

# Yurii Chinenov

## List of Publications by Year in descending order

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34  
papers

3,882  
citations

218677

26  
h-index

377865

34  
g-index

35  
all docs

35  
docs citations

35  
times ranked

6362  
citing authors

#	ARTICLE	IF	CITATIONS
1	In vitro responses to platelet-rich-plasma are associated with variable clinical outcomes in patients with knee osteoarthritis. <i>Scientific Reports</i> , 2021, 11, 11493.	3.3	12
2	Altered function and differentiation of age-associated B cells contribute to the female bias in lupus mice. <i>Nature Communications</i> , 2021, 12, 4813.	12.8	47
3	<scp>RNA</scp>â€œ<scp>seq</scp> Analysis of <scp>Periâ€œImplant</scp> Tissue Shows Differences in Immune, Notch, Wnt, and Angiogenesis Pathways in Aged Versus Young Mice. <i>JBMR Plus</i> , 2021, 5, e10535.	2.7	6
4	Transcription cofactor GRIP1 differentially affects myeloid cellâ€œdriven neuroinflammation and response to IFN- $\gamma$ therapy. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	4
5	Serine-threonine kinase ROCK2 regulates germinal center B cell positioning and cholesterol biosynthesis. <i>Journal of Clinical Investigation</i> , 2020, 130, 3654-3670.	8.2	26
6	Regulation of age-associated B cells by IRF5 in systemic autoimmunity. <i>Nature Immunology</i> , 2018, 19, 407-419.	14.5	105
7	Gene-specific mechanisms direct glucocorticoid-receptor-driven repression of inflammatory response genes in macrophages. <i>ELife</i> , 2018, 7, .	6.0	77
8	iRhom2 promotes lupus nephritis through TNF- $\alpha$ and EGFR signaling. <i>Journal of Clinical Investigation</i> , 2018, 128, 1397-1412.	8.2	66
9	Shifting diets and the rise of male-biased inequality on the Central Plains of China during Eastern Zhou. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 932-937.	7.1	69
10	Targeting the RhoA-ROCK pathway to reverse T-cell dysfunction in SLE. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 740-747.	0.9	73
11	The mTORC1-4E-BP-eIF4E axis controls de novo Bcl6 protein synthesis in T cells and systemic autoimmunity. <i>Nature Communications</i> , 2017, 8, 254.	12.8	46
12	Glucocorticoid-induced phosphorylation by CDK9 modulates the coactivator functions of transcriptional cofactor GRIP1 in macrophages. <i>Nature Communications</i> , 2017, 8, 1739.	12.8	28
13	The transcriptional coregulator GRIP1 controls macrophage polarization and metabolic homeostasis. <i>Nature Communications</i> , 2016, 7, 12254.	12.8	37
14	Regulation of Effector Treg Cells in Murine Lupus. <i>Arthritis and Rheumatology</i> , 2016, 68, 1454-1466.	5.6	15
15	Glucocorticoid Signaling: An Update from a Genomic Perspective. <i>Annual Review of Physiology</i> , 2016, 78, 155-180.	13.1	109
16	Glucocorticoid receptor coordinates transcription factor-dominated regulatory network in macrophages. <i>BMC Genomics</i> , 2014, 15, 656.	2.8	73
17	Nuclear receptors in inflammation control: Repression by GR and beyond. <i>Molecular and Cellular Endocrinology</i> , 2013, 380, 55-64.	3.2	56
18	Glucocorticoid receptor represses proinflammatory genes at distinct steps of the transcription cycle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 14616-14621.	7.1	55

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19	Glucocorticoid-Dependent Phosphorylation of the Transcriptional Coregulator GRIP1. <i>Molecular and Cellular Biology</i> , 2012, 32, 730-739.	2.3	26
20	Role of transcriptional coregulator GRIP1 in the anti-inflammatory actions of glucocorticoids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 11776-11781.	7.1	87
21	The Type I Interferon Signaling Pathway Is a Target for Glucocorticoid Inhibition. <i>Molecular and Cellular Biology</i> , 2010, 30, 4564-4574.	2.3	126
22	Immediate mediators of the inflammatory response are poised for gene activation through RNA polymerase II stalling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 18207-18212.	7.1	132
23	GRIP1-associated SET-domain methyltransferase in glucocorticoid receptor target gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 20185-20190.	7.1	26
24	Glucocorticoids and the innate immune system: Crosstalk with the Toll-like receptor signaling network. <i>Molecular and Cellular Endocrinology</i> , 2007, 275, 30-42.	3.2	109
25	The GRIP1:IRF3 interaction as a target for glucocorticoid receptor-mediated immunosuppression. <i>EMBO Journal</i> , 2006, 25, 108-117.	7.8	141
26	Comparative modeling of the N-terminal domain of the 67kDa laminin-binding protein: implications for putative ribosomal function. <i>Biochemical and Biophysical Research Communications</i> , 2003, 300, 161-166.	2.1	11
27	Visualization of Interactions among bZIP and Rel Family Proteins in Living Cells Using Bimolecular Fluorescence Complementation. <i>Molecular Cell</i> , 2002, 9, 789-798.	9.7	1,395
28	A second catalytic domain in the Elp3 histone acetyltransferases: a candidate for histone demethylase activity?. <i>Trends in Biochemical Sciences</i> , 2002, 27, 115-117.	7.5	75
29	Close encounters of many kinds: Fos-Jun interactions that mediate transcription regulatory specificity. <i>Oncogene</i> , 2001, 20, 2438-2452.	5.9	634
30	The $\hat{1}\pm$ and $\hat{1}^2$ Subunits of the GA-binding Protein Form a Stable Heterodimer in Solution. <i>Journal of Biological Chemistry</i> , 2000, 275, 7749-7756.	3.4	46
31	Isolation of a bi-directional promoter directing expression of the mouse GABP $\hat{1}\pm$ and ATP synthase coupling factor 6 genes. <i>Gene</i> , 2000, 261, 311-320.	2.2	14
32	Identification of Redox-sensitive Cysteines in GA-binding Protein- $\hat{1}\pm$ That Regulate DNA Binding and Heterodimerization. <i>Journal of Biological Chemistry</i> , 1998, 273, 6203-6209.	3.4	37
33	GA-binding Protein-dependent Transcription Initiator Elements. <i>Journal of Biological Chemistry</i> , 1997, 272, 29060-29067.	3.4	55
34	Redox Regulation of GA-binding Protein- $\hat{1}\pm$ DNA Binding Activity. <i>Journal of Biological Chemistry</i> , 1996, 271, 25617-25623.	3.4	63