

# Niek de Klein

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9403111/publications.pdf>

Version: 2024-02-01

13  
papers

896  
citations

1040056

9  
h-index

1125743

13  
g-index

20  
all docs

20  
docs citations

20  
times ranked

3699  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of context-dependent expression quantitative trait loci in whole blood. <i>Nature Genetics</i> , 2017, 49, 139-145.	21.4	363
2	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. <i>Nature Genetics</i> , 2021, 53, 1636-1648.	21.4	223
3	Association of maternal prenatal smoking GFI1-locus and cardio-metabolic phenotypes in 18,212 adults. <i>EBioMedicine</i> , 2018, 38, 206-216.	6.1	43
4	TMEM258 Is a Component of the Oligosaccharyltransferase Complex Controlling ER Stress and Intestinal Inflammation. <i>Cell Reports</i> , 2016, 17, 2955-2965.	6.4	42
5	Deconvolution of bulk blood eQTL effects into immune cell subpopulations. <i>BMC Bioinformatics</i> , 2020, 21, 243.	2.6	38
6	Fisetin protects against cardiac cell death through reduction of ROS production and caspases activity. <i>Scientific Reports</i> , 2020, 10, 2896.	3.3	37
7	Epigenome-wide association meta-analysis of DNA methylation with coffee and tea consumption. <i>Nature Communications</i> , 2021, 12, 2830.	12.8	35
8	Meta-analyses identify DNA methylation associated with kidney function and damage. <i>Nature Communications</i> , 2021, 12, 7174.	12.8	30
9	A Comprehensive Analysis of MicroProteins Reveals Their Potentially Widespread Mechanism of Transcriptional Regulation. <i>Plant Physiology</i> , 2014, 165, 149-159.	4.8	21
10	microProtein Prediction Program (miP3): A Software for Predicting microProteins and Their Target Transcription Factors. <i>International Journal of Genomics</i> , 2015, 2015, 1-4.	1.6	6
11	The genetic architecture of molecular traits. <i>Current Opinion in Systems Biology</i> , 2017, 1, 25-31.	2.6	6
12	Potential impact of celiac disease genetic risk factors on T cell receptor signaling in gluten-specific CD4+ T cells. <i>Scientific Reports</i> , 2021, 11, 9252.	3.3	6
13	FC 011KIDNEYNETWORK: USING KIDNEY DERIVED GENE EXPRESSION DATA TO PREDICT AND PRIORITIZE NOVEL GENES INVOLVED IN KIDNEY DISEASE. <i>Nephrology Dialysis Transplantation</i> , 2021, 36, .	0.7	0