

# Bernd Rupp

## List of Publications by Year in descending order

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11  
papers

268  
citations

1040056

9  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

651  
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural Characterization and Ligand/Inhibitor Identification Provide Functional Insights into the Mycobacterium tuberculosis Cytochrome P450 CYP126A1. <i>Journal of Biological Chemistry</i> , 2017, 292, 1310-1329.	3.4	13
2	11th German Conference on Chemoinformatics (GCC 2015). <i>Journal of Cheminformatics</i> , 2016, 8, 18.	6.1	1
3	Design of a Generalâ€Purpose European Compound Screening Library for EUâ€OPENSREEN. <i>ChemMedChem</i> , 2014, 9, 2309-2326.	3.2	29
4	DACS: from compound collections to rationally designed HTS library. <i>Journal of Cheminformatics</i> , 2014, 6, .	6.1	0
5	Substrate Transport Activation Is Mediated through Second Periplasmic Loop of Transmembrane Protein MalF in Maltose Transport Complex of Escherichia coli. <i>Journal of Biological Chemistry</i> , 2012, 287, 17040-17049.	3.4	16
6	Characterization of Structural Features Controlling the Receptiveness of Empty Class II MHC Molecules. <i>PLoS ONE</i> , 2011, 6, e18662.	2.5	31
7	Chronic oral LOEL prediction by using a commercially available computational QSAR tool. <i>Archives of Toxicology</i> , 2010, 84, 681-688.	4.2	17
8	Design of chemical libraries with potentially bioactive molecules applying a maximum common substructure concept. <i>Molecular Diversity</i> , 2010, 14, 401-408.	3.9	69
9	Chemoenzymatic Synthesis of a Glycolipid Library and Elucidation of the Antigenic Epitope for Construction of a Vaccine Against Lyme Disease. <i>Chemistry - A European Journal</i> , 2010, 16, 3536-3544.	3.3	20
10	Anchor Side Chains of Short Peptide Fragments Trigger Ligand-Exchange of Class II MHC Molecules. <i>PLoS ONE</i> , 2008, 3, e1814.	2.5	34
11	Molecular design of two sterol 14Î±-demethylase homology models and their interactions with the azole antifungals ketoconazole and bifonazole. <i>Journal of Computer-Aided Molecular Design</i> , 2005, 19, 149-163.	2.9	38