

Francesco Antonio Aprile

List of Publications by Year in descending order

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Version: 2024-02-01

51
papers

3,556
citations

257101

24
h-index

197535

49
g-index

58
all docs

58
docs citations

58
times ranked

4684
citing authors

#	ARTICLE	IF	CITATIONS
1	A Chemical Mutagenesis Approach to Insert Post-translational Modifications in Aggregation-Prone Proteins. <i>ACS Chemical Neuroscience</i> , 2022, 13, 1714-1718.	1.7	1
2	Systematic Activity Maturation of a Single-Domain Antibody with Non-canonical Amino Acids through Chemical Mutagenesis. <i>Cell Chemical Biology</i> , 2021, 28, 70-77.e5.	2.5	15
3	Rationally Designed Bicyclic Peptides Prevent the Conversion of A β 242 Assemblies Into Fibrillar Structures. <i>Frontiers in Neuroscience</i> , 2021, 15, 623097.	1.4	6
4	Comparative Studies in the A30P and A53T β -Synuclein <i>C. elegans</i> Strains to Investigate the Molecular Origins of Parkinson's Disease. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 552549.	1.8	12
5	The Diagnostic Potential of Amyloidogenic Proteins. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4128.	1.8	7
6	The binding of the small heat-shock protein β -crystallin to fibrils of β -synuclein is driven by entropic forces. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	15
7	Modulation of amyloid- β aggregation by metal complexes with a dual binding mode and their delivery across the blood-brain barrier using focused ultrasound. <i>Chemical Science</i> , 2021, 12, 9485-9493.	3.7	12
8	Man does not live by intrinsically unstructured proteins alone: The role of structured regions in aggregation. <i>BioEssays</i> , 2021, 43, e2100178.	1.2	3
9	The cellular modifier MOAG/SERF drives amyloid formation through charge complementation. <i>EMBO Journal</i> , 2021, 40, e107568.	3.5	15
10	A rationally designed bicyclic peptide remodels A β 242 aggregation in vitro and reduces its toxicity in a worm model of Alzheimer's disease. <i>Scientific Reports</i> , 2020, 10, 15280.	1.6	15
11	Small-molecule sequestration of amyloid- β as a drug discovery strategy for Alzheimer's disease. <i>Science Advances</i> , 2020, 6, .	4.7	95
12	Rational design of a conformation-specific antibody for the quantification of A β 2 oligomers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 13509-13518.	3.3	61
13	Rationally Designed Antibodies as Research Tools to Study the Structure-Toxicity Relationship of Amyloid- β Oligomers. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4542.	1.8	12
14	Rational Design of Conformation-Specific Antibodies for Tau Oligomers. <i>Biophysical Journal</i> , 2020, 118, 370a-371a.	0.2	1
15	Enhancement of the Anti-Aggregation Activity of a Molecular Chaperone Using a Rationally Designed Post-Translational Modification. <i>ACS Central Science</i> , 2019, 5, 1417-1424.	5.3	18
16	Soluble aggregates present in cerebrospinal fluid change in size and mechanism of toxicity during Alzheimer's disease progression. <i>Acta Neuropathologica Communications</i> , 2019, 7, 120.	2.4	64
17	Different soluble aggregates of A β 242 can give rise to cellular toxicity through different mechanisms. <i>Nature Communications</i> , 2019, 10, 1541.	5.8	140
18	<i>C. elegans</i> expressing D76N β 2-microglobulin: a model for in vivo screening of drug candidates targeting amyloidosis. <i>Scientific Reports</i> , 2019, 9, 19960.	1.6	14

