

# Stefan Ge Roberts

## List of Publications by Citations

**Source:** <https://exaly.com/author-pdf/6001515/stefan-ge-roberts-publications-by-citations.pdf>

**Version:** 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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	A core promoter element downstream of the TATA box that is recognized by TFIIB. <i>Genes and Development</i> , <b>2005</b> , 19, 2418-23	12.6	101
27	Transcriptional regulation by WT1 in development. <i>Current Opinion in Genetics and Development</i> , <b>2005</b> , 15, 542-7	4.9	82
26	The transcription cycle in eukaryotes: from productive initiation to RNA polymerase II recycling. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , <b>2012</b> , 1819, 391-400	6	75
25	Par4 is a coactivator for a splice isoform-specific transcriptional activation domain in WT1. <i>Genes and Development</i> , <b>2001</b> , 15, 328-39	12.6	66
24	The Wilmsctumor suppressor protein WT1 is processed by the serine protease HtrA2/Omi. <i>Molecular Cell</i> , <b>2010</b> , 37, 159-71	17.6	60
23	A role for activator-mediated TFIIB recruitment in diverse aspects of transcriptional regulation. <i>Current Biology</i> , <b>1995</b> , 5, 508-16	6.3	53
22	The role of human TFIIB in transcription start site selection in vitro and in vivo. <i>Journal of Biological Chemistry</i> , <b>1999</b> , 274, 14337-43	5.4	53
21	Phosphorylation of TFIIB links transcription initiation and termination. <i>Current Biology</i> , <b>2010</b> , 20, 548-53	6.3	50
20	Activator-mediated disruption of sequence-specific DNA contacts by the general transcription factor TFIIB. <i>Genes and Development</i> , <b>2001</b> , 15, 2945-9	12.6	50
19	Two molecular subgroups of Wilmsctumors with or without WT1 mutations. <i>Clinical Cancer Research</i> , <b>2003</b> , 9, 2005-14	12.9	46
18	Expression of the Oct-1 transcription factor and characterization of its interactions with the Bob1 coactivator. <i>Biochemistry</i> , <b>2001</b> , 40, 6580-8	3.2	29
17	The conformation of the transcription factor TFIIB modulates the response to transcriptional activators in vivo. <i>Current Biology</i> , <b>2000</b> , 10, 273-6	6.3	26
16	Regulation of the Wilmsctumour suppressor protein transcriptional activation domain. <i>Oncogene</i> , <b>1999</b> , 18, 6546-54	9.2	23
15	A role of WT1 in cell division and genomic stability. <i>Cell Cycle</i> , <b>2015</b> , 14, 1358-64	4.7	17
14	The modulation of WTI transcription function by cofactors. <i>Biochemical Society Symposia</i> , <b>2006</b> , 191-201		15
13	BASP1 interacts with oestrogen receptor and modifies the tamoxifen response. <i>Cell Death and Disease</i> , <b>2017</b> , 8, e2771	9.8	14
12	IDPpi: Protein-Protein Interaction Analyses of Human Intrinsically Disordered Proteins. <i>Scientific Reports</i> , <b>2018</b> , 8, 10563	4.9	10

11	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> , <b>2014</b> , 23, 3958-74	5.6	10
10	New insights into the role of TFIIB in transcription initiation. <i>Transcription</i> , <b>2010</b> , 1, 126-129	4.8	10
9	TRI_tool: a web-tool for prediction of protein-protein interactions in human transcriptional regulation. <i>Bioinformatics</i> , <b>2017</b> , 33, 289-291	7.2	9
8	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> , <b>2017</b> , 292, 11873-11883	5.4	8
7	HtrA2, taming the oncogenic activities of WT1. <i>Cell Cycle</i> , <b>2010</b> , 9, 2508-14	4.7	8
6	Purification and analysis of functional preinitiation complexes. <i>Methods in Enzymology</i> , <b>1996</b> , 273, 110-8	1.7	6
5	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> , <b>1991</b> , 19, 5205-11	20.1	6
4	The WT1-BASP1 complex is required to maintain the differentiated state of taste receptor cells. <i>Life Science Alliance</i> , <b>2019</b> , 2,	5.8	6
3	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> , <b>2020</b> , 1863, 194642	6	5
2	Cholesterol is required for transcriptional repression by BASP1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2021</b> , 118,	11.5	2
1	In Vitro Transcription to Study WT1 Function. <i>Methods in Molecular Biology</i> , <b>2016</b> , 1467, 137-54	1.4	