

Rudolf Grosschedl

List of Publications by Year in descending order

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49
papers

9,680
citations

145106

33
h-index

232693

48
g-index

50
all docs

50
docs citations

50
times ranked

13593
citing authors

#	ARTICLE	IF	CITATIONS
1	How to resist Notch-targeted T-leukemia therapy: Lineage- and MYC enhancer switch. <i>Molecular Cell</i> , 2022, 82, 884-886.	4.5	0
2	EBF1 promotes triple-negative breast cancer progression by surveillance of the HIF1 α pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	4
3	ZFP451-mediated SUMOylation of SATB2 drives embryonic stem cell differentiation. <i>Genes and Development</i> , 2021, 35, 1142-1160.	2.7	9
4	DNA methylation signatures reveal that distinct combinations of transcription factors specify human immune cell epigenetic identity. <i>Immunity</i> , 2021, 54, 2465-2480.e5.	6.6	31
5	EBF1 and Pax5 safeguard leukemic transformation by limiting IL-7 signaling, Myc expression, and folate metabolism. <i>Genes and Development</i> , 2020, 34, 1503-1519.	2.7	15
6	Interactions between lineage-associated transcription factors govern haematopoietic progenitor states. <i>EMBO Journal</i> , 2020, 39, e104983.	3.5	20
7	A Prion-like Domain in Transcription Factor EBF1 Promotes Phase Separation and Enables B Cell Programming of Progenitor Chromatin. <i>Immunity</i> , 2020, 53, 1151-1167.e6.	6.6	47
8	EBF1-deficient bone marrow stroma elicits persistent changes in HSC potential. <i>Nature Immunology</i> , 2020, 21, 261-273.	7.0	30
9	MZB1 enables efficient interferon γ secretion in stimulated plasmacytoid dendritic cells. <i>Scientific Reports</i> , 2020, 10, 21626.	1.6	12
10	Dynamic EBF1 occupancy directs sequential epigenetic and transcriptional events in B-cell programming. <i>Genes and Development</i> , 2018, 32, 96-111.	2.7	76
11	Defining B Cell Chromatin: Lessons from EBF1. <i>Trends in Genetics</i> , 2018, 34, 257-269.	2.9	35
12	Revisiting the role of IRF3 in inflammation and immunity by conditional and specifically targeted gene ablation in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 5253-5258.	3.3	77
13	Comprehensive Proteomic Investigation of <i>Ebf1</i> Heterozygosity in Pro-B Lymphocytes Utilizing Data Independent Acquisition. <i>Journal of Proteome Research</i> , 2018, 17, 76-85.	1.8	21
14	Active intermixing of indirect and direct neurons builds the striatal mosaic. <i>Nature Communications</i> , 2018, 9, 4725.	5.8	28
15	Cochaperone Mzb1 is a key effector of Blimp1 in plasma cell differentiation and β 2-integrin function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E9630-E9639.	3.3	52
16	Satb2 Is Required for the Development of a Spinal Exteroceptive Microcircuit that Modulates Limb Position. <i>Neuron</i> , 2016, 91, 763-776.	3.8	42
17	Enhancer decommissioning by Snail1-induced competitive displacement of TCF7L2 and down-regulation of transcriptional activators results in EPHB2 silencing. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016, 1859, 1353-1367.	0.9	18
18	Interaction of CCR4 α with EBF1 regulates gene-specific transcription and mRNA stability in B lymphopoiesis. <i>Genes and Development</i> , 2016, 30, 2310-2324.	2.7	29

