

# Mohammadreza Hajjari

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

36  
papers

634  
citations

11  
h-index

25  
g-index

38  
ext. papers

732  
ext. citations

2.9  
avg, IF

4.6  
L-index

#	Paper	IF	Citations
36	HOTAIR Induces the Downregulation of miR-200 Family Members in Gastric Cancer Cell Lines. <i>Iranian Biomedical Journal</i> , <b>2022</b> , 26, 77-84	2	1
35	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> , <b>2021</b> , 53, 65-78	0.6	0
34	Whole exome sequencing identified a novel nonsense INPP4A mutation in a family with intellectual disability. <i>European Journal of Medical Genetics</i> , <b>2020</b> , 63, 103846	2.6	0
33	SNHG1 Long Noncoding RNA is Potentially Up-Regulated in Colorectal Adenocarcinoma. <i>Asian Pacific Journal of Cancer Prevention</i> , <b>2020</b> , 21, 897-901	1.7	2
32	miR-485-3p suppresses colorectal cancer via targeting TPX2. <i>Bratislava Medical Journal</i> , <b>2020</b> , 121, 302-307		12
31	Association Between SNPs of Long Non-coding RNA and Risk of Different Cancers. <i>Frontiers in Genetics</i> , <b>2019</b> , 10, 113	4.5	11
30	Exome sequencing revealed a p.G299R mutation in the COMP gene in an Iranian family suffering from pseudoachondroplasia. <i>Journal of Gene Medicine</i> , <b>2019</b> , 21, e3103	3.5	1
29	Exome sequencing found a novel homozygous deletion in ADCK3 gene involved in autosomal recessive spinocerebellar ataxia. <i>Gene</i> , <b>2019</b> , 708, 10-13	3.8	3
28	The non-coding RNA rprA can increase the resistance to ampicillin in Escherichia coli. <i>Microbial Pathogenesis</i> , <b>2019</b> , 129, 266-270	3.8	1
27	Long Noncoding RNAs in Colorectal Adenocarcinoma; an in silico Analysis. <i>Pathology and Oncology Research</i> , <b>2019</b> , 25, 1387-1394	2.6	6
26	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> , <b>2019</b> , 22, 576-580	1.8	
25	A novel inframe deletion in MSH6 gene in glioma: Conversation on MSH6 mutations in brain tumors. <i>Journal of Cellular Physiology</i> , <b>2019</b> , 234, 11092-11102	7	8
24	Long Non-Coding RNA SNHG6 as a Potential Biomarker for Hepatocellular Carcinoma. <i>Pathology and Oncology Research</i> , <b>2018</b> , 24, 329-337	2.6	33
23	Exome sequencing revealed a novel deletion in the ERCC8 gene in an Iranian family with Cockayne syndrome. <i>Annals of Human Genetics</i> , <b>2018</b> , 82, 304-308	2.2	1
22	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> , <b>2018</b> , 9, 758	9.8	21
21	Long non-coding RNAs expression levels in diffuse large B-cell lymphoma: An in silico analysis. <i>Pathology Research and Practice</i> , <b>2018</b> , 214, 1462-1466	3.4	11
20	Circulating HOTAIR RNA Is Potentially Up-regulated in Coronary Artery Disease. <i>Genomics and Informatics</i> , <b>2018</b> , 16, e25	1.9	12

19	Identification of the mutation p.S867P in the PTPRQ gene in an Iranian family with hearing impairment. <i>Meta Gene</i> , <b>2017</b> , 13, 48-49	0.7	
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> , <b>2017</b> , 40, 759-762	2	5
17	: A Promising Long Non-coding RNA with Potential Role in Breast Invasive Carcinoma. <i>Frontiers in Genetics</i> , <b>2017</b> , 8, 170	4.5	17
16	Up-Regulation of and Is Associated with The Progression of Gastric-Type Adenocarcinoma. <i>Cell Journal</i> , <b>2017</b> , 19, 66-71	2.4	3
15	Long Non-coding RNA: Characterizing the Locus Features by the Approaches. <i>Genomics and Informatics</i> , <b>2017</b> , 15, 170-177	1.9	6
14	Cloning and over expression of non-coding RNA rprA in E.coli and its resistance to Kanamycin without osmotic shock. <i>Bioinformation</i> , <b>2017</b> , 13, 21-24	1.1	4
13	The potential role of PHF6 as an oncogene: a genotranscriptomic/proteomic meta-analysis. <i>Tumor Biology</i> , <b>2016</b> , 37, 5317-25	2.9	6
12	Tissue Specific Expression Levels of Apoptosis Involved Genes Have Correlations with Codon and Amino Acid Usage. <i>Genomics and Informatics</i> , <b>2016</b> , 14, 234-240	1.9	
11	HOTAIR: an oncogenic long non-coding RNA in different cancers. <i>Cancer Biology and Medicine</i> , <b>2015</b> , 12, 1-9	5.2	287
10	Compositional features are potentially involved in the regulation of gene expression of tumor suppressor genes in human tissues. <i>Gene</i> , <b>2014</b> , 553, 126-9	3.8	4
9	Molecular function and regulation of long non-coding RNAs: paradigms with potential roles in cancer. <i>Tumor Biology</i> , <b>2014</b> , 35, 10645-63	2.9	42
8	Characterizing the Retinoblastoma 1 locus: putative elements for Rb1 regulation by in silico analysis. <i>Frontiers in Genetics</i> , <b>2014</b> , 5, 2	4.5	3
7	In silico finding of Putative Cis-Acting Elements for the Tethering of Polycomb Repressive Complex2 in Human Genome. <i>Bioinformation</i> , <b>2014</b> , 10, 187-90	1.1	3
6	Up-regulation of HOTAIR long non-coding RNA in human gastric adenocarcinoma tissues. <i>Medical Oncology</i> , <b>2013</b> , 30, 670	3.7	87
5	Junctional adhesion molecules 2 and 3 may potentially be involved in progression of gastric adenocarcinoma tumors. <i>Medical Oncology</i> , <b>2013</b> , 30, 380	3.7	17
4	Potential long non-coding RNAs to be considered as biomarkers or therapeutic targets in gastric cancer. <i>Frontiers in Genetics</i> , <b>2013</b> , 4, 210	4.5	12
3	Long non-coding RNAs in hematologic malignancies: road to translational research. <i>Frontiers in Genetics</i> , <b>2013</b> , 4, 250	4.5	10
2	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> , <b>2011</b> , 54, 53-60	1.8	1

1 Translational selection on SHH genes. *Genetics and Molecular Biology*, **2010**, 33, 408-10

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