

# Garwin Pichler

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/524456/publications.pdf>

Version: 2024-02-01

14  
papers

3,744  
citations

840119

11  
h-index

1199166

12  
g-index

15  
all docs

15  
docs citations

15  
times ranked

8116  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomics Analysis of Monocyte-Derived Hepatocyte-Like Cells Identifies Integrin Beta 3 as a Specific Biomarker for Drug-Induced Liver Injury by Diclofenac. <i>Frontiers in Pharmacology</i> , 2018, 9, 699.	1.6	23
2	Proteomik für die klinische Diagnostik. <i>Nachrichten Aus Der Chemie</i> , 2016, 64, 540-541.	0.0	0
3	Ultra-deep and quantitative saliva proteome reveals dynamics of the oral microbiome. <i>Genome Medicine</i> , 2016, 8, 44.	3.6	170
4	Circular non-coding RNA ANRIL modulates ribosomal RNA maturation and atherosclerosis in humans. <i>Nature Communications</i> , 2016, 7, 12429.	5.8	859
5	Plasma Proteome Profiling to Assess Human Health and Disease. <i>Cell Systems</i> , 2016, 2, 185-195.	2.9	549
6	Versatile and Efficient Site-Specific Protein Functionalization by Tubulin Tyrosine Ligase. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 13787-13791.	7.2	82
7	DNA methylation requires a DNMT1 ubiquitin interacting motif (UIM) and histone ubiquitination. <i>Cell Research</i> , 2015, 25, 911-929.	5.7	201
8	Minimal, encapsulated proteomic-sample processing applied to copy-number estimation in eukaryotic cells. <i>Nature Methods</i> , 2014, 11, 319-324.	9.0	1,447
9	Establishment of a robust single axis of cell polarity by coupling multiple positive feedback loops. <i>Nature Communications</i> , 2013, 4, 1807.	5.8	99
10	Fluorescent Protein Specific Nanotraps to Study Protein-Protein Interactions and Histone-Tail Peptide Binding. , 2012, 911, 475-483.		12
11	Versatile Toolbox for High Throughput Biochemical and Functional Studies with Fluorescent Fusion Proteins. <i>PLoS ONE</i> , 2012, 7, e36967.	1.1	9
12	Cooperative DNA and histone binding by Uhrf2 links the two major repressive epigenetic pathways. <i>Journal of Cellular Biochemistry</i> , 2011, 112, 2585-2593.	1.2	62
13	Regulation of DNA methyltransferase 1 by interactions and modifications. <i>Nucleus</i> , 2011, 2, 392-402.	0.6	86
14	The multi-domain protein Np95 connects DNA methylation and histone modification. <i>Nucleic Acids Research</i> , 2010, 38, 1796-1804.	6.5	139