Jayant B Udgaonkar

List of Publications by Year in descending order

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81839 114418 4,846 112 39 63 citations g-index h-index papers 117 117 117 3276 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	NMR evidence for an early framework intermediate on the folding pathway of ribonuclease A. Nature, 1988, 335, 694-699.	13.7	633
2	Thermodynamics of Denaturation of Barstar: Evidence for Cold Denaturation and Evaluation of the Interaction with Guanidine Hydrochloride. Biochemistry, 1995, 34, 3286-3299.	1.2	221
3	Initial hydrophobic collapse in the folding of barstar. Nature, 1995, 377, 754-757.	13.7	215
4	Understanding the Kinetic Roles of the Inducer Heparin and of Rod-like Protofibrils during Amyloid Fibril Formation by Tau Protein. Journal of Biological Chemistry, 2011, 286, 38948-38959.	1.6	122
5	Direct evidence for a dry molten globule intermediate during the unfolding of a small protein. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 12289-12294.	3.3	121
6	Structure is lost incrementally during the unfolding of barstar. Nature Structural Biology, 2001, 8, 799-804.	9.7	109
7	The Folding Mechanism of Barstar: Evidence for Multiple Pathways and Multiple Intermediates. Journal of Molecular Biology, 1995, 247, 1013-1027.	2.0	107
8	Salt-Induced Modulation of the Pathway of Amyloid Fibril Formation by the Mouse Prion Protein. Biochemistry, 2010, 49, 7615-7624.	1.2	101
9	Multiple Routes and Structural Heterogeneity in Protein Folding. Annual Review of Biophysics, 2008, 37, 489-510.	4.5	91
10	Multiple intermediates and transition states during protein unfolding. Nature Structural Biology, 1997, 4, 1016-1024.	9.7	87
11	Continuous dissolution of structure during the unfolding of a small protein. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 11113-11118.	3.3	84
12	Evidence for Stepwise Formation of Amyloid Fibrils by the Mouse Prion Protein. Journal of Molecular Biology, 2008, 382, 1228-1241.	2.0	75
13	How cooperative are protein folding and unfolding transitions?. Protein Science, 2016, 25, 1924-1941.	3.1	70
14	Mechanism of Formation of Amyloid Protofibrils of Barstar from Soluble Oligomers: Evidence for Multiple Steps and Lateral Association Coupled to Conformational Conversion. Journal of Molecular Biology, 2007, 367, 1186-1204.	2.0	57
15	Evidence for the Existence of a Secondary Pathway for Fibril Growth during the Aggregation of Tau. Journal of Molecular Biology, 2012, 421, 296-314.	2.0	57
16	Mechanism of Formation of a Productive Molten Globule Form of Barstarâ€. Biochemistry, 2002, 41, 1710-1716.	1.2	55
17	Osmolytes Induce Structure in an Early Intermediate on the Folding Pathway of Barstar. Journal of Biological Chemistry, 2004, 279, 40303-40313.	1.6	55
18	Development of the Structural Core and of Conformational Heterogeneity during the Conversion of Oligomers of the Mouse Prion Protein to Worm-like Amyloid Fibrils. Journal of Molecular Biology, 2012, 423, 217-231.	2.0	54

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19	Increasing Stability Reduces Conformational Heterogeneity in a Protein Folding Intermediate Ensemble. Journal of Molecular Biology, 2004, 337, 699-711.	2.0	53
20	Rational Stabilization of Helix 2 of the Prion Protein Prevents Its Misfolding and Oligomerization. Journal of the American Chemical Society, 2014, 136, 16704-16707.	6.6	53
21	Molecular Mechanism of the Misfolding and Oligomerization of the Prion Protein: Current Understanding and Its Implications. Biochemistry, 2015, 54, 4431-4442.	1.2	53
22	Characterization of the Unfolding of Ribonuclease A by a Pulsed Hydrogen Exchange Study: Evidence for Competing Pathways for Unfoldingâ€. Biochemistry, 2002, 41, 2641-2654.	1.2	52
23	Polypeptide chain collapse and protein folding. Archives of Biochemistry and Biophysics, 2013, 531, 24-33.	1.4	52
