

Yuanjia Tang

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

4,475
citations

186209

28
h-index

289141

40
g-index

41
all docs

41
docs citations

41
times ranked

6153
citing authors

#	ARTICLE	IF	CITATIONS
1	MicroRNA-146a contributes to abnormal activation of the type I interferon pathway in human lupus by targeting the key signaling proteins. <i>Arthritis and Rheumatism</i> , 2009, 60, 1065-1075.	6.7	679
2	MicroRNA-21 and MicroRNA-148a Contribute to DNA Hypomethylation in Lupus CD4+ T Cells by Directly and Indirectly Targeting DNA Methyltransferase 1. <i>Journal of Immunology</i> , 2010, 184, 6773-6781.	0.4	499
3	The microRNA miR-23b suppresses IL-17-associated autoimmune inflammation by targeting TAB2, TAB3 and IKK- β . <i>Nature Medicine</i> , 2012, 18, 1077-1086.	15.2	397
4	Sex-specific association of X-linked Toll-like receptor 7 (TLR7) with male systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 15838-15843.	3.3	324
5	A Functional Variant in MicroRNA-146a Promoter Modulates Its Expression and Confers Disease Risk for Systemic Lupus Erythematosus. <i>PLoS Genetics</i> , 2011, 7, e1002128.	1.5	241
6	miR-155 and its star-form partner miR-155* cooperatively regulate type I interferon production by human plasmacytoid dendritic cells. <i>Blood</i> , 2010, 116, 5885-5894.	0.6	233
7	Identification of the long noncoding RNA NEAT1 as a novel inflammatory regulator acting through MAPK pathway in human lupus. <i>Journal of Autoimmunity</i> , 2016, 75, 96-104.	3.0	233
8	MicroRNA-125a contributes to elevated inflammatory chemokine RANTES levels via targeting KLF13 in systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2010, 62, 3425-3435.	6.7	212
9	MicroRNAs as novel regulators of systemic lupus erythematosus pathogenesis. <i>Nature Reviews Rheumatology</i> , 2012, 8, 701-709.	3.5	143
10	MiR-125a targets effector programs to stabilize Treg-mediated immune homeostasis. <i>Nature Communications</i> , 2015, 6, 7096.	5.8	133
11	miR-132 regulates the differentiation of dopamine neurons by directly targeting Nurr1 expression. <i>Journal of Cell Science</i> , 2012, 125, 1673-82.	1.2	132
12	Identification of 38 novel loci for systemic lupus erythematosus and genetic heterogeneity between ancestral groups. <i>Nature Communications</i> , 2021, 12, 772.	5.8	128
13	Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 632-640.	0.5	103
14	Identification of microRNA-31 as a novel regulator contributing to impaired interleukin-2 production in T cells from patients with systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 3715-3725.	6.7	97
15	Association of large intergenic noncoding RNA expression with disease activity and organ damage in systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2015, 17, 131.	1.6	92
16	The role of long non-coding RNAs in rheumatic diseases. <i>Nature Reviews Rheumatology</i> , 2017, 13, 657-669.	3.5	65
17	MicroRNA-130b Ameliorates Murine Lupus Nephritis Through Targeting the Type I Interferon Pathway on Renal Mesangial Cells. <i>Arthritis and Rheumatology</i> , 2016, 68, 2232-2243.	2.9	59
18	T-bet+CD11c+ B cells are critical for antichromatin immunoglobulin G production in the development of lupus. <i>Arthritis Research and Therapy</i> , 2017, 19, 225.	1.6	58

