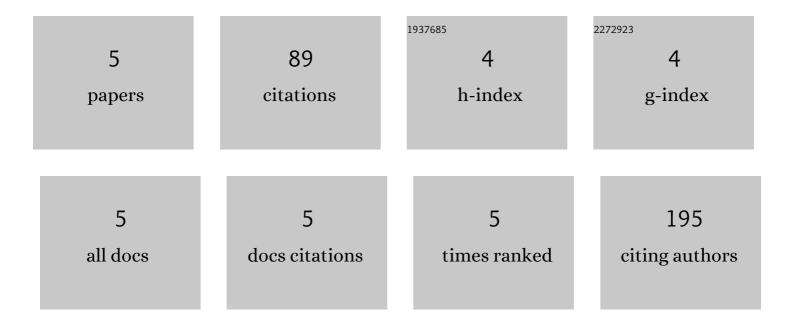
Tobias Depke

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6029703/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Clustering of MS2 spectra using unsupervised methods to aid the identification of secondary metabolites from Pseudomonas aeruginosa. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1071, 19-28.	2.3	37
2	Untargeted LC-MS Metabolomics Differentiates Between Virulent and Avirulent Clinical Strains of Pseudomonas aeruginosa. Biomolecules, 2020, 10, 1041.	4.0	21
3	The Alkylquinolone Repertoire of Pseudomonas aeruginosa is Linked to Structural Flexibility of the FabHâ€like 2â€Heptylâ€3â€hydroxyâ€4(1 H)â€quinolone (PQS) Biosynthesis Enzyme PqsBC. ChemBioChem, 20 1531-1544.)18,19,	17
4	CluMSID: an R package for similarity-based clustering of tandem mass spectra to aid feature annotation in metabolomics. Bioinformatics, 2019, 35, 3196-3198.	4.1	14
5	The Peptide Chain Release Factor Methyltransferase PrmC Influences the Pseudomonas aeruginosa PA14 Endo- and Exometabolome. Metabolites, 2020, 10, 417.	2.9	0