

# Stefan Wilkening

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11722665/publications.pdf>

Version: 2024-02-01

20  
papers

1,470  
citations

567281

15  
h-index

752698

20  
g-index

20  
all docs

20  
docs citations

20  
times ranked

2728  
citing authors

#	ARTICLE	IF	CITATIONS
1	COMPARISON OF PRIMARY HUMAN HEPATOCYTES AND HEPATOMA CELL LINE HEPG2 WITH REGARD TO THEIR BIOTRANSFORMATION PROPERTIES. <i>Drug Metabolism and Disposition</i> , 2003, 31, 1035-1042.	3.3	662
2	MDM2 SNP309 and cancer risk: a combined analysis. <i>Carcinogenesis</i> , 2007, 28, 2262-2267.	2.8	98
3	An efficient method for genome-wide polyadenylation site mapping and RNA quantification. <i>Nucleic Acids Research</i> , 2013, 41, e65-e65.	14.5	98
4	Influence of culture time on the expression of drug-metabolizing enzymes in primary human hepatocytes and hepatoma cell line HepG2. <i>Journal of Biochemical and Molecular Toxicology</i> , 2003, 17, 207-213.	3.0	92
5	Alternative polyadenylation diversifies post-transcriptional regulation by selective <i>scRNA</i> protein interactions. <i>Molecular Systems Biology</i> , 2014, 10, 719.	7.2	91
6	An Evaluation of High-Throughput Approaches to QTL Mapping in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2014, 196, 853-865.	2.9	86
7	Interleukin promoter polymorphisms and prognosis in colorectal cancer. <i>Carcinogenesis</i> , 2008, 29, 1202-1206.	2.8	63
8	Genotyping 1000 yeast strains by next-generation sequencing. <i>BMC Genomics</i> , 2013, 14, 90.	2.8	47
9	Determination of allele frequency in pooled DNA: comparison of three PCR-based methods. <i>BioTechniques</i> , 2005, 39, 853-858.	1.8	41
10	Quantitative real-time polymerase chain reaction: methodical analysis and mathematical model. <i>Journal of Biomolecular Techniques</i> , 2004, 15, 107-111.	1.5	35
11	The Single Nucleotide Polymorphism IVS1+309 in <i>Mouse Double Minute 2</i> Does Not Affect Risk of Familial Breast Cancer. <i>Cancer Research</i> , 2006, 66, 646-648.	0.9	33
12	Negative feedback buffers effects of regulatory variants. <i>Molecular Systems Biology</i> , 2015, 11, 785.	7.2	33
13	Candidate Genetic Modifiers for Breast and Ovarian Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 308-316.	2.5	22
14	Genome-Wide Polyadenylation Site Mapping. <i>Methods in Enzymology</i> , 2012, 513, 271-296.	1.0	21
15	Allelotyping of pooled DNA with 250 K SNP microarrays. <i>BMC Genomics</i> , 2007, 8, 77.	2.8	16
16	Induced Mutations in Yeast Cell Populations Adapting to an Unforeseen Challenge. <i>PLoS ONE</i> , 2014, 9, e111133.	2.5	10
17	Differential Regulation of CYP3A4 and CYP3A7 by Dimethylsulfoxide in Primary Human Hepatocytes. <i>Basic and Clinical Pharmacology and Toxicology</i> , 2004, 95, 92-93.	0.0	9
18	STR Markers for Kinship Analysis. <i>Human Biology</i> , 2006, 78, 1-8.	0.2	7

#	ARTICLE	IF	CITATIONS
19	Polyglutamine repeat length in the NCOA3 does not affect risk in familial breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 291-2.	2.5	4
20	Genome-Wide Identification of Alternative Polyadenylation Events Using 3'UT-Fill. <i>Methods in Molecular Biology</i> , 2016, 1358, 295-302.	0.9	2