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19	In Vivo Therapeutic Success of MicroRNA-155 Antagomir in a Mouse Model of Lupus Alveolar Hemorrhage. <i>Arthritis and Rheumatology</i> , 2016, 68, 953-964.	2.9	57
20	MiR-125a-5p ameliorates monocrotaline-induced pulmonary arterial hypertension by targeting the TGF- β 1 and IL-6/STAT3 signaling pathways. <i>Experimental and Molecular Medicine</i> , 2018, 50, 1-11.	3.2	56
21	Identification of LncRNA Linc00513 Containing Lupus-Associated Genetic Variants as a Novel Regulator of Interferon Signaling Pathway. <i>Frontiers in Immunology</i> , 2018, 9, 2967.	2.2	56
22	MicroRNA-125a-Loaded Polymeric Nanoparticles Alleviate Systemic Lupus Erythematosus by Restoring Effector/Regulatory T Cells Balance. <i>ACS Nano</i> , 2020, 14, 4414-4429.	7.3	53
23	Type I Interferon Inhibition of MicroRNA-146a Maturation Through Up-Regulation of Monocyte Chemotactic Protein-1 Induced Protein 1 in Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2015, 67, 3209-3218.	2.9	51
24	Exome-wide association study identifies four novel loci for systemic lupus erythematosus in Han Chinese population. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 417-417.	0.5	50
25	SLE non-coding genetic risk variant determines the epigenetic dysfunction of an immune cell specific enhancer that controls disease-critical microRNA expression. <i>Nature Communications</i> , 2021, 12, 135.	5.8	48
26	Zirconia Hybrid Nanoshells for Nutrient and Toxin Detection. <i>Small</i> , 2020, 16, e2003902.	5.2	37
27	Identification of Cyclin-Dependent Kinase 1 as a Novel Regulator of Type I Interferon Signaling in Systemic Lupus Erythematosus. <i>Arthritis and Rheumatology</i> , 2016, 68, 1222-1232.	2.9	35
28	Enhanced transfection of polyplexes based on pluronic-polypropylenimine dendrimer for gene transfer. <i>Archives of Pharmacal Research</i> , 2009, 32, 1045-1054.	2.7	28
29	MicroRNA-125b/Lin28 Pathway Contributes to the Mesendodermal Fate Decision of Embryonic Stem Cells. <i>Stem Cells and Development</i> , 2012, 21, 1524-1537.	1.1	25
30	miR-744 enhances type I interferon signaling pathway by targeting PTP1B in primary human renal mesangial cells. <i>Scientific Reports</i> , 2015, 5, 12987.	1.6	23
31	MiR-125a Is a critical modulator for neutrophil development. <i>PLoS Genetics</i> , 2017, 13, e1007027.	1.5	19
32	Genetic polymorphism in the 3' untranslated region of the E-cadherin gene is associated with risk of different cancers. <i>Molecular Carcinogenesis</i> , 2011, 50, 857-862.	1.3	17
33	SARS-CoV-2-Encoded MiRNAs Inhibit Host Type I Interferon Pathway and Mediate Allelic Differential Expression of Susceptible Gene. <i>Frontiers in Immunology</i> , 2021, 12, 767726.	2.2	17
34	Lupus enhancer risk variant causes dysregulation of IRF8 through cooperative lncRNA and DNA methylation machinery. <i>Nature Communications</i> , 2022, 13, 1855.	5.8	16
35	Interferon- γ exacerbates neuropsychiatric phenotypes in lupus-prone mice. <i>Arthritis Research and Therapy</i> , 2019, 21, 205.	1.6	14
36	Paradoxical effects of very low dose MK-801. <i>European Journal of Pharmacology</i> , 2006, 537, 77-84.	1.7	12

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37	miR-152 Attenuates the Severity of Lupus Nephritis Through the Downregulation of Macrophage Migration Inhibitory Factor (MIF)-Induced Expression of COL1A1. <i>Frontiers in Immunology</i> , 2019, 10, 158.	2.2	12
38	A Novel Vector-Based Method for Exclusive Overexpression of Star-Form MicroRNAs. <i>PLoS ONE</i> , 2012, 7, e41504.	1.1	9
39	Protective Role of microRNA-31 in Acetaminophen-Induced Liver Injury: A Negative Regulator of c-Jun N-Terminal Kinase (JNK) Signaling Pathway. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2021, 12, 1789-1807.	2.3	6
40	The MicroRNA <i>miR-22</i> Represses Th17 Cell Pathogenicity by Targeting PTEN-Regulated Pathways. <i>ImmunoHorizons</i> , 2020, 4, 308-318.	0.8	6
41	Epigenetics of Lupus. , 2019, , 69-85.		0