

# Giulia Culetta

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2186019/publications.pdf>

Version: 2024-02-01

8  
papers

113  
citations

1478505

6  
h-index

1588992

8  
g-index

8  
all docs

8  
docs citations

8  
times ranked

139  
citing authors

#	ARTICLE	IF	CITATIONS
1	Exploring the SARS-CoV-2 Proteome in the Search of Potential Inhibitors via Structure-Based Pharmacophore Modeling/Docking Approach. <i>Computation</i> , 2020, 8, 77.	2.0	30
2	Evaluation of the IKK $\beta$ Binding of Indicaxanthin by Induced-Fit Docking, Binding Pose Metadynamics, and Molecular Dynamics. <i>Frontiers in Pharmacology</i> , 2021, 12, 701568.	3.5	24
3	Indicaxanthin, a multi-target natural compound from <i>Opuntia ficus-indica</i> fruit: From its poly-pharmacological effects to biochemical mechanisms and molecular modelling studies. <i>European Journal of Medicinal Chemistry</i> , 2019, 179, 753-764.	5.5	22
4	Pharmacophore-Based Design of New Chemical Scaffolds as Translational Readthrough-Inducing Drugs (TRIDs). <i>ACS Medicinal Chemistry Letters</i> , 2020, 11, 747-753.	2.8	13
5	In Silico Design, Synthesis, and Biological Evaluation of Anticancer Arylsulfonamide Endowed with Anti-Telomerase Activity. <i>Pharmaceuticals</i> , 2022, 15, 82.	3.8	11
6	Comparing molecular dynamics-derived pharmacophore models with docking: A study on CDK-2 inhibitors. <i>Chemical Data Collections</i> , 2020, 28, 100485.	2.3	8
7	Immunoproteasome and Non-Covalent Inhibition: Exploration by Advanced Molecular Dynamics and Docking Methods. <i>Molecules</i> , 2021, 26, 4046.	3.8	3
8	A Definitive Pharmacophore Modelling Study on CDK2 ATP Pocket Binders: Tracing the Path of New Virtual High-Throughput Screenings. <i>Current Drug Discovery Technologies</i> , 2020, 17, 740-747.	1.2	2