

# Yaser Mansoori

## List of Publications by Year in descending order

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Version: 2024-02-01

44  
papers

610  
citations

687363

13  
h-index

642732

23  
g-index

47  
all docs

47  
docs citations

47  
times ranked

801  
citing authors

#	ARTICLE	IF	CITATIONS
1	Competing endogenous RNA (ceRNA) cross talk and language in ceRNA regulatory networks: A new look at hallmarks of breast cancer. <i>Journal of Cellular Physiology</i> , 2019, 234, 10080-10100.	4.1	208
2	Expression pattern of miR-21, miR-25 and PTEN in peripheral blood mononuclear cells of patients with significant or insignificant coronary stenosis. <i>Gene</i> , 2019, 698, 170-178.	2.2	34
3	miRNA Polymorphisms and Risk of Cardio-Cerebrovascular Diseases: A Systematic Review and Meta-Analysis. <i>International Journal of Molecular Sciences</i> , 2019, 20, 293.	4.1	32
4	miRNA-Related Polymorphisms in miR-423 (rs6505162) and <i>PEX6</i> (rs1129186) and Risk of Esophageal Squamous Cell Carcinoma in an Iranian Cohort. <i>Genetic Testing and Molecular Biomarkers</i> , 2017, 21, 382-390.	0.7	26
5	Association between polymorphism of GSTP1, GSTT1, GSTM1 and CYP2E1 genes and susceptibility to benzene-induced hematotoxicity. <i>Archives of Toxicology</i> , 2018, 92, 1983-1990.	4.2	21
6	Dysregulated expression of STAT1, miR-150, and miR-223 in peripheral blood mononuclear cells of coronary artery disease patients with significant or insignificant stenosis. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 19810-19824.	2.6	20
7	Epigenetic Changes of the <i>ESR1</i> Gene in Breast Tissue of Healthy Women: A Missing Link with Breast Cancer Risk Factors?. <i>Genetic Testing and Molecular Biomarkers</i> , 2017, 21, 464-470.	0.7	17
8	Expression levels of breast cancer-related GAS5 and LSINCT5 lncRNAs in cancer-free breast tissue: Molecular associations with age at menarche and obesity. <i>Breast Journal</i> , 2018, 24, 876-882.	1.0	17
9	The intricate role of miR-155 in carcinogenesis: potential implications for esophageal cancer research. <i>Biomarkers in Medicine</i> , 2019, 13, 147-159.	1.4	16
10	Circular RNA hsa_circ_0044234 as distinct molecular signature of triple negative breast cancer: a potential regulator of GATA3. <i>Cancer Cell International</i> , 2021, 21, 312.	4.1	16
11	Integrative analyses of triple negative dysregulated transcripts compared with non-triple negative tumors and their functional and molecular interactions. <i>Journal of Cellular Physiology</i> , 2019, 234, 22386-22399.	4.1	15
12	Significance of a common variant in the CDKAL1 gene with susceptibility to type 2 diabetes mellitus in Iranian population. <i>Advanced Biomedical Research</i> , 2015, 4, 45.	0.5	15
13	Individual genetic variability mainly of Proinflammatory cytokines, cytokine receptors, and toll-like receptors dictates pathophysiology of COVID-19 disease. <i>Journal of Medical Virology</i> , 2022, 94, 4088-4096.	5.0	15
14	Association of VDR gene polymorphisms with risk of relapsing-remitting multiple sclerosis in an Iranian Kurdish population. <i>International Journal of Neuroscience</i> , 2018, 128, 505-511.	1.6	14
15	Circular RNA-associated ceRNA network involved in HIF-1 signalling in triple-negative breast cancer: circ_0047303 as a potential key regulator. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 11322-11332.	3.6	13
16	Association of interleukin-18 gene variants with susceptibility to visceral leishmaniasis in Iranian population. <i>Molecular Biology Reports</i> , 2013, 40, 4009-4014.	2.3	11
17	Evidences from a Systematic Review and Meta-Analysis Unveil the Role of MiRNA Polymorphisms in the Predisposition to Female Neoplasms. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5088.	4.1	9
18	Epi-miRNAs: Regulators of the Histone Modification Machinery in Human Cancer. <i>Journal of Oncology</i> , 2022, 2022, 1-22.	1.3	9

