

Niek de Klein

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

896
citations

1040056

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3699
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| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | 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 |
| 2 | Epigenome-wide association meta-analysis of DNA methylation with coffee and tea consumption. <i>Nature Communications</i> , 2021, 12, 2830. | 12.8 | 35 |
| 3 | 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 |
| 4 | 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 |
| 5 | Meta-analyses identify DNA methylation associated with kidney function and damage. <i>Nature Communications</i> , 2021, 12, 7174. | 12.8 | 30 |
| 6 | Deconvolution of bulk blood eQTL effects into immune cell subpopulations. <i>BMC Bioinformatics</i> , 2020, 21, 243. | 2.6 | 38 |
| 7 | Fisetin protects against cardiac cell death through reduction of ROS production and caspases activity. <i>Scientific Reports</i> , 2020, 10, 2896. | 3.3 | 37 |
| 8 | 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 |
| 9 | Identification of context-dependent expression quantitative trait loci in whole blood. <i>Nature Genetics</i> , 2017, 49, 139-145. | 21.4 | 363 |
| 10 | The genetic architecture of molecular traits. <i>Current Opinion in Systems Biology</i> , 2017, 1, 25-31. | 2.6 | 6 |
| 11 | 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 |
| 12 | 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 |
| 13 | A Comprehensive Analysis of MicroProteins Reveals Their Potentially Widespread Mechanism of Transcriptional Regulation. <i>Plant Physiology</i> , 2014, 165, 149-159. | 4.8 | 21 |