

# Mohammadreza Hajjari

## List of Publications by Year in descending order

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Version: 2024-02-01

38  
papers

825  
citations

759055

12  
h-index

501076

28  
g-index

38  
all docs

38  
docs citations

38  
times ranked

1567  
citing authors

#	ARTICLE	IF	CITATIONS
1	HOTAIR: an oncogenic long non-coding RNA in different cancers. <i>Cancer Biology and Medicine</i> , 2015, 12, 1-9.	1.4	360
2	Up-regulation of HOTAIR long non-coding RNA in human gastric adenocarcinoma tissues. <i>Medical Oncology</i> , 2013, 30, 670.	1.2	100
3	Molecular function and regulation of long non-coding RNAs: paradigms with potential roles in cancer. <i>Tumor Biology</i> , 2014, 35, 10645-10663.	0.8	48
4	Long Non-Coding RNA SNHG6 as a Potential Biomarker for Hepatocellular Carcinoma. <i>Pathology and Oncology Research</i> , 2018, 24, 329-337.	0.9	42
5	Downregulation of miR-130a, antagonized doxorubicin-induced cardiotoxicity via increasing the PPAR $\alpha$ expression in mESCs-derived cardiac cells. <i>Cell Death and Disease</i> , 2018, 9, 758.	2.7	31
6	miR-485-3p suppresses colorectal cancer via targeting TPX2. <i>Bratislava Medical Journal</i> , 2020, 121, 302-307.	0.4	23
7	Junctional adhesion molecules 2 and 3 may potentially be involved in progression of gastric adenocarcinoma tumors. <i>Medical Oncology</i> , 2013, 30, 380.	1.2	22
8	HOTAIR: A Promising Long Non-coding RNA with Potential Role in Breast Invasive Carcinoma. <i>Frontiers in Genetics</i> , 2017, 8, 170.	1.1	22
9	Association Between SNPs of Long Non-coding RNA HOTAIR and Risk of Different Cancers. <i>Frontiers in Genetics</i> , 2019, 10, 113.	1.1	15
10	Circulating HOTAIR LncRNA Is Potentially Up-regulated in Coronary Artery Disease. <i>Genomics and Informatics</i> , 2018, 16, e25.	0.4	15
11	Long non-coding RNAs in hematologic malignancies: road to translational research. <i>Frontiers in Genetics</i> , 2013, 4, 250.	1.1	14
12	Long non-coding RNAs expression levels in diffuse large B-cell lymphoma: An in silico analysis. <i>Pathology Research and Practice</i> , 2018, 214, 1462-1466.	1.0	13
13	Potential long non-coding RNAs to be considered as biomarkers or therapeutic targets in gastric cancer. <i>Frontiers in Genetics</i> , 2013, 4, 210.	1.1	12
14	HOTAIR Long Non-coding RNA: Characterizing the Locus Features by the In Silico Approaches. <i>Genomics and Informatics</i> , 2017, 15, 170-177.	0.4	12
15	Long Noncoding RNAs in Colorectal Adenocarcinoma; an in silico Analysis. <i>Pathology and Oncology Research</i> , 2019, 25, 1387-1394.	0.9	11
16	A novel infram deletion in MSH6 gene in glioma: Conversation on MSH6 mutations in brain tumors. <i>Journal of Cellular Physiology</i> , 2019, 234, 11092-11102.	2.0	10
17	The potential role of PHF6 as an oncogene: a genotranscriptomic/proteomic meta-analysis. <i>Tumor Biology</i> , 2016, 37, 5317-5325.	0.8	9
18	Identification of a novel mutation in ARSA gene in three patients of an Iranian family with metachromatic leukodystrophy disorder. <i>Genetics and Molecular Biology</i> , 2017, 40, 759-762.	0.6	8

