## **Koos Rooijers**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2592046/publications.pdf

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687220 996849 1,922 16 13 15 citations h-index g-index papers 16 16 16 4240 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Alcohol-derived DNA crosslinks are repaired by two distinct mechanisms. Nature, 2020, 579, 603-608.	13.7	82
2	Simultaneous quantification of protein–DNA interactions and transcriptomes in single cells with scDam&T-seq. Nature Protocols, 2020, 15, 1922-1953.	5.5	25
3	Simultaneous quantification of protein–DNA contacts and transcriptomes in single cells. Nature Biotechnology, 2019, 37, 766-772.	9.4	86
4	<scp>TGF</scp> β1â€induced leucine limitation uncovered by differential ribosome codon reading. EMBO Reports, 2017, 18, 549-557.	2.0	8
5	Using mitoribosomal profiling to investigate human mitochondrial translation. Wellcome Open Research, 2017, 2, 116.	0.9	4
6	Tumour-specific proline vulnerability uncovered by differential ribosome codon reading. Nature, 2016, 530, 490-494.	13.7	202
7	Parasiteâ€induced <scp>ER</scp> stress response in hepatocytes facilitates <i>Plasmodium</i> liver stage infection. EMBO Reports, 2015, 16, 955-964.	2.0	46
8	Exposure to arginine analog canavanine induces aberrant mitochondrial translation products, mitoribosome stalling, and instability of the mitochondrial proteome. International Journal of Biochemistry and Cell Biology, 2015, 65, 268-274.	1,2	16
9	Genome-wide profiling of p53-regulated enhancer RNAs uncovers a subset of enhancers controlled by a lncRNA. Nature Communications, 2015, 6, 6520.	5.8	149
10	Human bone marrow- and adipose-mesenchymal stem cells secrete exosomes enriched in distinctive miRNA and tRNA species. Stem Cell Research and Therapy, 2015, 6, 127.	2.4	599
11	p53 induces transcriptional and translational programs to suppress cell proliferation and growth. Genome Biology, 2013, 14, R32.	13.9	97
12	Ribosome profiling reveals features of normal and disease-associated mitochondrial translation. Nature Communications, 2013, 4, 2886.	5.8	87
13	Alternative Cleavage and Polyadenylation during Colorectal Cancer Development. Clinical Cancer Research, 2012, 18, 5256-5266.	3.2	108
14	The Poly(A)-Binding Protein Nuclear 1 Suppresses Alternative Cleavage and Polyadenylation Sites. Cell, 2012, 149, 538-553.	13.5	309
15	An iterative workflow for mining the human intestinal metaproteome. BMC Genomics, 2011, 12, 6.	1.2	93
16	Using mitoribosomal profiling to investigate human mitochondrial translation. Wellcome Open Research, 0, 2, 116.	0.9	11