

# Janosch Achenbach

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10650747/publications.pdf>

Version: 2024-02-01

15  
papers

366  
citations

933447

10  
h-index

996975

15  
g-index

15  
all docs

15  
docs citations

15  
times ranked

648  
citing authors

#	ARTICLE	IF	CITATIONS
1	Computational tools for polypharmacology and repurposing. <i>Future Medicinal Chemistry</i> , 2011, 3, 961-968.	2.3	66
2	Dual-Target Virtual Screening by Pharmacophore Elucidation and Molecular Shape Filtering. <i>ACS Medicinal Chemistry Letters</i> , 2012, 3, 155-158.	2.8	43
3	VAMMPIRE: A Matched Molecular Pairs Database for Structure-Based Drug Design and Optimization. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 5203-5207.	6.4	43
4	Investigation of imatinib and other approved drugs as starting points for antidiabetic drug discovery with FXR modulating activity. <i>Biochemical Pharmacology</i> , 2012, 83, 1674-1681.	4.4	36
5	DrugBank screening revealed alitretinoin and bexarotene as liver X receptor modulators. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2017, 27, 1193-1198.	2.2	36
6	Exploring the Chemical Space of Multitarget Ligands Using Aligned Self-Organizing Maps. <i>ACS Medicinal Chemistry Letters</i> , 2013, 4, 1169-1172.	2.8	33
7	Multi-dimensional target profiling of N,4-diaryl-1,3-thiazole-2-amines as potent inhibitors of eicosanoid metabolism. <i>European Journal of Medicinal Chemistry</i> , 2014, 84, 302-311.	5.5	29
8	Evaluation of structure-derived pharmacophore of soluble epoxide hydrolase inhibitors by virtual screening. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2012, 22, 6762-6765.	2.2	22
9	Synthetic cannabinoids: In silico prediction of the cannabinoid receptor 1 affinity by a quantitative structure-activity relationship model. <i>Toxicology Letters</i> , 2016, 245, 1-6.	0.8	18
10	VAMMPIRE-LORD: A Web Server for Straightforward Lead Optimization Using Matched Molecular Pairs. <i>Journal of Chemical Information and Modeling</i> , 2015, 55, 207-213.	5.4	14
11	Identification of novel farnesoid X receptor modulators using a combined ligand- and structure-based virtual screening. <i>MedChemComm</i> , 2013, 4, 920.	3.4	10
12	Computer-Aided Selective Optimization of Side Activities of Talinolol. <i>ACS Medicinal Chemistry Letters</i> , 2019, 10, 899-903.	2.8	9
13	Pyrazolo[1,5a]pyrimidines as a new class of FUSE binding protein 1 (FUBP1) inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2016, 24, 5717-5729.	3.0	5
14	Fragment-based identification of multi-target ligands by self-organizing map alignment. <i>Journal of Cheminformatics</i> , 2012, 4, .	6.1	1
15	11th German Conference on Chemoinformatics (GCC 2015). <i>Journal of Cheminformatics</i> , 2016, 8, 18.	6.1	1