24	Dependence of the Size of the Initially Collapsed Form During the Refolding of Barstar on Denaturant Concentration: Evidence for a Continuous Transition. Journal of Molecular Biology, 2005, 353, 704-718.	2.0	51
25	Dissecting the Non-specific and Specific Components of the Initial Folding Reaction of Barstar by Multi-site FRET Measurements. Journal of Molecular Biology, 2007, 370, 385-405.	2.0	51
26	Mechanistic Studies Unravel the Complexity Inherent in Tau Aggregation Leading to Alzheimer's Disease and the Tauopathies. Biochemistry, 2013, 52, 4107-4126.	1.2	51
27	Dissection of Conformational Conversion Events during Prion Amyloid Fibril Formation Using Hydrogen Exchange and Mass Spectrometry. Journal of Molecular Biology, 2013, 425, 3510-3521.	2.0	49
28	Quantitative analysis of the kinetics of denaturation and renaturation of barstar in the folding transition zone. Protein Science, 1994, 3, 1409-1417.	3.1	48
29	Unfolding Rates of Barstar Determined in Native and Low Denaturant Conditions Indicate the Presence of Intermediates. Biochemistry, 2002, 41, 1568-1578.	1.2	48
30	Conformational Conversion May Precede or Follow Aggregate Elongation on Alternative Pathways of Amyloid Protofibril Formation. Journal of Molecular Biology, 2009, 385, 1266-1276.	2.0	48
31	Structurally Distinct Amyloid Protofibrils Form on Separate Pathways of Aggregation of a Small Protein. Biochemistry, 2009, 48, 6441-6449.	1.2	47
32	Observation of Multistate Kinetics during the Slow Folding and Unfolding of Barstar. Biochemistry, 1999, 38, 9158-9168.	1.2	46
33	The slow folding reaction of barstar: the core tryptophan region attains tight packing before substantial secondary and tertiary structure formation and final compaction of the polypeptide chain 1 IEdited by C. R. Matthews. Journal of Molecular Biology, 2000, 302, 479-495.	2.0	46
34	Characterization of the Folding and Unfolding Reactions of Single-Chain Monellin:  Evidence for Multiple Intermediates and Competing Pathways. Biochemistry, 2007, 46, 11727-11743.	1.2	46
35	Differential Salt-induced Stabilization of Structure in the Initial Folding Intermediate Ensemble of Barstar. Journal of Molecular Biology, 2002, 324, 331-347.	2.0	43
36	Characterization of Intra-molecular Distances and Site-specific Dynamics in Chemically Unfolded Barstar: Evidence for Denaturant-dependent Non-random Structure. Journal of Molecular Biology, 2006, 359, 174-189.	2.0	42

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37	Exploring the Cooperativity of the Fast Folding Reaction of a Small Protein Using Pulsed Thiol Labeling and Mass Spectrometry. Journal of Biological Chemistry, 2007, 282, 37479-37491.	1.6	42
38	Evidence for Initial Non-specific Polypeptide Chain Collapse During the Refolding of the SH3 Domain of PI3 Kinase. Journal of Molecular Biology, 2010, 403, 430-445.	2.0	42
39	Partially Unfolded Forms of the Prion Protein Populated under Misfolding-promoting Conditions. Journal of Biological Chemistry, 2015, 290, 25227-25240.	1.6	42
40	Dynamics of the Core Tryptophan during the Formation of a Productive Molten Globule Intermediate of Barstar. Biochemistry, 2003, 42, 7986-8000.	1.2	41
41	Folding of Tryptophan Mutants of Barstar:  Evidence for an Initial Hydrophobic Collapse on the Folding Pathway. Biochemistry, 1997, 36, 8602-8610.	1.2	39
42	Unfolding of a Small Protein Proceeds via Dry and Wet Globules and a Solvated Transition State. Biophysical Journal, 2013, 105, 2392-2402.	0.2	39
43	Surface Expansion Is Independent of and Occurs Faster than Core Solvation during the Unfolding of Barstarâ€. Biochemistry, 2003, 42, 1551-1563.	1.2	37
44	Diffusional Barrier in the Unfolding of a Small Protein. Journal of Molecular Biology, 2007, 366, 1016-1028.	2.0	37
