

# Filippo Pullara

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3488711/publications.pdf>

Version: 2024-02-01

12  
papers

508  
citations

1478505

6  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

926  
citing authors

#	ARTICLE	IF	CITATIONS
1	Aggregation kinetics of bovine serum albumin studied by FTIR spectroscopy and light scattering. <i>Biophysical Chemistry</i> , 2004, 107, 175-187.	2.8	266
2	Crystal Structure of a Transcribing RNA Polymerase II Complex Reveals a Complete Transcription Bubble. <i>Molecular Cell</i> , 2015, 59, 258-269.	9.7	98
3	Explainable AI (xAI) for Anatomic Pathology. <i>Advances in Anatomic Pathology</i> , 2020, 27, 241-250.	4.3	46
4	WNT5A Inhibits Hepatocyte Proliferation and Concludes $\beta$ -Catenin Signaling in Liver Regeneration. <i>American Journal of Pathology</i> , 2015, 185, 2194-2205.	3.8	29
5	Spatial domain analysis predicts risk of colorectal cancer recurrence and infers associated tumor microenvironment networks. <i>Nature Communications</i> , 2020, 11, 3515.	12.8	24
6	A general path for large-scale solubilization of cellular proteins: From membrane receptors to multiprotein complexes. <i>Protein Expression and Purification</i> , 2013, 87, 111-119.	1.3	17
7	Why protein conformers in molecular dynamics simulations differ from their crystal structures: a thermodynamic insight. <i>Turkish Journal of Chemistry</i> , 2019, 43, 394-403.	1.2	9
8	Early stages of $\alpha$ -microglobulin aggregation and the inhibiting action of $\beta$ -crystallin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 73, 1037-1046.	2.6	5
9	Mechanisms of Activation and Subunit Release in $Ca^{2+}$ /Calmodulin-Dependent Protein Kinase II. <i>Journal of Physical Chemistry B</i> , 2017, 121, 10344-10352.	2.6	5
10	Network Proteomics: From Protein Structure to Protein-Protein Interaction. <i>BioMed Research International</i> , 2017, 2017, 1-1.	1.9	4
11	In situ functional cell phenotyping reveals microdomain networks in colorectal cancer recurrence. <i>Cell Reports Methods</i> , 2021, 1, 100072.	2.9	3
12	Population reversal driven by unrestrained interactions in molecular dynamics simulations: A dialanine model. <i>AIP Advances</i> , 2015, 5, .	1.3	2