

David Arenillas

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12120392/publications.pdf>

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9
papers

1,747
citations

1163117

8
h-index

1474206

9
g-index

9
all docs

9
docs citations

9
times ranked

4731
citing authors

#	ARTICLE	IF	CITATIONS
1	Update of the FANTOM web resource: high resolution transcriptome of diverse cell types in mammals. <i>Nucleic Acids Research</i> , 2017, 45, D737-D743.	14.5	116
2	A SNP in the HTT promoter alters NF- κ B binding and is a bidirectional genetic modifier of Huntington disease. <i>Nature Neuroscience</i> , 2015, 18, 807-816.	14.8	113
3	Combined serial analysis of gene expression and transcription factor binding site prediction identifies novel-candidate-target genes of Nr2e1 in neocortex development. <i>BMC Genomics</i> , 2015, 16, 545.	2.8	9
4	MIR@NT@N: a framework integrating transcription factors, microRNAs and their targets to identify sub-network motifs in a meta-regulation network model. <i>BMC Bioinformatics</i> , 2011, 12, 67.	2.6	64
5	Global mapping of binding sites for Nrf2 identifies novel targets in cell survival response through ChIP-Seq profiling and network analysis. <i>Nucleic Acids Research</i> , 2010, 38, 5718-5734.	14.5	653
6	JASPAR 2010: the greatly expanded open-access database of transcription factor binding profiles. <i>Nucleic Acids Research</i> , 2010, 38, D105-D110.	14.5	529
7	The PAZAR database of gene regulatory information coupled to the ORCA toolkit for the study of regulatory sequences. <i>Nucleic Acids Research</i> , 2009, 37, D54-D60.	14.5	97
8	Mechanisms underlying p53 regulation of PIK3CA transcription in ovarian surface epithelium and in ovarian cancer. <i>Journal of Cell Science</i> , 2008, 121, 664-674.	2.0	72
9	In Silico Detection of Sequence Variations Modifying Transcriptional Regulation. <i>PLoS Computational Biology</i> , 2008, 4, e5.	3.2	94