

Stefan Ge Roberts

List of Publications by Year in Descending Order

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Version: 2024-04-28

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28
papers

841
citations

15
h-index

29
g-index

34
ext. papers

941
ext. citations

7.4
avg, IF

4.18
L-index

#	Paper	IF	Citations
28	Cholesterol is required for transcriptional repression by BASP1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	2
27	WT1 activates transcription of the splice factor kinase SRPK1 gene in PC3 and K562 cancer cells in the absence of corepressor BASP1. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2020 , 1863, 194642	6	5
26	The WT1-BASP1 complex is required to maintain the differentiated state of taste receptor cells. <i>Life Science Alliance</i> , 2019 , 2,	5.8	6
25	IDPpi: Protein-Protein Interaction Analyses of Human Intrinsically Disordered Proteins. <i>Scientific Reports</i> , 2018 , 8, 10563	4.9	10
24	BASP1 interacts with oestrogen receptor β and modifies the tamoxifen response. <i>Cell Death and Disease</i> , 2017 , 8, e2771	9.8	14
23	A transcription factor IIA-binding site differentially regulates RNA polymerase II-mediated transcription in a promoter context-dependent manner. <i>Journal of Biological Chemistry</i> , 2017 , 292, 11873-11885	5.4	8
22	TRI_tool: a web-tool for prediction of protein-protein interactions in human transcriptional regulation. <i>Bioinformatics</i> , 2017 , 33, 289-291	7.2	9
21	In Vitro Transcription to Study WT1 Function. <i>Methods in Molecular Biology</i> , 2016 , 1467, 137-54	1.4	
20	A role of WT1 in cell division and genomic stability. <i>Cell Cycle</i> , 2015 , 14, 1358-64	4.7	17
19	Classification of a frameshift/extended and a stop mutation in WT1 as gain-of-function mutations that activate cell cycle genes and promote Wilms tumour cell proliferation. <i>Human Molecular Genetics</i> , 2014 , 23, 3958-74	5.6	10
18	The transcription cycle in eukaryotes: from productive initiation to RNA polymerase II recycling. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2012 , 1819, 391-400	6	75
17	HtrA2, taming the oncogenic activities of WT1. <i>Cell Cycle</i> , 2010 , 9, 2508-14	4.7	8
16	New insights into the role of TFIIB in transcription initiation. <i>Transcription</i> , 2010 , 1, 126-129	4.8	10
15	The Wilms tumor suppressor protein WT1 is processed by the serine protease HtrA2/Omi. <i>Molecular Cell</i> , 2010 , 37, 159-71	17.6	60
14	Phosphorylation of TFIIB links transcription initiation and termination. <i>Current Biology</i> , 2010 , 20, 548-53	6.3	50
13	The modulation of WTI transcription function by cofactors. <i>Biochemical Society Symposia</i> , 2006 , 191-201		15
12	Transcriptional regulation by WT1 in development. <i>Current Opinion in Genetics and Development</i> , 2005 , 15, 542-7	4.9	82

11	A core promoter element downstream of the TATA box that is recognized by TFIIB. <i>Genes and Development</i> , 2005 , 19, 2418-23	12.6	101
10	Two molecular subgroups of Wilmsctumors with or without WT1 mutations. <i>Clinical Cancer Research</i> , 2003 , 9, 2005-14	12.9	46
9	Activator-mediated disruption of sequence-specific DNA contacts by the general transcription factor TFIIB. <i>Genes and Development</i> , 2001 , 15, 2945-9	12.6	50
8	Par4 is a coactivator for a splice isoform-specific transcriptional activation domain in WT1. <i>Genes and Development</i> , 2001 , 15, 328-39	12.6	66
7	Expression of the Oct-1 transcription factor and characterization of its interactions with the Bob1 coactivator. <i>Biochemistry</i> , 2001 , 40, 6580-8	3.2	29
6	The conformation of the transcription factor TFIIB modulates the response to transcriptional activators in vivo. <i>Current Biology</i> , 2000 , 10, 273-6	6.3	26
5	Regulation of the Wilmsctumour suppressor protein transcriptional activation domain. <i>Oncogene</i> , 1999 , 18, 6546-54	9.2	23
4	The role of human TFIIB in transcription start site selection in vitro and in vivo. <i>Journal of Biological Chemistry</i> , 1999 , 274, 14337-43	5.4	53
3	Purification and analysis of functional preinitiation complexes. <i>Methods in Enzymology</i> , 1996 , 273, 110-8	1.7	6
2	A role for activator-mediated TFIIB recruitment in diverse aspects of transcriptional regulation. <i>Current Biology</i> , 1995 , 5, 508-16	6.3	53
1	The mouse proline-rich protein MP6 promoter binds isoprenaline-inducible parotid nuclear proteins via a highly conserved NFkB/rel-like site. <i>Nucleic Acids Research</i> , 1991 , 19, 5205-11	20.1	6