45	Mechanism of aggregation and membrane interactions of mammalian prion protein. Biochimica Et Biophysica Acta - Biomembranes, 2018, 1860, 1927-1935.	1.4	37
46	A five-residue motif for the design of domain swapping in proteins. Nature Communications, 2019, 10, 452.	5.8	37
47	Native state dynamics drive the unfolding of the SH3 domain of PI3 kinase at high denaturant concentration. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 20711-20716.	3.3	36
48	Unraveling the Molecular Mechanism of pH-Induced Misfolding and Oligomerization of the Prion Protein. Journal of Molecular Biology, 2016, 428, 1345-1355.	2.0	36
49	pH-Jump-Induced Folding and Unfolding Studies of Barstar:  Evidence for Multiple Folding and Unfolding Pathways. Biochemistry, 2001, 40, 15267-15279.	1.2	35
50	Two structural subdomains of barstar detected by rapid mixing NMR measurement of amide hydrogen exchange., 1998, 30, 295-308.		34
51	Resonance Raman Spectroscopic Measurements Delineate the Structural Changes that Occur during Tau Fibril Formation. Biochemistry, 2014, 53, 6550-6565.	1.2	34
52	Structural Effects of Multiple Pathogenic Mutations Suggest a Model for the Initiation of Misfolding of the Prion Protein. Angewandte Chemie - International Edition, 2015, 54, 7529-7533.	7.2	34
53	Folding subdomains of thioredoxin characterized by native-state hydrogen exchange. Protein Science, 2003, 12, 1719-1731.	3.1	29
54	Protein dynamics control proton transfer from bulk solvent to protein interior: A case study with a green fluorescent protein. Protein Science, 2005, 14, 1787-1799.	3.1	29

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55	Barrierless evolution of structure during the submillisecond refolding reaction of a small protein. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7998-8003.	3.3	29
56	Native and nonnative conformational preferences in the urea-unfolded state of barstar. Protein Science, 2009, 13, 3085-3091.	3.1	28
57	Defining the Pathway of Worm-like Amyloid Fibril Formation by the Mouse Prion Protein by Delineation of the Productive and Unproductive Oligomerization Reactions. Biochemistry, 2011, 50, 1153-1161.	1.2	28
58	Identification of Multiple Folding Pathways of Monellin Using Pulsed Thiol Labeling and Mass Spectrometry. Biochemistry, 2011, 50, 3062-3074.	1.2	28
59	High-Energy Intermediates in Protein Unfolding Characterized by Thiol Labeling under Nativelike Conditions. Biochemistry, 2014, 53, 3608-3620.	1.2	28
60	Microsecond Rearrangements of Hydrophobic Clusters in an Initially Collapsed Globule Prime Structure Formation during the Folding of a Small Protein. Journal of Molecular Biology, 2016, 428, 3102-3117.	2.0	28
61	Modulation of the extent of structural heterogeneity in \hat{l}_{\pm} -synuclein fibrils by the small molecule thioflavin T. Journal of Biological Chemistry, 2017, 292, 16891-16903.	1.6	28
62	Thermodynamic Characterization of the Unfolding of the Prion Protein. Biophysical Journal, 2014, 106, 410-420.	0.2	26
63	Tuning Cooperativity on the Free Energy Landscape of Protein Folding. Biochemistry, 2015, 54, 3431-3441.	1.2	26
64	Salt-Mediated Oligomerization of the Mouse Prion Protein Monitored by Real-Time NMR. Journal of Molecular Biology, 2017, 429, 1852-1872.	2.0	26
65	HX-ESI-MS and Optical Studies of the Unfolding of Thioredoxin Indicate Stabilization of a Partially Unfolded, Aggregation-Competent Intermediate at Low pHâ€. Biochemistry, 2006, 45, 11226-11238.	1.2	25
66	Heterogeneity in Protein Folding and Unfolding Reactions. Chemical Reviews, 2022, 122, 8911-8935.	23.0	25
67	Revealing a Concealed Intermediate that Forms after the Rate-limiting Step of Refolding of the SH3 Domain of PI3 Kinase. Journal of Molecular Biology, 2009, 387, 348-362.	2.0	24
68	Effect of Salt on the Urea-Unfolded Form of Barstar Probed bymValue Measurementsâ€. Biochemistry, 2004, 43, 11393-11402.	1.2	23
