

Iris Jonkers

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8284499/publications.pdf>

Version: 2024-02-01

14
papers

1,465
citations

840776

11
h-index

1058476

14
g-index

17
all docs

17
docs citations

17
times ranked

4518
citing authors

#	ARTICLE	IF	CITATIONS
1	Gut mucosa dissociation protocols influence cell type proportions and single-cell gene expression levels. <i>Scientific Reports</i> , 2022, 12, .	3.3	23
2	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
3	Inflammatory Protein Profiles in Plasma of Candidaemia Patients and the Contribution of Host Genetics to Their Variability. <i>Frontiers in Immunology</i> , 2021, 12, 662171.	4.8	6
4	A Combined mRNA- and miRNA-Sequencing Approach Reveals miRNAs as Potential Regulators of the Small Intestinal Transcriptome in Celiac Disease. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11382.	4.1	6
5	Molecular Biomarkers for Celiac Disease: Past, Present and Future. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8528.	4.1	13
6	Deconvolution of bulk blood eQTL effects into immune cell subpopulations. <i>BMC Bioinformatics</i> , 2020, 21, 243.	2.6	38
7	Systematic Prioritization of Candidate Genes in Disease Loci Identifies TRAFD1 as a Master Regulator of IFN γ Signaling in Celiac Disease. <i>Frontiers in Genetics</i> , 2020, 11, 562434.	2.3	20
8	A Genome-Wide Functional Genomics Approach Identifies Susceptibility Pathways to Fungal Bloodstream Infection in Humans. <i>Journal of Infectious Diseases</i> , 2019, 220, 862-872.	4.0	17
9	A liver-specific long noncoding RNA with a role in cell viability is elevated in human nonalcoholic steatohepatitis. <i>Hepatology</i> , 2017, 66, 794-808.	7.3	80
10	Disease variants alter transcription factor levels and methylation of their binding sites. <i>Nature Genetics</i> , 2017, 49, 131-138.	21.4	390
11	An integrative genomics approach identifies novel pathways that influence candidaemia susceptibility. <i>PLoS ONE</i> , 2017, 12, e0180824.	2.5	24
12	Understanding human immune function using the resources from the Human Functional Genomics Project. <i>Nature Medicine</i> , 2016, 22, 831-833.	30.7	63
13	Understanding Celiac Disease by Genomics. <i>Trends in Genetics</i> , 2016, 32, 295-308.	6.7	78
14	Getting up to speed with transcription elongation by RNA polymerase II. <i>Nature Reviews Molecular Cell Biology</i> , 2015, 16, 167-177.	37.0	692