14, 67-75.	0.9	46
40	Effect of Self-Association on the Structural Organization of Partially Folded Proteins: Inactivated Actin. <i>Biophysical Journal</i> , 1999, 77, 2788-2800.	0.2	45
41	The Place of Inactivated Actin and Its Kinetic Predecessor in Actin Foldingâˆ“Unfolding. <i>Biochemistry</i> , 2002, 41, 13127-13132.	1.2	45
42	Proteomic analysis of the 20S proteasome (PSMA3)-interacting proteins reveals a functional link between the proteasome and mRNA metabolism. <i>Biochemical and Biophysical Research Communications</i> , 2011, 416, 258-265.	1.0	45
43	Liquidâ€“liquid phase separation as an organizing principle of intracellular space: overview of the evolution of the cell compartmentalization concept. <i>Cellular and Molecular Life Sciences</i> , 2022, 79, 251.	2.4	42
44	The environment of the tryptophan residue in <i>Pseudomonas aeruginosa</i> azurin and its fluorescence properties. <i>Biophysical Chemistry</i> , 1985, 23, 79-89.	1.5	41
45	Contribution of separate tryptophan residues to intrinsic fluorescence of actin. Analysis of 3D structure. <i>FEBS Letters</i> , 1999, 452, 205-210.	1.3	39
46	Conformational changes in subdomain I of actin induced by proteolytic cleavage within the DNase I-binding loop: energy transfer from tryptophan to AEDANS. <i>FEBS Letters</i> , 1996, 383, 105-108.	1.3	37
47	Kinetics of Actin Unfolding Induced by Guanidine Hydrochlorideâ€“. <i>Biochemistry</i> , 2002, 41, 1014-1019.	1.2	37
48	Changes of structure and intramolecular mobility in the course of actin denaturation. <i>Biophysical Chemistry</i> , 1988, 32, 73-78.	1.5	36
49	The Structure and Dynamics of Partially Folded Actinâ€“. <i>Biochemistry</i> , 1999, 38, 6261-6269.	1.2	36
50	Simultaneous processing and degradation of mitochondrial RNAs revealed by circularized RNA sequencing. <i>Nucleic Acids Research</i> , 2017, 45, 5487-5500.	6.5	36
51	Adult-onset obesity is triggered by impaired mitochondrial gene expression. <i>Science Advances</i> , 2017, 3, e1700677.	4.7	36
52	Lung Cancer Risk from Plutonium: A Pooled Analysis of the Mayak and Sellafield Worker Cohorts. <i>Radiation Research</i> , 2017, 188, 725.	0.7	36
53	Investigation of Î±-Synuclein Amyloid Fibrils Using the Fluorescent Probe Thioflavin T. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2486.	1.8	36
54	Allosteric effects of chromophore interaction with dimeric near-infrared fluorescent proteins engineered from bacterial phytochromes. <i>Scientific Reports</i> , 2016, 6, 18750.	1.6	35

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55	Hydrophobic interactions and ionic networks play an important role in thermal stability and denaturation mechanism of the porcine odorant-binding protein. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 71, 35-44.	1.5	32
56	High Fluorescence Anisotropy of Thioflavin T in Aqueous Solution Resulting from Its Molecular Rotor Nature. <i>Analytical Chemistry</i> , 2016, 88, 718-724.	3.2	32
57	Oncolytic influenza A virus expressing interleukin-15 decreases tumor growth in vivo. <i>Surgery</i> , 2017, 161, 735-746.	1.0	31
58	Protein unfolding in crowded milieu: what crowding can do to a protein undergoing unfolding?. <i>Journal of Biomolecular Structure and Dynamics</i> , 2016, 34, 2155-2170.	2.0	28
59	Unfolding and Refolding of the Glutamine-Binding Protein from <i>Escherichia coli</i> and Its Complex with Glutamine Induced by Guanidine Hydrochloride. <i>Biochemistry</i> , 2005, 44, 5625-5633.	1.2	27
60	Stability and Dynamics of the Porcine Odorant-Binding Protein. <i>Biochemistry</i> , 2007, 46, 11120-11127.	1.2	27
61	Different conditions of fibrillogenesis cause polymorphism of lysozyme amyloid fibrils. <i>Journal of Molecular Structure</i> , 2017, 1140, 52-58.	1.8	27
62	Stoichiometry and Affinity of Thioflavin T Binding to Sup35p Amyloid Fibrils. <i>PLoS ONE</i> , 2016, 11, e0156314.	1.1	23
