

# Vigo Heissmeyer

## List of Publications by Year in descending order

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Version: 2024-02-01

41  
papers

4,262  
citations

218677

26  
h-index

276875

41  
g-index

43  
all docs

43  
docs citations

43  
times ranked

7353  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cooperation of RNA-Binding Proteins – a Focus on Roquin Function in T Cells. <i>Frontiers in Immunology</i> , 2022, 13, 839762.	4.8	4
2	Post-transcriptional control of T-cell development in the thymus. <i>Immunology Letters</i> , 2022, 247, 1-12.	2.5	3
3	Defining the RBPome of primary T helper cells to elucidate higher-order Roquin-mediated mRNA regulation. <i>Nature Communications</i> , 2021, 12, 5208.	12.8	23
4	Disrupting Roquin-1 interaction with Regnase-1 induces autoimmunity and enhances antitumor responses. <i>Nature Immunology</i> , 2021, 22, 1563-1576.	14.5	22
5	TRAF6 prevents fatal inflammation by homeostatic suppression of MALT1 protease. <i>Science Immunology</i> , 2021, 6, eabh2095.	11.9	17
6	Validation strategies for antibodies targeting modified ribonucleotides. <i>Rna</i> , 2020, 26, 1489-1506.	3.5	18
7	Elevated Exhaustion Levels of NK and CD8+ T Cells as Indicators for Progression and Prognosis of COVID-19 Disease. <i>Frontiers in Immunology</i> , 2020, 11, 580237.	4.8	96
8	Immune homeostasis and regulation of the interferon pathway require myeloid-derived Regnase-3. <i>Journal of Experimental Medicine</i> , 2019, 216, 1700-1723.	8.5	29
9	Production and Application of Stable Isotope-Labeled Internal Standards for RNA Modification Analysis. <i>Genes</i> , 2019, 10, 26.	2.4	38
10	A translational silencing function of MCPIP1/Regnase-1 specified by the target site context. <i>Nucleic Acids Research</i> , 2018, 46, 4256-4270.	14.5	20
11	Posttranscriptional regulation of T helper cell fate decisions. <i>Journal of Cell Biology</i> , 2018, 217, 2615-2631.	5.2	29
12	Binding of NUFIP2 to Roquin promotes recognition and regulation of ICOS mRNA. <i>Nature Communications</i> , 2018, 9, 299.	12.8	27
13	Roquin targets mRNAs in a 3' UTR-specific manner by different modes of regulation. <i>Nature Communications</i> , 2018, 9, 3810.	12.8	40
14	Posttranscriptional Gene Regulation of T Follicular Helper Cells by RNA-Binding Proteins and microRNAs. <i>Frontiers in Immunology</i> , 2018, 9, 1794.	4.8	17
15	Roquin Suppresses the PI3K-mTOR Signaling Pathway to Inhibit T Helper Cell Differentiation and Conversion of Treg to Tfr Cells. <i>Immunity</i> , 2017, 47, 1067-1082.e12.	14.3	109
16	Roquin recognizes a non-canonical hexaloop structure in the 3' UTR of Ox40. <i>Nature Communications</i> , 2016, 7, 11032.	12.8	38
17	Epstein-Barr viral miRNAs inhibit antiviral CD4+ T cell responses targeting IL-12 and peptide processing. <i>Journal of Experimental Medicine</i> , 2016, 213, 2065-2080.	8.5	108
18	Alternative splicing of MALT1 controls signalling and activation of CD4+ T cells. <i>Nature Communications</i> , 2016, 7, 11292.	12.8	94

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19	<scp>RNA</scp> recognition by Roquin in posttranscriptional gene regulation. Wiley Interdisciplinary Reviews RNA, 2016, 7, 455-469.	6.4	15
20	OX40L blockade protects against inflammation-driven fibrosis. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E3901-10.	7.1	50
21	InÂVivo Killing Capacity of Cytotoxic T Cells Is Limited and Involves Dynamic Interactions and T Cell Cooperativity. Immunity, 2016, 44, 233-245.	14.3	199
22	Regulation of T cell signaling and autoimmunity by RNA-binding proteins. Current Opinion in Immunology, 2016, 39, 127-135.	5.5	29
23	Roquin binds microRNA-146a and Argonaute2 to regulate microRNA homeostasis. Nature Communications, 2015, 6, 6253.	12.8	59
24	The Transcription Factor NFAT Promotes Exhaustion of Activated CD8 + T Cells. Immunity, 2015, 42, 265-278.	14.3	555
25	Induced miRâ€99a expression represses <i>Mtor</i> cooperatively with miRâ€150 to promote regulatory Tâ€cell differentiation. EMBO Journal, 2015, 34, 1195-1213.	7.8	83
26	Uncoupling Malt1 Threshold Function from Paracaspase Activity Results in Destructive Autoimmune Inflammation. Cell Reports, 2014, 9, 1292-1305.	6.4	133
27	Degradation of oligouridylated histone <scp>mRNAs</scp>: see <scp>UUUUU</scp> and goodbye. Wiley Interdisciplinary Reviews RNA, 2014, 5, 577-589.	6.4	23
28	Tfh Cell Differentiation: Missing Stat3 Uncovers Interferonsâ€™ Interference. Immunity, 2014, 40, 307-309.	14.3	3
29	Cleavage of roquin and regnase-1 by the paracaspase MALT1 releases their cooperatively repressed targets to promote TH17 differentiation. Nature Immunology, 2014, 15, 1079-1089.	14.5	238
30	Structural basis for RNA recognition in roquin-mediated post-transcriptional gene regulation. Nature Structural and Molecular Biology, 2014, 21, 671-678.	8.2	77
31	Roquin Paralogs 1 and 2 Redundantly Repress the Icos and Ox40 Costimulator mRNAs and Control Follicular Helper T Cell Differentiation. Immunity, 2013, 38, 655-668.	14.3	178
32	Adenoviral Transduction of Naive CD4 T Cells to Study Treg Differentiation. Journal of Visualized Experiments, 2013, , .	0.3	6
33	T cell activation induces proteasomal degradation of Argonaute and rapid remodeling of the microRNA repertoire. Journal of Experimental Medicine, 2013, 210, 417-432.	8.5	180
34	Molecular control of Tfhâ€cell differentiation by Roquin family proteins. Immunological Reviews, 2013, 253, 273-289.	6.0	42
35	Eri1 degrades the stem-loop of oligouridylated histone mRNAs to induce replication-dependent decay. Nature Structural and Molecular Biology, 2013, 20, 73-81.	8.2	68
36	Eri1 regulates microRNA homeostasis and mouse lymphocyte development and antiviral function. Blood, 2012, 120, 130-142.	1.4	61

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37	Roquin binds inducible costimulator mRNA and effectors of mRNA decay to induce microRNA-independent post-transcriptional repression. <i>Nature Immunology</i> , 2010, 11, 725-733.	14.5	159
38	Six RNA Viruses and Forty-One Hosts: Viral Small RNAs and Modulation of Small RNA Repertoires in Vertebrate and Invertebrate Systems. <i>PLoS Pathogens</i> , 2010, 6, e1000764.	4.7	234
39	MicroRNAs grow up in the immune system. <i>Current Opinion in Immunology</i> , 2008, 20, 281-287.	5.5	63
40	Mouse Eri1 interacts with the ribosome and catalyzes 5.8S rRNA processing. <i>Nature Structural and Molecular Biology</i> , 2008, 15, 523-530.	8.2	53
41	FOXP3 Controls Regulatory T Cell Function through Cooperation with NFAT. <i>Cell</i> , 2006, 126, 375-387.	28.9	1,019