Celia P Martinez Jimenez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6957534/publications.pdf

Version: 2024-02-01

22 papers 1,961 citations

623734 14 h-index 677142 22 g-index

27 all docs

27 docs citations

times ranked

27

4382 citing authors

#	Article	IF	CITATIONS
1	Five-Vertebrate ChIP-seq Reveals the Evolutionary Dynamics of Transcription Factor Binding. Science, 2010, 328, 1036-1040.	12.6	663
2	Aging increases cell-to-cell transcriptional variability upon immune stimulation. Science, 2017, 355, 1433-1436.	12.6	265
3	Staged developmental mapping and X chromosome transcriptional dynamics during mouse spermatogenesis. Nature Communications, 2019, 10, 1251.	12.8	177
4	Hepatocyte cell lines: their use, scope and limitations in drug metabolism studies. Expert Opinion on Drug Metabolism and Toxicology, 2006, 2, 183-212.	3.3	173
5	Hepatocyte Nuclear Factor 4α Coordinates a Transcription Factor Network Regulating Hepatic Fatty Acid Metabolism. Molecular and Cellular Biology, 2010, 30, 565-577.	2.3	132
6	Transcriptional Regulation and Expression of CYP3A4 in Hepatocytes. Current Drug Metabolism, 2007, 8, 185-194.	1.2	122
7	Transcriptional Activation of CYP2C9, CYP1A1, and CYP1A2 by Hepatocyte Nuclear Factor 4α Requires Coactivators Peroxisomal Proliferator Activated Receptor-γ Coactivator 1α and Steroid Receptor Coactivator 1. Molecular Pharmacology, 2006, 70, 1681-1692.	2.3	63
8	Transcriptional Regulation of the Human Hepatic CYP3A4: Identification of a New Distal Enhancer Region Responsive to CCAAT/Enhancer-Binding Protein \hat{I}^2 Isoforms (Liver Activating Protein and Liver) Tj ETQq0 C) 02 <i>g</i> BT /C	Overstock 10 Tf
9	Underexpressed Coactivators PGC1α AND SRC1 Impair Hepatocyte Nuclear Factor 4α Function and Promote Dedifferentiation in Human Hepatoma Cells. Journal of Biological Chemistry, 2006, 281, 29840-29849.	3.4	55
10	Dominant and Redundant Functions of TFIID Involved in the Regulation of Hepatic Genes. Molecular Cell, 2008, 31, 531-543.	9.7	54
11	Single-nucleus RNA-seq2 reveals functional crosstalk between liver zonation and ploidy. Nature Communications, 2021, 12, 4264.	12.8	46
12	<scp>IL</scp> â€7â€dependent compositional changes within the γδT cell pool in lymph nodes during ageing lead toÂan unbalanced antiâ€ŧumour response. EMBO Reports, 2019, 20, e47379.	4.5	38
13	Innovative approach for self-management and social welfare of children with cystic fibrosis in Europe: development, validation and implementation of an mHealth tool (MyCyFAPP). BMJ Open, 2017, 7, e014931.	1.9	28
14	Comprehensive analysis of interacting proteins and genomeâ€wide location studies of the Sas3â€dependent NuA3 histone acetyltransferase complex. FEBS Open Bio, 2014, 4, 996-1006.	2.3	15
15	Epigenetic crosstalk a molecular language in human metabolic disorders. Frontiers in Bioscience - Scholar, 2015, 7, 46-57.	2.1	13
16	Inactivation of the Nuclear Orphan Receptor COUP-TFII by Small Chemicals. ACS Chemical Biology, 2017, 12, 654-663.	3.4	13
17	Advances of single-cell genomics and epigenomics in human disease: where are we now?. Mammalian Genome, 2020, 31, 170-180.	2.2	9
18	Species-specific mechanisms for cholesterol 7α-hydroxylase (CYP7A1) regulation by drugs and bile acids. Archives of Biochemistry and Biophysics, 2005, 434, 75-85.	3.0	8

#	Article	IF	CITATIONS
19	The mechanisms shaping the single-cell transcriptional landscape. Current Opinion in Genetics and Development, 2016, 37, 27-35.	3.3	7
20	Can Hepatoma Cell Lines be Re-differentiated to be Used in Drug Metabolism Studies?. ATLA Alternatives To Laboratory Animals, 2004, 32, 65-74.	1.0	5
21	Transcriptional Elongation and mRNA Export Are Coregulated Processes. Genetics Research International, 2011, 2011, 1-10.	2.0	3
22	Hepatocyte-specific activity of TSC22D4 triggers progressive NAFLD by impairing mitochondrial function. Molecular Metabolism, 2022, 60, 101487.	6.5	3