63	Two Novel Amyloid Proteins, RopA and RopB, from the Root Nodule Bacterium <i>Rhizobium leguminosarum</i> . <i>Biomolecules</i> , 2019, 9, 694.	1.8	23
64	Stress signaling and cellular proliferation reverse the effects of mitochondrial mistranslation. <i>EMBO Journal</i> , 2019, 38, e102155.	3.5	21
65	A knot in the protein structure – probing the near-infrared fluorescent protein iRFP designed from a bacterial phytochrome. <i>FEBS Journal</i> , 2014, 281, 2284-2298.	2.2	20
66	Prion-associated proteins in yeast: comparative analysis of isogenic [PSI ⁺] and [psi ⁺] strains. <i>Yeast</i> , 2009, 26, 611-631.	0.8	19
67	Establishment of a Chimeric, Replication-Deficient Influenza A Virus Vector by Modulation of Splicing Efficiency. <i>Journal of Virology</i> , 2011, 85, 2469-2473.	1.5	19
68	Distinct Effects of Guanidine Thiocyanate on the Structure of Superfolder GFP. <i>PLoS ONE</i> , 2012, 7, e48809.	1.1	19
69	Sensitivity of Superfolder GFP to Ionic Agents. <i>PLoS ONE</i> , 2014, 9, e110750.	1.1	18
70	Conformational Change of the Dimeric DsbC Molecule Induced by GdnHCl. A Study by Intrinsic Fluorescence. <i>Biochemistry</i> , 2004, 43, 5296-5303.	1.2	17
71	Structural Features of Amyloid Fibrils Formed from the Full-Length and Truncated Forms of Beta-2-Microglobulin Probed by Fluorescent Dye Thioflavin T. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2762.	1.8	17
72	The Role of Non-Specific Interactions in Canonical and ALT-Associated PML-Bodies Formation and Dynamics. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5821.	1.8	17

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73	Structural Analogue of Thioflavin T, DMASEBT, as a Tool for Amyloid Fibrils Study. <i>Analytical Chemistry</i> , 2019, 91, 3131-3140.	3.2	16
74	Spectral characteristics of the mutant form GGBP/H152C of D-glucose/D-galactose-binding protein labeled with fluorescent dye BADAN: influence of external factors. <i>PeerJ</i> , 2014, 2, e275.	0.9	16
75	Fluorescence Properties of Glutamine-Binding Protein from <i>Escherichia coli</i> and Its Complex with Glutamine. <i>Journal of Proteome Research</i> , 2005, 4, 417-423.	1.8	15
76	Understanding the role of Arg96 in structure and stability of green fluorescent protein. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 73, 539-551.	1.5	15
77	Binding Stoichiometry and Affinity of Fluorescent Dyes to Proteins in Different Structural States. <i>Methods in Molecular Biology</i> , 2012, 895, 441-460.	0.4	15
78	Spectral Manifestations of Thioflavin T Aggregation. <i>Journal of Applied Spectroscopy</i> , 2015, 82, 33-39.	0.3	15
79	Denaturant effect on amyloid fibrils: Declusterization, depolymerization, denaturation and reassembly. <i>International Journal of Biological Macromolecules</i> , 2020, 150, 681-694.	3.6	15
80	Actinous enigma or enigmatic actin. <i>Intrinsically Disordered Proteins</i> , 2014, 2, e34500.	1.9	14
81	Stabilization of structure in near-infrared fluorescent proteins by binding of biliverdin chromophore. <i>Journal of Molecular Structure</i> , 2017, 1140, 22-31.	1.8	14
82	Trans-2-[4-(dimethylamino)styryl]-3-ethyl-1,3-benzothiazolium perchlorate - New fluorescent dye for testing of amyloid fibrils and study of their structure. <i>Dyes and Pigments</i> , 2018, 157, 385-395.	2.0	14
83	Trypsin Induced Degradation of Amyloid Fibrils. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4828.	1.8	14
84	New Insight in Protein-Ligand Interactions. 2. Stability and Properties of Two Mutant Forms of the <i>d</i> -Galactose- <i>d</i> -Glucose-Binding Protein from <i>E. coli</i> . <i>Journal of Physical Chemistry B</i> , 2011, 115, 9022-9032.	1.2	13
