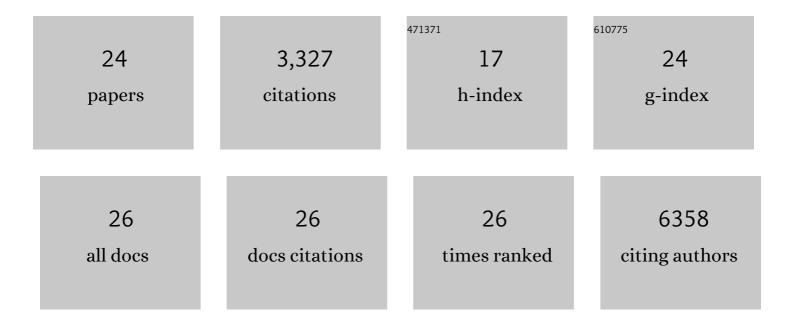
Javier RodrÃ-guez-Ubreva

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9434946/publications.pdf

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#	Article	IF	CITATIONS
1	Autophagy maintains stemness by preventing senescence. Nature, 2016, 529, 37-42.	13.7	1,013
2	Geriatric muscle stem cells switch reversible quiescence into senescence. Nature, 2014, 506, 316-321.	13.7	785
3	Changes in the pattern of DNA methylation associate with twin discordance in systemic lupus erythematosus. Genome Research, 2010, 20, 170-179.	2.4	569
4	PU.1 target genes undergo Tet2-coupled demethylation and DNMT3b-mediated methylation in monocyte-to-osteoclast differentiation. Genome Biology, 2013, 14, R99.	13.9	177
5	IL-4 orchestrates STAT6-mediated DNA demethylation leading to dendritic cell differentiation. Genome Biology, 2016, 17, 4.	3.8	122
6	Prostaglandin E2 Leads to the Acquisition of DNMT3A-Dependent Tolerogenic Functions in Human Myeloid-Derived Suppressor Cells. Cell Reports, 2017, 21, 154-167.	2.9	116
7	Monozygotic twins discordant for common variable immunodeficiency reveal impaired DNA demethylation during naÃ ⁻ ve-to-memory B-cell transition. Nature Communications, 2015, 6, 7335.	5.8	81
8	Epigenetic interplay between immune, stromal and cancer cells in the tumor microenvironment. Clinical Immunology, 2018, 196, 64-71.	1.4	61
9	Inflammatory cytokines shape a changing DNA methylome in monocytes mirroring disease activity in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2019, 78, 1505-1516.	0.5	47
10	In vivo conditional deletion of HDAC7 reveals its requirement to establish proper B lymphocyte identity and development. Journal of Experimental Medicine, 2016, 213, 2591-2601.	4.2	39
11	Pre-B cell to macrophage transdifferentiation without significant promoter DNA methylation changes. Nucleic Acids Research, 2012, 40, 1954-1968.	6.5	37
12	TET2- and TDG-mediated changes are required for the acquisition of distinct histone modifications in divergent terminal differentiation of myeloid cells. Nucleic Acids Research, 2017, 45, 10002-10017.	6.5	36
13	Chromatin Immunoprecipitation. Methods in Molecular Biology, 2014, 1094, 309-318.	0.4	32
14	Targeting aberrant DNA methylation in mesenchymal stromal cells as a treatment for myeloma bone disease. Nature Communications, 2021, 12, 421.	5.8	29
15	NF-κB directly mediates epigenetic deregulation of common microRNAs in Epstein-Barr virus-mediated transformation of B-cells and in lymphomas. Nucleic Acids Research, 2014, 42, 11025-11039.	6.5	27
16	C/EBPa-Mediated Activation of MicroRNAs 34a and 223 Inhibits Lef1 Expression To Achieve Efficient Reprogramming into Macrophages. Molecular and Cellular Biology, 2014, 34, 1145-1157.	1.1	26
17	STATegra, a comprehensive multi-omics dataset of B-cell differentiation in mouse. Scientific Data, 2019, 6, 256.	2.4	26
18	Single-cell Atlas of common variable immunodeficiency shows germinal center-associated epigenetic dysregulation in B-cell responses. Nature Communications. 2022, 13, 1779.	5.8	25

#	Article	IF	CITATIONS
19	Impaired CpG Demethylation in Common Variable Immunodeficiency Associates With B Cell Phenotype and Proliferation Rate. Frontiers in Immunology, 2019, 10, 878.	2.2	19
20	Prediction of the Progression of Undifferentiated Arthritis to Rheumatoid Arthritis Using DNA Methylation Profiling. Arthritis and Rheumatology, 2021, 73, 2229-2239.	2.9	16
21	Epigenetic mechanisms of myeloid differentiation in the tumor microenvironment. Current Opinion in Pharmacology, 2017, 35, 20-29.	1.7	12
22	The synovial and blood monocyte DNA methylomes mirror prognosis, evolution, and treatment in early arthritis. JCI Insight, 2022, 7, .	2.3	11
23	Dissecting Epigenetic Dysregulation of Primary Antibody Deficiencies. Journal of Clinical Immunology, 2016, 36, 48-56.	2.0	10
24	Activation-induced cytidine deaminase targets SUV4-20-mediated histone H4K20 trimethylation to class-switch recombination sites. Scientific Reports, 2017, 7, 7594.	1.6	10