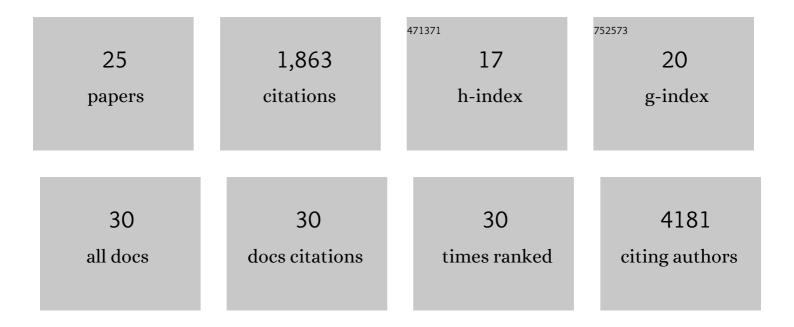
Monique G P Van Der Wijst

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7908824/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Gut mucosa dissociation protocols influence cell type proportions and single-cell gene expression levels. Scientific Reports, 2022, 12, .	1.6	23
2	Single-cell RNA-sequencing of peripheral blood mononuclear cells reveals widespread, context-specific gene expression regulation upon pathogenic exposure. Nature Communications, 2022, 13, .	5.8	39
3	Genetic, parental and lifestyle factors influence telomere length. Communications Biology, 2022, 5, .	2.0	23
4	Type I interferon autoantibodies are associated with systemic immune alterations in patients with COVID-19. Science Translational Medicine, 2021, 13, eabh2624.	5.8	155
5	Large-scale cis- and trans-eQTL analyses identify thousands of genetic loci and polygenic scores that regulate blood gene expression. Nature Genetics, 2021, 53, 1300-1310.	9.4	590
6	Deconvolution of bulk blood eQTL effects into immune cell subpopulations. BMC Bioinformatics, 2020, 21, 243.	1.2	38
7	Integrating GWAS with bulk and single-cell RNA-sequencing reveals a role for LY86 in the anti-Candida host response. PLoS Pathogens, 2020, 16, e1008408.	2.1	18
8	The single-cell eQTLGen consortium. ELife, 2020, 9, .	2.8	150
9	Title is missing!. , 2020, 16, e1008408.		0
10	Title is missing!. , 2020, 16, e1008408.		0
11	Title is missing!. , 2020, 16, e1008408.		0
12	Title is missing!. , 2020, 16, e1008408.		0
13	Title is missing!. , 2020, 16, e1008408.		Ο
14	Single-cell RNA sequencing identifies celltype-specific cis-eQTLs and co-expression QTLs. Nature Genetics, 2018, 50, 493-497.	9.4	289
15	An integrative approach for building personalized gene regulatory networks for precision medicine. Genome Medicine, 2018, 10, 96.	3.6	49
16	Importance of Metal-Ion Exchange for the Biological Activity of Coordination Complexes of the Biomimetic Ligand N4Py. Inorganic Chemistry, 2018, 57, 7748-7756.	1.9	23
17	Experimental mitochondria-targeted DNA methylation identifies GpC methylation, not CpG methylation, as potential regulator of mitochondrial gene expression. Scientific Reports, 2017, 7, 177.	1.6	72
18	Regulation of mitochondrial gene expression the epigenetic enigma. Frontiers in Bioscience - Landmark. 2017. 22. 1099-1113.	3.0	69

#	Article	IF	CITATIONS
19	Re-expression of Selected Epigenetically Silenced Candidate Tumor Suppressor Genes in Cervical Cancer by TET2-directed Demethylation. Molecular Therapy, 2016, 24, 536-547.	3.7	33
20	Local chromatin microenvironment determines DNMT activity: from DNA methyltransferase to DNA denverse de demethylase or DNA dehydroxymethylase. Epigenetics, 2015, 10, 671-676.	1.3	72
21	Mitochondrial epigenetics: an overlooked layer of regulation?. Trends in Genetics, 2015, 31, 353-356.	2.9	85
22	Targeting Nrf2 in healthy and malignant ovarian epithelial cells: Protection versus promotion. Molecular Oncology, 2015, 9, 1259-1273.	2.1	17
23	Prolonged re-expression of the hypermethylated gene <i>EPB41L3</i> using artificial transcription factors and epigenetic drugs. Epigenetics, 2015, 10, 384-396.	1.3	28
24	Nrf2, the master redox switch: The Achilles' heel of ovarian cancer?. Biochimica Et Biophysica Acta: Reviews on Cancer, 2014, 1846, 494-509.	3.3	36
25	Efficient Nuclear DNA Cleavage in Human Cancer Cells by Synthetic Bleomycin Mimics. ACS Chemical Biology, 2014, 9, 1044-1051.	1.6	23