85	New Insight into Protein-Ligand Interactions. The Case of the <i>d</i> -Galactose/ <i>d</i> -Glucose-Binding Protein from <i>Escherichia coli</i> . <i>Journal of Physical Chemistry B</i> , 2011, 115, 2765-2773.	1.2	13
86	Photophysical Properties of Fluorescent Probe Thioflavin T in Crowded Milieu. <i>Journal of Spectroscopy</i> , 2017, 2017, 1-10.	0.6	13
87	New findings on GFP-like protein application as fluorescent tags: Fibrillogenesis, oligomerization, and amorphous aggregation. <i>International Journal of Biological Macromolecules</i> , 2021, 192, 1304-1310.	3.6	13
88	Correlation between Polymerizability and Conformation in Scallop β -Like Actin and Rabbit Skeletal Muscle β -Actin. <i>Archives of Biochemistry and Biophysics</i> , 1999, 368, 105-111.	1.4	12
89	Monitoring of Actin Unfolding by Room Temperature Tryptophan Phosphorescence. <i>Biochemistry</i> , 2003, 42, 13551-13557.	1.2	12
90	Highly UV-Absorbing Complex in Selenomethionine-Substituted Alcohol Dehydrogenase from <i>Sulfolobus solfataricus</i> . <i>Journal of Proteome Research</i> , 2004, 3, 613-620.	1.8	12

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91	Different disturbances of one pathway of protein unfolding. Actin folding-unfolding and misfolding. <i>Cell Biology International</i> , 2007, 31, 405-412.	1.4	12
92	The effect of red pigment on the amyloidization of yeast proteins. <i>Yeast</i> , 2011, 28, 505-526.	0.8	12
93	Peculiarities of the Super-Folder GFP Folding in a Crowded Milieu. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1805.	1.8	12
94	Targeting an Oncolytic Influenza A Virus to Tumor Tissue by Elastase. <i>Molecular Therapy - Oncolytics</i> , 2017, 7, 37-44.	2.0	12
95	Expression of recombinant GFP-actin fusion protein in the methylotrophic yeast. <i>FEMS Yeast Research</i> , 2003, 3, 105-111.	1.1	11
96	Spectral properties and factors determining high quantum yield of thioflavin T incorporated in amyloid fibrils. <i>Spectroscopy</i> , 2010, 24, 169-172.	0.8	11
97	Adaptive mutation in nuclear export protein allows stable transgene expression in a chimaeric influenza A virus vector. <i>Journal of General Virology</i> , 2014, 95, 337-349.	1.3	11
98	Interaction of Biliverdin Chromophore with Near-Infrared Fluorescent Protein BphP1-FP Engineered from Bacterial Phytochrome. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1009.	1.8	11
99	Structure and Conformational Properties of d-Glucose/d-Galactose-Binding Protein in Crowded Milieu. <i>Molecules</i> , 2017, 22, 244.	1.7	11
100	M60-like metalloprotease domain of the Escherichia coli YghJ protein forms amyloid fibrils. <i>PLoS ONE</i> , 2018, 13, e0191317.	1.1	11
101	Folding of poly-amino acids and intrinsically disordered proteins in overcrowded milieu induced by pH change. <i>International Journal of Biological Macromolecules</i> , 2019, 125, 244-255.	3.6	11
102	Mitochondrial mistranslation modulated by metabolic stress causes cardiovascular disease and reduced lifespan. <i>Aging Cell</i> , 2021, 20, e13408.	3.0	11
103	β -Barrels and Amyloids: Structural Transitions, Biological Functions, and Pathogenesis. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11316.	1.8	11
104	Native globular actin has a thermodynamically unstable quasi-stationary structure with elements of intrinsic disorder. <i>FEBS Journal</i> , 2016, 283, 438-445.	2.2	10
105	Protein folding and stability in the presence of osmolytes. <i>Biophysics (Russian Federation)</i> , 2016, 61, 185-192.	0.2	10
106	Structural and functional characteristics of various forms of red pigment of yeast <i>Saccharomyces cerevisiae</i> and its synthetic analog. <i>Cell and Tissue Biology</i> , 2013, 7, 86-94.	0.2	9
107	Investigation of the kinetics of insulin amyloid fibrils formation. <i>Cell and Tissue Biology</i> , 2014, 8, 186-191.	0.2	9
108	The trehalose/maltose-binding protein as the sensitive element of a glucose biosensor. <i>Optical Materials</i> , 2014, 36, 1676-1679.	1.7	9

