

Irina Vlasova-St Louis

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

1,645
citations

471061

17
h-index

500791

28
g-index

35
all docs

35
docs citations

35
times ranked

2096
citing authors

#	ARTICLE	IF	CITATIONS
1	Three functional variants of IFN regulatory factor 5 (IRF5) define risk and protective haplotypes for human lupus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 6758-6763.	3.3	428
2	Conserved GU-Rich Elements Mediate mRNA Decay by Binding to CUG-Binding Protein 1. <i>Molecular Cell</i> , 2008, 29, 263-270.	4.5	216
3	Tristetraprolin Down-Regulates <i>IL-2</i> Gene Expression through AU-Rich Element-Mediated mRNA Decay. <i>Journal of Immunology</i> , 2005, 174, 953-961.	0.4	190
4	Analysis of CUGBP1 Targets Identifies GU-Repeat Sequences That Mediate Rapid mRNA Decay. <i>Molecular and Cellular Biology</i> , 2010, 30, 3970-3980.	1.1	120
5	Tristetraprolin Mediates Interferon- β mRNA Decay. <i>Journal of Biological Chemistry</i> , 2009, 284, 11216-11223.	1.6	109
6	Posttranscriptional regulation of gene networks by GU-rich elements and CELF proteins. <i>RNA Biology</i> , 2008, 5, 201-207.	1.5	81
7	Coordinate regulation of mRNA decay networks by GU-rich elements and CELF1. <i>Current Opinion in Genetics and Development</i> , 2011, 21, 444-451.	1.5	68
8	Patterns of coordinate down-regulation of ARE-containing transcripts following immune cell activation. <i>Genomics</i> , 2004, 84, 1002-1013.	1.3	57
9	Regulation of CUG-binding Protein 1 (CUGBP1) Binding to Target Transcripts upon T Cell Activation. <i>Journal of Biological Chemistry</i> , 2012, 287, 950-960.	1.6	49
10	CELFish ways to modulate mRNA decay. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013, 1829, 695-707.	0.9	43
11	Post-Transcriptional Regulation of Cytokine Signaling by AU-Rich and GU-Rich Elements. <i>Journal of Interferon and Cytokine Research</i> , 2014, 34, 233-241.	0.5	43
12	Global assessment of GU-rich regulatory content and function in the human transcriptome. <i>RNA Biology</i> , 2011, 8, 681-691.	1.5	39
13	Post-transcriptional regulation of cytokine and growth factor signaling in cancer. <i>Cytokine and Growth Factor Reviews</i> , 2017, 33, 83-93.	3.2	32
14	Coordinate stabilization of growth-regulatory transcripts in T cell malignancies. <i>Genomics</i> , 2005, 86, 159-171.	1.3	29
15	Antiretroviral Therapy Down-Regulates Innate Antiviral Response Genes in Patients With AIDS in Sub-Saharan Africa. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2010, 55, 428-438.	0.9	23
16	Transcriptomic Predictors of Paradoxical Cryptococcosis-Associated Immune Reconstitution Inflammatory Syndrome. <i>Open Forum Infectious Diseases</i> , 2018, 5, ofy157.	0.4	23
17	Feedback Regulation of Kinase Signaling Pathways by AREs and GREs. <i>Cells</i> , 2016, 5, 4.	1.8	17
18	Altered CELF1 binding to target transcripts in malignant T cells. <i>Rna</i> , 2015, 21, 1757-1769.	1.6	15

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19	Viral manipulation of host mRNA decay. <i>Future Virology</i> , 2018, 13, 211-223.	0.9	12
20	Cryptococcal Immune Reconstitution Inflammatory Syndrome: From Blood and Cerebrospinal Fluid Biomarkers to Treatment Approaches. <i>Life</i> , 2021, 11, 95.	1.1	11
21	The hepatitis C viral nonstructural protein 5A stabilizes growth-regulatory human transcripts. <i>Nucleic Acids Research</i> , 2018, 46, 2537-2547.	6.5	8
22	Immune Reconstitution Disorders: Spotlight on Interferons. <i>International Journal of Biomedical Investigation</i> , 2019, 2, 1-21.	0.7	8
23	Transcriptomic biomarker pathways associated with death in HIV-infected patients with cryptococcal meningitis. <i>BMC Medical Genomics</i> , 2021, 14, 108.	0.7	6
24	Post-transcriptional regulation of cytokine expression and signaling. <i>Current Trends in Immunology</i> , 2018, 19, 33-40.	4.0	5
25	Reovirus infection induces stabilization and up-regulation of cellular transcripts that encode regulators of TGF- β signaling. <i>PLoS ONE</i> , 2018, 13, e0204622.	1.1	4
26	Cytomegalovirus-Specific Immunity Recovers More Slowly after Cord Blood Transplantation Compared with Matched Sibling Donor Allogeneic Transplantation. <i>Transplantation and Cellular Therapy</i> , 2021, 27, 187.e1-187.e4.	0.6	4
27	Kinetics of immune reconstitution and immune complications after cell and organ transplantation. <i>Integrative Cancer Science and Therapeutics</i> , 2020, 7, .	0.1	2
28	Post-transcriptional Regulation of Cytokine Signaling During Inflammatory Responses. , 2016, , 55-70.		1
29	Mammalian Cis-Acting RNA Sequence Elements. , 2018, , .		0
30	Diagnostic Applications for RNA-Seq Technology and Transcriptome Analyses in Human Diseases Caused by RNA Viruses. , 0, , .		0
31	SÃndrome inflamatÃ³ria da reconstituiÃ§Ã£o imune associada Ã meninge criptococÃ³tica: fatores de risco e biomarcadores. <i>Arquivos De CiÃªncias Da SaÃde</i> , 2021, 28, .	0.3	0
32	Introductory Chapter: Applications of RNA-Seq Diagnostics in Biology and Medicine. , 0, , .		0