

# Simon Cross

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6576110/publications.pdf>

Version: 2024-02-01

9  
papers

340  
citations

1478280

6  
h-index

1474057

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

611  
citing authors

#	ARTICLE	IF	CITATIONS
1	FragExplorer: GRID-Based Fragment Growing and Replacement. <i>Journal of Chemical Information and Modeling</i> , 2022, 62, 1224-1235.	2.5	4
2	Combining machine learning and quantum mechanics yields more <scp>chemically aware</scp> molecular descriptors for medicinal chemistry applications. <i>Journal of Computational Chemistry</i> , 2021, 42, 2068-2078.	1.5	6
3	Playing with Opening and Closing of Heterocycles: Using the Cusmano-Ruccia Reaction to Develop a Novel Class of Oxadiazolothiazinones, Active as Calcium Channel Modulators and P-Glycoprotein Inhibitors. <i>Molecules</i> , 2014, 19, 16543-16572.	1.7	6
4	Molecular interaction fields in drug discovery: recent advances and future perspectives. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2013, 3, 594-613.	6.2	36
5	GRID-Based Three-Dimensional Pharmacophores II: PharmBench, a Benchmark Data Set for Evaluating Pharmacophore Elucidation Methods. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 2599-2608.	2.5	36
6	GRID-Based Three-Dimensional Pharmacophores I: FLAPpharm, a Novel Approach for Pharmacophore Elucidation. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 2587-2598.	2.5	76
7	Molecular fields in drug discovery: getting old or reaching maturity?. <i>Drug Discovery Today</i> , 2010, 15, 23-32.	3.2	67
8	FLAP: GRID Molecular Interaction Fields in Virtual Screening. Validation using the DUD Data Set. <i>Journal of Chemical Information and Modeling</i> , 2010, 50, 1442-1450.	2.5	94
9	Grid-derived structure-based 3D pharmacophores and their performance compared to docking. <i>Drug Discovery Today: Technologies</i> , 2010, 7, e213-e219.	4.0	15