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109	The Quaternary Structure of the Recombinant Bovine Odorant-Binding Protein Is Modulated by Chemical Denaturants. PLoS ONE, 2014, 9, e85169.	1.1	9
110	New Evidence of the Importance of Weak Interactions in the Formation of PML-Bodies. International Journal of Molecular Sciences, 2022, 23, 1613.	1.8	9
111	Effects of low urea concentrations on protein-water interactions. Journal of Biomolecular Structure and Dynamics, 2017, 35, 207-218.	2.0	8
112	The effects of crowding agents Dextran-70k and PEG-8k on actin structure and unfolding reaction. Journal of Molecular Structure, 2017, 1140, 46-51.	1.8	8
113	Near-Infrared Markers based on Bacterial Phytochromes with Phycocyanobilin as a Chromophore. International Journal of Molecular Sciences, 2019, 20, 6067.	1.8	8
114	Effect of red pigment on amyloidization of yeast. Cell and Tissue Biology, 2010, 4, 152-166.	0.2	7
115	Thioflavin T Interaction with Acetylcholinesterase: New Evidence of 1:1 Binding Stoichiometry Obtained with Samples Prepared by Equilibrium Microdialysis. ACS Chemical Neuroscience, 2018, 9, 1793-1801.	1.7	7
116	Folding perspectives of an intrinsically disordered transactivation domain and its single mutation breaking the folding propensity. International Journal of Biological Macromolecules, 2020, 155, 1359-1372.	3.6	7
117	Alpha-B-Crystallin Effect on Mature Amyloid Fibrils: Different Degradation Mechanisms and Changes in Cytotoxicity. International Journal of Molecular Sciences, 2020, 21, 7659.	1.8	7
118	OmicsVolcano: Software for intuitive visualization and interactive exploration of high-throughput biological data. STAR Protocols, 2021, 2, 100279.	0.5	7
119	Photo-dependent membrane-less organelles formed from plant phyB and PIF6 proteins in mammalian cells. International Journal of Biological Macromolecules, 2021, 176, 325-331.	3.6	7
120	Stress-Induced Membraneless Organelles in Eukaryotes and Prokaryotes: Bird's-Eye View. International Journal of Molecular Sciences, 2022, 23, 5010.	1.8	7
121	New Evidence on a Distinction between A β 240 and A β 242 Amyloids: Thioflavin T Binding Modes, Clustering Tendency, Degradation Resistance, and Cross-Seeding. International Journal of Molecular Sciences, 2022, 23, 5513.	1.8	7
122	What causes the depolarization of trypsin and trypsinogen fluorescence. Biophysical Chemistry, 1986, 25, 315-323.	1.5	6
123	ThT as an instrument for testing and investigation of amyloid and amyloid-like fibrils. Proceedings of SPIE, 2007, , .	0.8	6
124	Tryptophan Residue of the D-Galactose/D-Glucose-Binding Protein from E. Coli Localized in its Active Center Does not Contribute to the Change in Intrinsic Fluorescence Upon Glucose Binding. Journal of Fluorescence, 2015, 25, 87-94.	1.3	6
125	Inelastic Incoherent Neutron Scattering in Some Proteins. Ferroelectrics, 2007, 348, 154-160.	0.3	5
126	Determination of homogeneous and inhomogeneous broadenings of quantum-well excitons by 2DFTS: An experiment-theory comparison. Physica Status Solidi C: Current Topics in Solid State Physics, 2009, 6, 445-448.	0.8	5

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127	Structure and stability of recombinant bovine odorant-binding protein: III. Peculiarities of the wild type bOBP unfolding in crowded milieu. PeerJ, 2016, 4, e1642.	0.9	5
128	sfGFP throws light on the early stages of Î²-barrel amyloidogenesis. International Journal of Biological Macromolecules, 2022, 215, 224-234.	3.6	5
129	EFFECTS OF SOME BIOLOGICALLY ACTIVE COMPOUNDS ON PHAGOSOMEâ€“LYSOSOME FUSION IN PERITONEAL MACROPHAGES OF MICE. Cell Biology International, 1998, 22, 465-472.	1.4	4
130	Expression of recombinant GFP-actin fusion protein in the methylotrophic yeast Pichia pastoris. FEMS Yeast Research, 2003, 3, 105-111.	1.1	4