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19	Whole exome sequencing identified a novel nonsense INPP4A mutation in a family with intellectual disability. <i>European Journal of Medical Genetics</i> , 2020, 63, 103846.	0.7	8
20	Exome sequencing found a novel homozygous deletion in ADCK3 gene involved in autosomal recessive spinocerebellar ataxia. <i>Gene</i> , 2019, 708, 10-13.	1.0	6
21	Characterizing the Retinoblastoma 1 locus: putative elements for Rb1 regulation by in silico analysis. <i>Frontiers in Genetics</i> , 2014, 5, 2.	1.1	5
22	SNHG1 Long Noncoding RNA is Potentially Up-Regulated in Colorectal Adenocarcinoma. <i>Asian Pacific Journal of Cancer Prevention</i> , 2020, 21, 897-901.	0.5	5
23	Compositional features are potentially involved in the regulation of gene expression of tumor suppressor genes in human tissues. <i>Gene</i> , 2014, 553, 126-129.	1.0	4
24	Cloning and over expression of non-coding RNA rprA in E. coli and its resistance to Kanamycin without osmotic shock. <i>Bioinformation</i> , 2017, 13, 21-24.	0.2	4
25	HOTAIR Induces the Downregulation of miR-200 Family Members in Gastric Cancer Cell Lines. <i>Iranian Biomedical Journal</i> , 2022, 26, 77-84.	0.4	4
26	Editorial: Molecular Function and Regulation of Non-coding RNAs in Multifactorial Diseases. <i>Frontiers in Genetics</i> , 2016, 7, 9.	1.1	3
27	Exome sequencing revealed a novel deletion in the ERCC8 gene in an Iranian family with Cockayne syndrome. <i>Annals of Human Genetics</i> , 2018, 82, 304-308.	0.3	3
28	The non-coding RNA rprA can increase the resistance to ampicillin in Escherichia coli. <i>Microbial Pathogenesis</i> , 2019, 129, 266-270.	1.3	3
29	Up-Regulation of and Is Associated with The Progression of Gastric-Type Adenocarcinoma. <i>Cell Journal</i> , 2017, 19, 66-71.	0.2	3
30	SNHG7 and FAIM2 are up-regulated and co-expressed in colorectal adenocarcinoma tissues. <i>Klinicka Onkologie</i> , 2020, 33, .	0.1	3
31	In silico finding of Putative Cis-Acting Elements for the Tethering of Polycomb Repressive Complex2 in Human Genome. <i>Bioinformation</i> , 2014, 10, 187-190.	0.2	3
32	SNHG7 has an oncogenic role in colorectal cancer via potential sponging of MIR-485-5P and MIR-193A-5P; in silico approach. <i>Genetika</i> , 2021, 53, 65-78.	0.1	2
33	The construction of a short gene by a very fast, modified, and simplified gene synthesis and the analysis of various effects on this synthesis. <i>Brazilian Archives of Biology and Technology</i> , 2011, 54, 53-60.	0.5	1
34	Exome sequencing revealed a p.G299R mutation in the COMP gene in an Iranian family suffering from pseudoachondroplasia. <i>Journal of Gene Medicine</i> , 2019, 21, e3103.	1.4	1
35	Translational selection on SHH genes. <i>Genetics and Molecular Biology</i> , 2010, 33, 408-410.	0.6	0
36	Identification of the mutation p.S867P in the PTPRQ gene in an Iranian family with hearing impairment. <i>Meta Gene</i> , 2017, 13, 48-49.	0.3	0

#	ARTICLE	IF	CITATIONS
37	Tissue Specific Expression Levels of Apoptosis Involved Genes Have Correlations with Codon and Amino Acid Usage. <i>Genomics and Informatics</i> , 2016, 14, 234.	0.4	0
38	Whole exome sequencing revealed a novel dystrophin-related protein-2 () deletion in an Iranian family with symptoms of polyneuropathy. <i>Iranian Journal of Basic Medical Sciences</i> , 2019, 22, 576-580.	1.0	0