

# Wolfgang Kaisers

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3316559/publications.pdf>

Version: 2024-02-01

11  
papers

195  
citations

1163117

8  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

452  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of amino acid determinants in CYP4B1 for optimal catalytic processing of 4-ipomeanol. <i>Biochemical Journal</i> , 2015, 465, 103-114.	3.7	46
2	Analysis of Documentation Speed Using Web-Based Medical Speech Recognition Technology: Randomized Controlled Trial. <i>Journal of Medical Internet Research</i> , 2015, 17, e247.	4.3	30
3	Age, gender and UV-exposition related effects on gene expression in in vivo aged short term cultivated human dermal fibroblasts. <i>PLoS ONE</i> , 2017, 12, e0175657.	2.5	29
4	Ranking noncanonical 5â€™ splice site usage by genome-wide RNA-seq analysis and splicing reporter assays. <i>Genome Research</i> , 2018, 28, 1826-1840.	5.5	22
5	Biomechanical assessment of healthy and keratoconic corneas (with/without crosslinking) using dynamic ultrahigh-speed Scheimpflug technology and the relevance of the parameter (A1L~A2L). <i>British Journal of Ophthalmology</i> , 2019, 103, 558-564.	3.9	18
6	Succession of splicing regulatory elements determines cryptic 5â€™ss functionality. <i>Nucleic Acids Research</i> , 2017, 45, gkw1317.	14.5	16
7	rbamtools: an R interface to samtools enabling fast accumulative tabulation of splicing events over multiple RNA-seq samples. <i>Bioinformatics</i> , 2015, 31, 1663-1664.	4.1	12
8	High Concentration of Low-Density Lipoprotein Results in Disturbances in Mitochondrial Transcription and Functionality in Endothelial Cells. <i>Oxidative Medicine and Cellular Longevity</i> , 2019, 2019, 1-12.	4.0	8
9	Validation of Splicing Events in Transcriptome Sequencing Data. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1110.	4.1	6
10	Hierarchical Clustering of DNA k-mer Counts in RNAseq Fastq Files Identifies Sample Heterogeneities. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3687.	4.1	5
11	Sample Size Estimation for Detection of Splicing Events in Transcriptome Sequencing Data. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1900.	4.1	3