131	Actin and amphiphilic polymers influence on channel formation by Syringomycin E in lipid bilayers. European Biophysics Journal, 2006, 35, 382-392.	1.2	4
132	Comparative assay of amyloid and prion contents in yeast cells. Cell and Tissue Biology, 2008, 2, 71-80.	0.2	4
133	Structure and stability of D-galactose/D-glucose-binding protein. The role of D-glucose binding and Ca ion depletion. Spectroscopy, 2010, 24, 355-359.	0.8	4
134	Denaturation of proteins with beta-barrel topology induced by guanidine hydrochloride. Spectroscopy, 2010, 24, 367-373.	0.8	4
135	Spectral properties of BADAN in solutions with different polarities. Journal of Molecular Structure, 2015, 1090, 107-111.	1.8	4
136	Point mutations affecting yeast prion propagation change the structure of its amyloid fibrils. Journal of Molecular Liquids, 2020, 314, 113618.	2.3	4
137	Structure and stability of recombinant bovine odorant-binding protein: I. Design and analysis of monomeric mutants. PeerJ, 2016, 4, e1933.	0.9	4
138	The Combined Use of Fluorescence Spectroscopy and X-Ray Crystallography Greatly Contributes to Elucidating Structure and Dynamics of Proteins. , 2005, , 25-61.		3
139	Interaction between linker histone H1 and non-histone chromatin protein HMGB1. Spectroscopy, 2010, 24, 165-168.	0.8	3
140	Ligand-Binding Proteins: Structure, Stability and Practical Application. , 0, , .		3
141	Photophysical Properties of Trans-2-[4-(dimethylamino)styryl]-3-ethyl-1,3-benzothiazolium Perchlorate, a New Structural Analog of Thioflavin T. Journal of Applied Spectroscopy, 2014, 81, 205-213.	0.3	3
142	Formation of trans-2-[4-(Dimethylamino)Styryl]-3-Ethyl-1,3-Benzothiazolium Perchlorate Dimers in the Presence of Sodium Polystyrene Sulfonate. Journal of Applied Spectroscopy, 2017, 83, 917-923.	0.3	3
143	The Pathways of the iRFP713 Unfolding Induced by Different Denaturants. International Journal of Molecular Sciences, 2018, 19, 2776.	1.8	3
144	Photophysical Properties of BADAN Revealed in the Study of GGBP Structural Transitions. International Journal of Molecular Sciences, 2021, 22, 11113.	1.8	3

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145	Comparison of crude lysate pellets from isogenic strains of yeast with different prion composition: Identification of prion-associated proteins. <i>Cell and Tissue Biology</i> , 2010, 4, 36-53.	0.2	2
146	Protein-Ligand Interactions of the D-Galactose/D-Glucose-Binding Protein as a Potential Sensing Probe of Glucose Biosensors. <i>Spectroscopy</i> , 2012, 27, 373-379.	0.8	2
147	Circularized Visualisation of Genetic Interactions. , 2017, , .		2
148	Osmolyte-Like Stabilizing Effects of Low GdnHCl Concentrations on d-Glucose/d-Galactose-Binding Protein. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2008.	1.8	2
149	Probing the allostery in dimeric near-infrared biomarkers derived from the bacterial phytochromes: The impact of the T204A substitution on the inter-monomer interaction. <i>International Journal of Biological Macromolecules</i> , 2020, 162, 894-902.	3.6	2
150	Structure and stability of recombinant bovine odorant-binding protein: II. Unfolding of the monomeric forms. <i>PeerJ</i> , 2016, 4, e1574.	0.9	2
151	What causes the variation of polarization degree across the emission spectrum of proteins?. <i>Biophysical Chemistry</i> , 1986, 24, 327-335.	1.5	1
152	High stability of trehalose/maltose binding protein from <i>Thermococcus litoralis</i> makes it a good candidate as a sensitive element in biosensor systems for sugar control. <i>Spectroscopy</i> , 2010, 24, 349-353.	0.8	1
153	Structural Perturbation of Superfolder GFP in the Presence of Guanidine Thiocyanate. <i>Spectroscopy</i> , 2012, 27, 381-386.	0.8	1
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