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19	Pioneering Activity of the C-Terminal Domain of EBF1 Shapes the Chromatin Landscape for B Cell Programming. <i>Immunity</i> , 2016, 44, 527-541.	6.6	102
20	Satb2 Regulates the Differentiation of Both Callosal and Subcerebral Projection Neurons in the Developing Cerebral Cortex. <i>Cerebral Cortex</i> , 2015, 25, 3406-3419.	1.6	137
21	MZB1 is a GRP94 cochaperone that enables proper immunoglobulin heavy chain biosynthesis upon ER stress. <i>Genes and Development</i> , 2014, 28, 1165-1178.	2.7	95
22	Integrated genomic analysis identifies recurrent mutations and evolution patterns driving the initiation and progression of follicular lymphoma. <i>Nature Genetics</i> , 2014, 46, 176-181.	9.4	624
23	The regulatory network of B cell differentiation: a focused view of early B cell factor 1 function. <i>Immunological Reviews</i> , 2014, 261, 102-115.	2.8	113
24	Transcription factor EBF1 is essential for the maintenance of B cell identity and prevention of alternative fates in committed cells. <i>Nature Immunology</i> , 2013, 14, 867-875.	7.0	168
25	Establishment and Maintenance of B Cell Identity. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2013, 78, 23-30.	2.0	13
26	Transcription factor Ebf1 regulates differentiation stage-specific signaling, proliferation, and survival of B cells. <i>Genes and Development</i> , 2012, 26, 668-682.	2.7	134
27	Structure of an Ebf1:DNA complex reveals unusual DNA recognition and structural homology with Rel proteins. <i>Genes and Development</i> , 2010, 24, 2270-2275.	2.7	47
28	Early B Cell Factor 1 Regulates B Cell Gene Networks by Activation, Repression, and Transcription-Independent Poising of Chromatin. <i>Immunity</i> , 2010, 32, 714-725.	6.6	191
29	Transcription control of early B cell differentiation. <i>Current Opinion in Immunology</i> , 2010, 22, 161-167.	2.4	117
30	Mzb1 Protein Regulates Calcium Homeostasis, Antibody Secretion, and Integrin Activation in Innate-like B Cells. <i>Immunity</i> , 2010, 33, 723-735.	6.6	92
31	Early B Cell Factor 2 Regulates Hematopoietic Stem Cell Homeostasis in a Cell-Nonautonomous Manner. <i>Cell Stem Cell</i> , 2010, 7, 496-507.	5.2	44
32	Satb1 and Satb2 regulate embryonic stem cell differentiation and <i>Nanog</i> expression. <i>Genes and Development</i> , 2009, 23, 2625-2638.	2.7	125
33	Transcription factor EBF restricts alternative lineage options and promotes B cell fate commitment independently of Pax5. <i>Nature Immunology</i> , 2008, 9, 203-215.	7.0	215
34	Satb2 Regulates Callosal Projection Neuron Identity in the Developing Cerebral Cortex. <i>Neuron</i> , 2008, 57, 364-377.	3.8	581
35	Distinct Promoters Mediate the Regulation of Ebf1 Gene Expression by Interleukin-7 and Pax5. <i>Molecular and Cellular Biology</i> , 2007, 27, 579-594.	1.1	150
36	Dynamics and interplay of nuclear architecture, genome organization, and gene expression. <i>Genes and Development</i> , 2007, 21, 3027-3043.	2.7	358

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37	SATB2 Is a Multifunctional Determinant of Craniofacial Patterning and Osteoblast Differentiation. <i>Cell</i> , 2006, 125, 971-986.	13.5	458
38	Role of transcription factors in commitment and differentiation of early B lymphoid cells. <i>Seminars in Immunology</i> , 2006, 18, 12-19.	2.7	20
39	EBF2 Regulates Osteoblast-Dependent Differentiation of Osteoclasts. <i>Developmental Cell</i> , 2005, 9, 757-767.	3.1	107
40	Assembling a Gene Regulatory Network for Specification of the B Cell Fate. <i>Developmental Cell</i> , 2004, 7, 607-617.	3.1	212
41	SUMO modification of a novel MAR-binding protein, SATB2, modulates immunoglobulin λ gene expression. <i>Genes and Development</i> , 2003, 17, 3048-3061.	2.7	233
42	Identification of the regions involved in DNA binding by the mouse PEBP2 β protein. <i>FEBS Letters</i> , 2000, 470, 125-130.	1.3	15
43	Coordinate Regulation of B Cell Differentiation by the Transcription Factors EBF and E2A. <i>Immunity</i> , 1999, 11, 21-31.	6.6	293
44	EBF and E47 Collaborate to Induce Expression of the Endogenous Immunoglobulin Surrogate Light Chain Genes. <i>Immunity</i> , 1997, 7, 25-36.	6.6	247
45	Extension of chromatin accessibility by nuclear matrix attachment regions. <i>Nature</i> , 1997, 385, 269-272.	13.7	237
46	Functional interaction of β -catenin with the transcription factor LEF-1. <i>Nature</i> , 1996, 382, 638-642.	13.7	2,720
47	Failure of B-cell differentiation in mice lacking the transcription factor EBF. <i>Nature</i> , 1995, 376, 263-267.	13.7	603
48	Structural basis for DNA bending by the architectural transcription factor LEF-1. <i>Nature</i> , 1995, 376, 791-795.	13.7	582
49	Depletion of the predominant B-cell population in immunoglobulin μ heavy-chain transgenic mice. <i>Nature</i> , 1987, 329, 71-73.	13.7	101