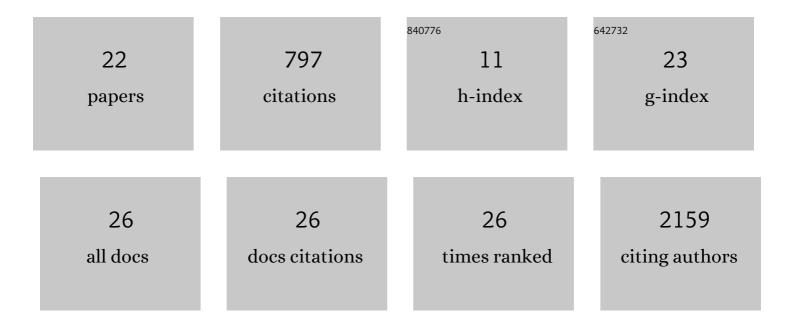
## Dennis J Montoya

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2778936/publications.pdf Version: 2024-02-01



DENNIS I MONTOVA

#	Article	IF	CITATIONS
1	DNA methylation profiles in pneumonia patients reflect changes in cell types and pneumonia severity. Epigenetics, 2022, 17, 1646-1660.	2.7	5
2	The Gene Expression Deconvolution Interactive Tool (GEDIT): accurate cell type quantification from gene expression data. GigaScience, 2021, 10, .	6.4	33
3	Effect of the p53 P72R Polymorphism on Mutant <i>TP53</i> Allele Selection in Human Cancer. Journal of the National Cancer Institute, 2021, 113, 1246-1257.	6.3	16
4	Human DNA methylation signatures differentiate persistent from resolving MRSA bacteremia. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	14
5	Systematic evaluation of transcriptomics-based deconvolution methods and references using thousands of clinical samples. Briefings in Bioinformatics, 2021, 22, .	6.5	10
6	Quinacrine Induces Nucleolar Stress in Treatment-Refractory Ovarian Cancer Cell Lines. Cancers, 2021, 13, 4645.	3.7	7
7	The P72R Polymorphism in R248Q/W p53 Mutants Modifies the Mutant Effect on Epithelial to Mesenchymal Transition Phenotype and Cell Invasion via CXCL1 Expression. International Journal of Molecular Sciences, 2020, 21, 8025.	4.1	4
8	Integrative Analysis of Glucometabolic Traits, Adipose Tissue DNA Methylation, and Gene Expression Identifies Epigenetic Regulatory Mechanisms of Insulin Resistance and Obesity in African Americans. Diabetes, 2020, 69, 2779-2793.	0.6	8
9	Profiling immunoglobulin repertoires across multiple human tissues using RNA sequencing. Nature Communications, 2020, 11, 3126.	12.8	44
10	The cell fate regulator NUPR1 is induced by Mycobacterium leprae via type I interferon in human leprosy. PLoS Neglected Tropical Diseases, 2019, 13, e0007589.	3.0	7
11	Vitamin A Metabolism by Dendritic Cells Triggers an Antimicrobial Response against Mycobacterium tuberculosis. MSphere, 2019, 4, .	2.9	14
12	Dual RNA-Seq of Human Leprosy Lesions Identifies Bacterial Determinants Linked to Host Immune Response. Cell Reports, 2019, 26, 3574-3585.e3.	6.4	38
13	Plasticity of antimicrobial and phagocytic programs in human macrophages. Immunology, 2019, 156, 164-173.	4.4	20
14	Genetic Mechanisms of Immune Evasion in Colorectal Cancer. Cancer Discovery, 2018, 8, 730-749.	9.4	367
15	Epigenome-wide association in adipose tissue from the METSIM cohort. Human Molecular Genetics, 2018, 27, 1830-1846.	2.9	38
16	Prenatal Growth Patterns and Birthweight Are Associated With Differential DNA Methylation and Gene Expression of Cardiometabolic Risk Genes in Human Placentas: A Discovery-Based Approach. Reproductive Sciences, 2018, 25, 523-539.	2.5	41
17	A phylogenomic study quantifies competing mechanisms for pseudogenization in prokaryotes—The Mycobacterium leprae case. PLoS ONE, 2018, 13, e0204322.	2.5	3
18	SaVanT: a web-based tool for the sample-level visualization of molecular signatures in gene expression profiles. BMC Genomics, 2017, 18, 824.	2.8	32

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#	Article	IF	CITATIONS
19	Epigenetic changes in Tâ€eell and monocyte signatures and production of neurotoxic cytokines in ALS patients. FASEB Journal, 2016, 30, 3461-3473.	0.5	41
20	Jagged1 Instructs Macrophage Differentiation in Leprosy. PLoS Pathogens, 2016, 12, e1005808.	4.7	32
21	Anti-inflammatory therapies of amyotrophic lateral sclerosis guided by immune pathways. American Journal of Neurodegenerative Disease, 2015, 4, 28-39.	0.1	6
22	ILâ€1beta triggers monocytes to differentiate into CD209+ macrophages. FASEB Journal, 2008, 22, 539-539.	0.5	0