69	Critical Evaluation of the Two-State Model Describing the Equilibrium Unfolding of the PI3K SH3 Domain by Time-Resolved Fluorescence Resonance Energy Transfer. Biochemistry, 2013, 52, 9482-9496.	1.2	23
70	Multistage Unfolding of an SH3 Domain: An Initial Urea-Filled Dry Molten Globule Precedes a Wet Molten Globule with Non-Native Structure. Journal of Physical Chemistry B, 2014, 118, 6380-6392.	1.2	23
71	Multiple Kinetic Intermediates Accumulate during the Unfolding of Horse Cytochromecin the Oxidized Stateâ€. Biochemistry, 1998, 37, 9147-9155.	1.2	22
72	Identification and Structural Characterization of the Precursor Conformation of the Prion Protein which Directly Initiates Misfolding and Oligomerization. Journal of Molecular Biology, 2017, 429, 886-899.	2.0	22

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73	Amyloid Fibril Formation by the Chain B Subunit of Monellin Occurs by a Nucleation-Dependent Polymerization Mechanism. Biochemistry, 2014, 53, 1206-1217.	1.2	21
74	Pathogenic Mutations within the Disordered Palindromic Region of the Prion Protein Induce Structure Therein and Accelerate the Formation of Misfolded Oligomers. Journal of Molecular Biology, 2016, 428, 3935-3947.	2.0	21
75	Site-specific time-resolved FRET reveals local variations in the unfolding mechanism in an apparently two-state protein unfolding transition. Physical Chemistry Chemical Physics, 2018, 20, 3216-3232.	1.3	21
76	The Osmolyte TMAO Modulates Protein Folding Cooperativity by Altering Global Protein Stability. Biochemistry, 2018, 57, 5851-5863.	1.2	21
77	Equilibrium Unfolding Studies of Monellin: The Double-Chain Variant Appears To Be More Stable Than the Single-Chain Variant. Biochemistry, 2011, 50, 2434-2444.	1.2	20
78	Rise of the Helix from a Collapsed Globule during the Folding of Monellin. Biochemistry, 2015, 54, 5356-5365.	1.2	20
79	The Pathogenic Mutation T182A Converts the Prion Protein into a Molten Globule-like Conformation Whose Misfolding to Oligomers but Not to Fibrils Is Drastically Accelerated. Biochemistry, 2016, 55, 459-469.	1.2	20
80	The G126V Mutation in the Mouse Prion Protein Hinders Nucleation-Dependent Fibril Formation by Slowing Initial Fibril Growth and by Increasing the Critical Concentration. Biochemistry, 2017, 56, 5931-5942.	1.2	20
81	Structural mechanisms of oligomer and amyloid fibril formation by the prion protein. Chemical Communications, 2018, 54, 6230-6242.	2.2	20
82	Binding-induced folding under unfolding conditions: Switching between induced fit and conformational selection mechanisms. Journal of Biological Chemistry, 2019, 294, 16942-16952.	1.6	19
83	Kinetic Studies of the Folding of Heterodimeric Monellin: Evidence for Switching between Alternative Parallel Pathways. Journal of Molecular Biology, 2012, 420, 235-250.	2.0	18
84	Secondary Structural Change Can Occur Diffusely and Not Modularly during Protein Folding and Unfolding Reactions. Journal of the American Chemical Society, 2016, 138, 5866-5878.	6.6	18
85	Mechanistic and Structural Origins of the Asymmetric Barrier to Prion-like Cross-Seeding between Tau-3R and Tau-4R. Journal of Molecular Biology, 2018, 430, 5304-5312.	2.0	18
86	Monitoring site-specific conformational changes in real-time reveals a misfolding mechanism of the prion protein. ELife, $2019, 8, .$	2.8	18
87	Heterologous expression, purification and characterization of heterodimeric monellin. Protein Expression and Purification, 2011, 76, 248-253.	0.6	17
88	Four-State Folding of a SH3 Domain: Salt-Induced Modulation of the Stabilities of the Intermediates and Native State. Biochemistry, 2012, 51, 4723-4734.	1.2	17
89	Transient Non-Native Burial of a Trp Residue Occurs Initially during the Unfolding of a SH3 Domain. Biochemistry, 2012, 51, 8226-8234.	1.2	16