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19	Effects of Genetic Polymorphism on Susceptibility to Nephrotoxic Properties of BTEXs Compounds. <i>Journal of Occupational and Environmental Medicine</i> , 2018, 60, e377-e382.	1.7	8
20	Expression and clinicopathological significance of AOC4P, PRNCR1, and PCAT1 lncRNAs in breast cancer. <i>Pathology Research and Practice</i> , 2020, 216, 153131.	2.3	8
21	Expression profiles and functional prediction of long non-coding RNAs LINC01133, ZEB1-AS1 and ABHD11-AS1 in the luminal subtype of breast cancer. <i>Journal of Translational Medicine</i> , 2021, 19, 364.	4.4	8
22	Investigation of BRCAness associated miRNA-gene axes in breast cancer: cell-free miR-182-5p as a potential expression signature of BRCAness. <i>BMC Cancer</i> , 2022, 22, .	2.6	8
23	The HHEX rs1111875A/G gene polymorphism is associated with susceptibility to type 2 diabetes in the Iranian population. <i>Molecular Biology</i> , 2015, 49, 535-542.	1.3	7
24	Influences of L-1b-3953 C>T and MMP-9-1562C<math>\to>T</math> Gene Variants on Myocardial Infarction Susceptibility in a Subset of the Iranian Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2017, 21, 33-38.	0.7	7
25	Breast cancer-linked lncRNA u&Eleanor is upregulated in breast of healthy women with lack or short duration of breastfeeding. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 9869-9876.	2.6	7
26	Peripheral Blood Mononuclear Cells Expression Levels of miR-196a and miR-100 in Coronary Artery Disease Patients. <i>Immunological Investigations</i> , 2021, 50, 914-924.	2.0	7
27	Critical roles of microRNA-196 in normal physiology and non-malignant diseases: Diagnostic and therapeutic implications. <i>Experimental and Molecular Pathology</i> , 2021, 122, 104664.	2.1	6
28	Methylation of progesterone receptor isoform A promoter in normal breast tissue: An epigenetic link between early age at menarche and risk of breast cancer?. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 12393-12401.	2.6	4
29	The Effect of Mesenchymal Stem Cells on the Expression of IDO and Qa2 Molecules in Dendritic Cells. <i>Advanced Pharmaceutical Bulletin</i> , 2019, 9, 56-63.	1.4	4
30	Potential roles of hsa_circ_000839 and hsa_circ_0005986 in breast cancer. <i>Journal of Clinical Laboratory Analysis</i> , 2022, 36, e24263.	2.1	4
31	Re. <i>Journal of Occupational and Environmental Medicine</i> , 2018, 60, e560-e561.	1.7	3
32	Faculty retention in regional medical schools in Iran: a qualitative content analysis. <i>BMC Medical Education</i> , 2021, 21, 24.	2.4	3
33	Dysregulated Expression of miR-146a and Its Associated Immune Effectors in Peripheral Blood Mononuclear Cells of Esophageal Carcinoma Patients. <i>Immunological Investigations</i> , 2020, , 1-11.	2.0	2
34	A methylation signature at the CpG island promoter of estrogen receptor beta (ER- $\beta$ ) in breasts of women may be an early footmark of lack of breastfeeding and nulliparity. <i>Pathology Research and Practice</i> , 2021, 218, 153328.	2.3	2
35	Expression signature of lncRNA APTR in clinicopathology of breast cancer: Its potential oncogenic function in dysregulation of ErbB signaling pathway. <i>Gene Reports</i> , 2021, 23, 101116.	0.8	2
36	lncRNA&miRNA&mRNA Networks of Gastrointestinal Cancers Representing Common and Specific lncRNAs and mRNAs. <i>Frontiers in Genetics</i> , 2021, 12, 791919.	2.3	2

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37	Potential Roles of Long Noncoding RNAs as Therapeutic Targets in Organ Transplantation. <i>Frontiers in Immunology</i> , 2022, 13, 835746.	4.8	2
38	A link between expression level of long-non-coding RNA ZFAS1 in breast tissue of healthy women and obesity. <i>International Journal of Biological Markers</i> , 2018, 33, 500-506.	1.8	1
39	Genetic polymorphisms of base excision repair gene XRCC1 and susceptibility to benzene among employees of chemical industries. <i>Gene Reports</i> , 2021, 23, 101081.	0.8	1
40	Perturbation of miR-146b and relevant inflammatory elements in esophageal carcinoma patients supports an immune downregulatory mechanism. <i>Pathology Research and Practice</i> , 2021, 225, 153560.	2.3	1
41	A Proposed TUSC7/miR-211/Nurr1 ceRNET Might Potentially be Disturbed by a cer-SNP rs2615499 in Breast Cancer. <i>Biochemical Genetics</i> , 2022, 60, 2200-2225.	1.7	1
42	Genetic variations in <i>ATM</i> and <i>H2AX</i> loci contribute to risk of hematological abnormalities in individuals exposed to BTEX chemicals. <i>Journal of Clinical Laboratory Analysis</i> , 2022, 36, e24321.	2.1	1
43	The SDF1 A/G Gene Variant: A Susceptibility Variant for Myocardial Infarction. <i>Genetic Testing and Molecular Biomarkers</i> , 2017, 21, 506-511.	0.7	0
44	Experimental and Bioinformatic Clues to the Potential Roles of hsa_circ_0013958 and hsa_circ_0003028 in Clinopathophysiology of Breast Cancer. <i>Galen</i> , 0, 10, e2064.	0.6	0