Mohammadreza Hajjari

List of Publications by Citations

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: an oncogenic long non-coding RNA in different cancers. *Cancer Biology and Medicine*, **2015**, 12, 1-9	5.2	287
35	Up-regulation of HOTAIR long non-coding RNA in human gastric adenocarcinoma tissues. *Medical Oncology*, **2013**, 30, 670	3.7	87
34	Molecular function and regulation of long non-coding RNAs: paradigms with potential roles in cancer. *Tumor Biology*, **2014**, 35, 10645-63	2.9	42
33	Long Non-Coding RNA SNHG6 as a Potential Biomarker for Hepatocellular Carcinoma. *Pathology and Oncology Research*, **2018**, 24, 329-337	2.6	33
32	Downregulation of miR-130a, antagonized doxorubicin-induced cardiotoxicity via increasing the PPAR expression in mESCs-derived cardiac cells. *Cell Death and Disease*, **2018**, 9, 758	9.8	21
31	Junctional adhesion molecules 2 and 3 may potentially be involved in progression of gastric adenocarcinoma tumors. *Medical Oncology*, **2013**, 30, 380	3.7	17
30	: A Promising Long Non-coding RNA with Potential Role in Breast Invasive Carcinoma. *Frontiers in Genetics*, **2017**, 8, 170	4.5	17
29	Potential long non-coding RNAs to be considered as biomarkers or therapeutic targets in gastric cancer. *Frontiers in Genetics*, **2013**, 4, 210	4.5	12
28	miR-485-3p suppresses colorectal cancer via targeting TPX2. *Bratislava Medical Journal*, **2020**, 121, 302-307	1.7	12
27	Circulating HOTAIR RNA Is Potentially Up-regulated in Coronary Artery Disease. *Genomics and Informatics*, **2018**, 16, e25	1.9	12
26	Association Between SNPs of Long Non-coding RNA and Risk of Different Cancers. *Frontiers in Genetics*, **2019**, 10, 113	4.5	11
25	Long non-coding RNAs expression levels in diffuse large B-cell lymphoma: An in silico analysis. *Pathology Research and Practice*, **2018**, 214, 1462-1466	3.4	11
24	Long non-coding RNAs in hematologic malignancies: road to translational research. *Frontiers in Genetics*, **2013**, 4, 250	4.5	10
23	A novel infram deletion in MSH6 gene in glioma: Conversation on MSH6 mutations in brain tumors. *Journal of Cellular Physiology*, **2019**, 234, 11092-11102	7	8
22	The potential role of PHF6 as an oncogene: a genotranscriptomic/proteomic meta-analysis. *Tumor Biology*, **2016**, 37, 5317-25	2.9	6
21	Long Noncoding RNAs in Colorectal Adenocarcinoma; an in silico Analysis. *Pathology and Oncology Research*, **2019**, 25, 1387-1394	2.6	6
20	Long Non-coding RNA: Characterizing the Locus Features by the Approaches. *Genomics and Informatics*, **2017**, 15, 170-177	1.9	6

19	Identification of a novel mutation in ARSA gene in three patients of an Iranian family with metachromatic leukodystrophy disorder. *Genetics and Molecular Biology*, **2017**, 40, 759-762	2	5
18	Compositional features are potentially involved in the regulation of gene expression of tumor suppressor genes in human tissues. *Gene*, **2014**, 553, 126-9	3.8	4
17	Cloning and over expression of non-coding RNA rprA in E.coli and its resistance to Kanamycin without osmotic shock. *Bioinformation*, **2017**, 13, 21-24	1.1	4
16	Exome sequencing found a novel homozygous deletion in ADCK3 gene involved in autosomal recessive spinocerebellar ataxia. *Gene*, **2019**, 708, 10-13	3.8	3
15	Characterizing the Retinoblastoma 1 locus: putative elements for Rb1 regulation by in silico analysis. *Frontiers in Genetics*, **2014**, 5, 2	4.5	3
14	Up-Regulation of and Is Associated with The Progression of Gastric-Type Adenocarcinoma. *Cell Journal*, **2017**, 19, 66-71	2.4	3
13	In silico finding of Putative Cis-Acting Elements for the Tethering of Polycomb Repressive Complex2 in Human Genome. *Bioinformation*, **2014**, 10, 187-90	1.1	3
12	SNHG1 Long Noncoding RNA is Potentially Up-Regulated in Colorectal Adenocarcinoma. *Asian Pacific Journal of Cancer Prevention*, **2020**, 21, 897-901	1.7	2
11	Exome sequencing revealed a p.G299R mutation in the COMP gene in an Iranian family suffering from pseudoachondroplasia. *Journal of Gene Medicine*, **2019**, 21, e3103	3.5	1
10	The non-coding RNA rprA can increase the resistance to ampicillin in Escherichia coli. *Microbial Pathogenesis*, **2019**, 129, 266-270	3.8	1
9	Exome sequencing revealed a novel deletion in the ERCC8 gene in an Iranian family with Cockayne syndrome. *Annals of Human Genetics*, **2018**, 82, 304-308	2.2	1
8	The construction of a short gene by a very fast, modified, and simplified gene synthesis and the analysis of various effects on this synthesis. *Brazilian Archives of Biology and Technology*, **2011**, 54, 53-60	1.8	1
7	HOTAIR Induces the Downregulation of miR-200 Family Members in Gastric Cancer Cell Lines. *Iranian Biomedical Journal*, **2022**, 26, 77-84	2	1
6	Whole exome sequencing identified a novel nonsense INPP4A mutation in a family with intellectual disability. *European Journal of Medical Genetics*, **2020**, 63, 103846	2.6	0
5	SNHG7 has an oncogenic role in colorectal cancer via potential sponging of MIR-485-5P and MIR-193A-5P; in silico approach. *Genetika*, **2021**, 53, 65-78	0.6	0
4	Identification of the mutation p.S867P in the PTPRQ gene in an Iranian family with hearing impairment. *Meta Gene*, **2017**, 13, 48-49	0.7	
3	Translational selection on SHH genes. *Genetics and Molecular Biology*, **2010**, 33, 408-10	2	
2	Whole exome sequencing revealed a novel dystrophin-related protein-2 () deletion in an Iranian family with symptoms of polyneuropathy. *Iranian Journal of Basic Medical Sciences*, **2019**, 22, 576-580	1.8	

1 Tissue Specific Expression Levels of Apoptosis Involved Genes Have Correlations with Codon and Amino Acid Usage. *Genomics and Informatics*, **2016**, 14, 234-240 1.9