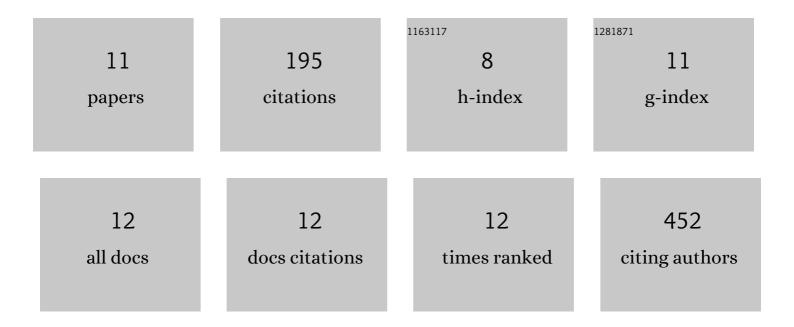
Wolfgang Kaisers

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification of amino acid determinants in CYP4B1 for optimal catalytic processing of 4-ipomeanol. Biochemical Journal, 2015, 465, 103-114.	3.7	46
2	Analysis of Documentation Speed Using Web-Based Medical Speech Recognition Technology: Randomized Controlled Trial. Journal of Medical Internet Research, 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. PLoS ONE, 2017, 12, e0175657.	2.5	29
4	Ranking noncanonical 5′ splice site usage by genome-wide RNA-seq analysis and splicing reporter assays. Genome Research, 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 (A1Lâ^'A2L). British Journal of Ophthalmology, 2019, 103, 558-564.	3.9	18
6	Succession of splicing regulatory elements determines cryptic 5′ss functionality. Nucleic Acids Research, 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. Bioinformatics, 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. Oxidative Medicine and Cellular Longevity, 2019, 2019, 1-12.	4.0	8
9	Validation of Splicing Events in Transcriptome Sequencing Data. International Journal of Molecular Sciences, 2017, 18, 1110.	4.1	6
10	Hierarchical Clustering of DNA k-mer Counts in RNAseq Fastq Files Identifies Sample Heterogeneities. International Journal of Molecular Sciences, 2018, 19, 3687.	4.1	5
11	Sample Size Estimation for Detection of Splicing Events in Transcriptome Sequencing Data. International Journal of Molecular Sciences, 2017, 18, 1900.	4.1	3