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19	Automated Behavioral Analysis of Large <i>C. elegans</i> Populations Using a Wide Field-of-view Tracking Platform. <i>Journal of Visualized Experiments</i> , 2018, , .	0.2	7
20	Third generation antibody discovery methods: <i>in silico</i> rational design. <i>Chemical Society Reviews</i> , 2018, 47, 9137-9157.	18.7	94
21	O ₂ : TARGETING AMYLOID FORMATION USING RATIONALLY DESIGNED ANTIBODIES. <i>Alzheimer's and Dementia</i> , 2018, 14, P611.	0.4	0
22	Targeting Amyloid Aggregation: An Overview of Strategies and Mechanisms. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2677.	1.8	103
23	Cooperative Assembly of Hsp70 Subdomain Clusters. <i>Biochemistry</i> , 2018, 57, 3641-3649.	1.2	13
24	Multistep Inhibition of α -Synuclein Aggregation and Toxicity <i>in Vitro</i> and <i>in Vivo</i> by Trodusquemine. <i>ACS Chemical Biology</i> , 2018, 13, 2308-2319.	1.6	86
25	A Rationally Designed Hsp70 Variant Rescues the Aggregation-Associated Toxicity of Human IAPP in Cultured Pancreatic Islet β -Cells. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1443.	1.8	14
26	A natural product inhibits the initiation of α -synuclein aggregation and suppresses its toxicity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E1009-E1017.	3.3	231
27	Inhibition of α -Synuclein Fibril Elongation by Hsp70 Is Governed by a Kinetic Binding Competition between α -Synuclein Species. <i>Biochemistry</i> , 2017, 56, 1177-1180.	1.2	47
28	Methods of probing the interactions between small molecules and disordered proteins. <i>Cellular and Molecular Life Sciences</i> , 2017, 74, 3225-3243.	2.4	56
29	Selective targeting of primary and secondary nucleation pathways in $A\beta^{242}$ aggregation using a rational antibody scanning method. <i>Science Advances</i> , 2017, 3, e1700488.	4.7	116
30	Identification of an RNA Polymerase III Regulator Linked to Disease-Associated Protein Aggregation. <i>Molecular Cell</i> , 2017, 65, 1096-1108.e6.	4.5	14
31	The polyglutamine protein ataxin-3 enables normal growth under heat shock conditions in the methylotrophic yeast <i>Pichia pastoris</i> . <i>Scientific Reports</i> , 2017, 7, 13417.	1.6	0
32	A Water-Bridged Cysteine-Cysteine Redox Regulation Mechanism in Bacterial Protein Tyrosine Phosphatases. <i>CheM</i> , 2017, 3, 665-677.	5.8	18
33	The molecular chaperones DNAJB6 and Hsp70 cooperate to suppress α -synuclein aggregation. <i>Scientific Reports</i> , 2017, 7, 9039.	1.6	67
34	Sequence Specificity in the Entropy-Driven Binding of a Small Molecule and a Disordered Peptide. <i>Journal of Molecular Biology</i> , 2017, 429, 2772-2779.	2.0	62
35	Delivery of Native Proteins into <i>C. elegans</i> Using a Transduction Protocol Based on Lipid Vesicles. <i>Scientific Reports</i> , 2017, 7, 15045.	1.6	16
36	Structure of a low-population binding intermediate in protein-RNA recognition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7171-7176.	3.3	54

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37	Microfluidic Diffusion Viscometer for Rapid Analysis of Complex Solutions. <i>Analytical Chemistry</i> , 2016, 88, 3488-3493.	3.2	29
38	Microfluidic Diffusion Analysis of the Sizes and Interactions of Proteins under Native Solution Conditions. <i>ACS Nano</i> , 2016, 10, 333-341.	7.3	105
39	Rational design of antibodies targeting specific epitopes within intrinsically disordered proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 9902-9907.	3.3	113
40	The CamSol Method of Rational Design of Protein Mutants with Enhanced Solubility. <i>Journal of Molecular Biology</i> , 2015, 427, 478-490.	2.0	341
41	Structure and Dynamics of the Integrin LFA-1 I-Domain in the Inactive State Underlie its Inside-Out/Outside-In Signaling and Allosteric Mechanisms. <i>Structure</i> , 2015, 23, 745-753.	1.6	15
42	Structural characterization of toxic oligomers that are kinetically trapped during α -synuclein fibril formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E1994-2003.	3.3	384
43	Biophysical approaches for the study of interactions between molecular chaperones and protein aggregates. <i>Chemical Communications</i> , 2015, 51, 14425-14434.	2.2	18
44	A Rational Design Strategy for the Selective Activity Enhancement of a Molecular Chaperone toward a Target Substrate. <i>Biochemistry</i> , 2015, 54, 5103-5112.	1.2	25
45	Cell surface localised Hsp70 is a cancer specific regulator of clathrin-independent endocytosis. <i>FEBS Letters</i> , 2015, 589, 2747-2753.	1.3	37
46	Structure of a low-population intermediate state in the release of an enzyme product. <i>ELife</i> , 2015, 4, .	2.8	33
47	NMR characterization of the conformational fluctuations of the human lymphocyte function-associated antigen-1 α domain. <i>Protein Science</i> , 2014, 23, 1596-1606.	3.1	8
48	Nanobodies Raised against Monomeric α -Synuclein Distinguish between Fibrils at Different Maturation Stages. <i>Journal of Molecular Biology</i> , 2013, 425, 2397-2411.	2.0	90
49	Hsp70 Oligomerization Is Mediated by an Interaction between the Interdomain Linker and the Substrate-Binding Domain. <i>PLoS ONE</i> , 2013, 8, e67961.	1.1	66
50	Direct Observation of the Interconversion of Normal and Toxic Forms of α -Synuclein. <i>Cell</i> , 2012, 149, 1048-1059.	13.5	755
51	The Relationship between Aggregation and Toxicity of Polyglutamine-Containing Ataxin-3 in the Intracellular Environment of <i>Escherichia coli</i> . <i>PLoS ONE</i> , 2012, 7, e51890.	1.1	20