

Stefan Ge Roberts

List of Publications by Year in descending order

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Version: 2024-02-01

30
papers

1,037
citations

516710

16
h-index

501196

28
g-index

34
all docs

34
docs citations

34
times ranked

1322
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Crosstalk between β -catenin and WT1 signaling activity in acute myeloid leukemia. <i>Haematologica</i> , 2023, 108, 283-289. | 3.5 | 4 |
| 2 | 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, . | 7.1 | 11 |
| 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> , 2020, 1863, 194642. | 1.9 | 14 |
| 4 | The WT1-BASP1 complex is required to maintain the differentiated state of taste receptor cells. <i>Life Science Alliance</i> , 2019, 2, e201800287. | 2.8 | 11 |
| 5 | IDPpi: Protein-Protein Interaction Analyses of Human Intrinsically Disordered Proteins. <i>Scientific Reports</i> , 2018, 8, 10563. | 3.3 | 18 |
| 6 | BASP1 interacts with oestrogen receptor β and modifies the tamoxifen response. <i>Cell Death and Disease</i> , 2017, 8, e2771-e2771. | 6.3 | 26 |
| 7 | 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. | 3.4 | 12 |
| 8 | TRI_tool: a web-tool for prediction of protein-protein interactions in human transcriptional regulation. <i>Bioinformatics</i> , 2017, 33, 289-291. | 4.1 | 17 |
| 9 | In Vitro Transcription to Study WT1 Function. <i>Methods in Molecular Biology</i> , 2016, 1467, 137-154. | 0.9 | 1 |
| 10 | A role of WT1 in cell division and genomic stability. <i>Cell Cycle</i> , 2015, 14, 1358-1364. | 2.6 | 24 |
| 11 | Abstract 3784: Regulation of chromatin condensation by mitotic checkpoint protein MAD2. , 2015, , . | | 0 |
| 12 | 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, 3958-3974. | 2.9 | 15 |
| 13 | 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. | 1.9 | 101 |
| 14 | Phosphorylation of TFIIB Links Transcription Initiation and Termination. <i>Current Biology</i> , 2010, 20, 548-553. | 3.9 | 62 |
| 15 | HtrA2, taming the oncogenic activities of WT1. <i>Cell Cycle</i> , 2010, 9, 2508-2514. | 2.6 | 10 |
| 16 | New insights into the role of TFIIB in transcription initiation. <i>Transcription</i> , 2010, 1, 126-129. | 3.1 | 11 |
| 17 | The Wilms' Tumor Suppressor Protein WT1 Is Processed by the Serine Protease HtrA2/Omi. <i>Molecular Cell</i> , 2010, 37, 159-171. | 9.7 | 69 |
| 18 | The modulation of WTI transcription function by cofactors. <i>Biochemical Society Symposia</i> , 2006, 73, 191-201. | 2.7 | 17 |

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|----|---|------|-----------|
| 19 | A core promoter element downstream of the TATA box that is recognized by TFIIB. <i>Genes and Development</i> , 2005, 19, 2418-2423. | 5.9 | 127 |
| 20 | Transcriptional regulation by WT1 in development. <i>Current Opinion in Genetics and Development</i> , 2005, 15, 542-547. | 3.3 | 91 |
| 21 | Two molecular subgroups of Wilms' tumors with or without WT1 mutations. <i>Clinical Cancer Research</i> , 2003, 9, 2005-14. | 7.0 | 49 |
| 22 | Expression of the Oct-1 Transcription Factor and Characterization of Its Interactions with the Bob1 Coactivator. <i>Biochemistry</i> , 2001, 40, 6580-6588. | 2.5 | 33 |
| 23 | Activator-mediated disruption of sequence-specific DNA contacts by the general transcription factor TFIIB. <i>Genes and Development</i> , 2001, 15, 2945-2949. | 5.9 | 57 |
| 24 | Par4 is a coactivator for a splice isoform-specific transcriptional activation domain in WT1. <i>Genes and Development</i> , 2001, 15, 328-339. | 5.9 | 76 |
| 25 | The conformation of the transcription factor TFIIB modulates the response to transcriptional activators in vivo. <i>Current Biology</i> , 2000, 10, 273-276. | 3.9 | 27 |
| 26 | Regulation of the Wilms' tumour suppressor protein transcriptional activation domain. <i>Oncogene</i> , 1999, 18, 6546-6554. | 5.9 | 27 |
| 27 | The Role of Human TFIIB in Transcription Start Site Selection in Vitro and in Vivo. <i>Journal of Biological Chemistry</i> , 1999, 274, 14337-14343. | 3.4 | 57 |
| 28 | [8] Purification and analysis of functional preinitiation complexes preinitiation complexes. <i>Methods in Enzymology</i> , 1996, 273, 110-118. | 1.0 | 6 |
| 29 | A role for activator-mediated TFIIB recruitment in diverse aspects of transcriptional regulation. <i>Current Biology</i> , 1995, 5, 508-516. | 3.9 | 57 |
| 30 | 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, 5205-5211. | 14.5 | 7 |