90	Ruggedness in the Free Energy Landscape Dictates Misfolding of the Prion Protein. Journal of Molecular Biology, 2019, 431, 807-824.	2.0	16

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91	Microsecond sub-domain motions and the folding and misfolding of the mouse prion protein. ELife, 2019, 8, .	2.8	16
92	The Utilization of Competing Unfolding Pathways of Monellin Is Dictated by Enthalpic Barriers. Biochemistry, 2013, 52, 5770-5779.	1.2	15
93	Stopped-flow NMR measurement of hydrogen exchange rates in reduced horse cytochromec under strongly destabilizing conditions. Proteins: Structure, Function and Bioinformatics, 1998, 32, 241-247.	1.5	14
94	Aminoâ€acid composition after loop deletion drives domain swapping. Protein Science, 2017, 26, 1994-2002.	3.1	13
95	Fluoroalcohol-Induced Modulation of the Pathway of Amyloid Protofibril Formation by Barstar. Biochemistry, 2011, 50, 805-819.	1.2	12
96	Difference in Fibril Core Stability between Two Tau Four-Repeat Domain Proteins: A Hydrogen–Deuterium Exchange Coupled to Mass Spectrometry Study. Biochemistry, 2013, 52, 8787-8789.	1.2	11
97	Stepwise Assembly of \hat{l}^2 -Sheet Structure during the Folding of an SH3 Domain Revealed by a Pulsed Hydrogen Exchange Mass Spectrometry Study. Biochemistry, 2017, 56, 3754-3769.	1.2	11
98	Observation of Continuous Contraction and a Metastable Misfolded State during the Collapse and Folding of a Small Protein. Journal of Molecular Biology, 2019, 431, 3814-3826.	2.0	11
99	Modulation of the Extent of Cooperative Structural Change During Protein Folding by Chemical Denaturant. Journal of Physical Chemistry B, 2017, 121, 8263-8275.	1.2	10
100	Mapping Distinct Sequences of Structure Formation Differentiating Multiple Folding Pathways of a Small Protein. Journal of the American Chemical Society, 2021, 143, 1447-1457.	6.6	9
101	Mechanistic approaches to understand the prion-like propagation of aggregates of the human tau protein. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2019, 1867, 922-932.	1.1	8
102	GroEL Can Unfold Late Intermediates Populated on the Folding Pathways of Monellin. Journal of Molecular Biology, 2009, 389, 759-775.	2.0	6
103	Chemical Denaturants Smoothen Ruggedness on the Free Energy Landscape of Protein Folding. Biochemistry, 2017, 56, 4053-4063.	1.2	6
104	A Dry Transition State More Compact Than the Native State Is Stabilized by Non-Native Interactions during the Unfolding of a Small Protein. Biochemistry, 2017, 56, 3699-3703.	1.2	5
105	Destabilization of polar interactions in the prion protein triggers misfolding and oligomerization. Protein Science, 2021, 30, 2258-2271.	3.1	5
106	The Lys 280 → Gln mutation mimicking diseaseâ€linked acetylation of Lys 280 in tau extends the struct core of fibrils and modulates their catalytic properties. Protein Science, 2021, 30, 785-803.	tural 3.1	4
107	Expression and purification of single cysteine-containing mutant variants of the mouse prion protein by oxidative refolding. Protein Expression and Purification, 2017, 140, 1-7.	0.6	3
108	Microsecond Dynamics During the Binding-induced Folding of an Intrinsically Disordered Protein. Journal of Molecular Biology, 2021, 433, 167254.	2.0	3

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109	Resolving Site-Specific Heterogeneity of the Unfolded State under Folding Conditions. Journal of Physical Chemistry Letters, 2021, 12, 3295-3302.	2.1	2
110	Structural Characterization of the Cooperativity of Unfolding of a Heterodimeric Protein using Hydrogen Exchange-Mass Spectrometry. Journal of Molecular Biology, 2021, 433, 167268.	2.0	2
111	Elongation of Fibrils Formed by a Tau Fragment is Inhibited by a Transient Dimeric Intermediate. Journal of Physical Chemistry B, 2022, 126, 3385-3397.	1.2	1
112	Introducing the Mechanical Forces in Biochemistry Special Issue. Biochemistry, 2019, 58, 4655-4656.	1.2	0