Iris Jonkers

List of Publications by Year in descending order

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IDIS LONKEDS

#	Article	IF	CITATIONS
1	Getting up to speed with transcription elongation by RNA polymerase II. Nature Reviews Molecular Cell Biology, 2015, 16, 167-177.	37.0	692
2	Disease variants alter transcription factor levels and methylation of their binding sites. Nature Genetics, 2017, 49, 131-138.	21.4	390
3	A liverâ€specific long noncoding RNA with a role in cell viability is elevated in human nonalcoholic steatohepatitis. Hepatology, 2017, 66, 794-808.	7.3	80
4	Understanding Celiac Disease by Genomics. Trends in Genetics, 2016, 32, 295-308.	6.7	78
5	Understanding human immune function using the resources from the Human Functional Genomics Project. Nature Medicine, 2016, 22, 831-833.	30.7	63
6	Deconvolution of bulk blood eQTL effects into immune cell subpopulations. BMC Bioinformatics, 2020, 21, 243.	2.6	38
7	An integrative genomics approach identifies novel pathways that influence candidaemia susceptibility. PLoS ONE, 2017, 12, e0180824.	2.5	24
8	Gut mucosa dissociation protocols influence cell type proportions and single-cell gene expression levels. Scientific Reports, 2022, 12, .	3.3	23
9	Systematic Prioritization of Candidate Genes in Disease Loci Identifies TRAFD1 as a Master Regulator of IFNÎ ³ Signaling in Celiac Disease. Frontiers in Genetics, 2020, 11, 562434.	2.3	20
10	A Genome-Wide Functional Genomics Approach Identifies Susceptibility Pathways to Fungal Bloodstream Infection in Humans. Journal of Infectious Diseases, 2019, 220, 862-872.	4.0	17
11	Molecular Biomarkers for Celiac Disease: Past, Present and Future. International Journal of Molecular Sciences, 2020, 21, 8528.	4.1	13
12	Potential impact of celiac disease genetic risk factors on T cell receptor signaling in gluten-specific CD4+ T cells. Scientific Reports, 2021, 11, 9252.	3.3	6
13	Inflammatory Protein Profiles in Plasma of Candidaemia Patients and the Contribution of Host Genetics to Their Variability. Frontiers in Immunology, 2021, 12, 662171.	4.8	6
14	A Combined mRNA- and miRNA-Sequencing Approach Reveals miRNAs as Potential Regulators of the Small Intestinal Transcriptome in Celiac Disease. International Journal of Molecular Sciences, 2021, 22, 11382.	4.1	6