## Alireza Pasdar

## List of Publications by Year in descending order

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304701 302107 1,783 87 22 39 h-index citations g-index papers 94 94 94 2985 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	PDâ€1/PDâ€1 pathway: Basic biology and role in cancer immunotherapy. Journal of Cellular Physiology, 2019, 234, 16824-16837.	4.1	279
2	Aptasensors as a new sensing technology developed for the detection of MUC1 mucin: A review. Biosensors and Bioelectronics, 2019, 130, 1-19.	10.1	103
3	The potential for circulating microRNAs in the diagnosis of myocardial infarction: a novel approach to disease diagnosis and treatment. Current Pharmaceutical Design, 2015, 22, 397-403.	1.9	83
4	Cytokine and growth factor profiling in patients with the metabolic syndrome. British Journal of Nutrition, 2015, 113, 1911-1919.	2.3	74
5	Therapeutic Potentials of BDNF/TrkB in Breast Cancer; Current Status and Perspectives. Journal of Cellular Biochemistry, 2017, 118, 2502-2515.	2.6	70
6	AutoMap is a high performance homozygosity mapping tool using next-generation sequencing data. Nature Communications, 2021, 12, 518.	12.8	68
7	Current insights into the metastasis of epithelial ovarian cancer - hopes and hurdles. Cellular Oncology (Dordrecht), 2020, 43, 515-538.	4.4	65
8	An imbalance in serum concentrations of inflammatory and anti-inflammatory cytokines in hypertension. Journal of the American Society of Hypertension, 2014, 8, 614-623.	2.3	59
9	Curcumin in tissue engineering: A traditional remedy for modern medicine. BioFactors, 2019, 45, 135-151.	5.4	53
10	<p>Antioxidant and toxicity studies of biosynthesized cerium oxide nanoparticles in rats</p> . International Journal of Nanomedicine, 2019, Volume 14, 2915-2926.	6.7	46
11	Paraoxonase gene polymorphisms and haplotype analysis in a stroke population. BMC Medical Genetics, 2006, 7, 28.	2.1	45
12	Circulating exosomal miRNAs in cardiovascular disease pathogenesis: New emerging hopes. Journal of Cellular Physiology, 2019, 234, 21796-21809.	4.1	43
13	Regulators and mechanisms of anoikis in triple-negative breast cancer (TNBC): A review. Critical Reviews in Oncology/Hematology, 2019, 140, 17-27.	4.4	40
14	The lipoprotein lipase S447X and cholesteryl ester transfer protein rs5882 polymorphisms and their relationship with lipid profile in human serum of obese individuals. Gene, 2015, 558, 195-199.	2.2	34
15	Familial combined hyperlipidemia: An overview of the underlying molecular mechanisms and therapeutic strategies. IUBMB Life, 2019, 71, 1221-1229.	3.4	34
16	Association of tumor necrosis factor-α promoter G-308A gene polymorphism with increased triglyceride level of subjects with metabolic syndrome. Gene, 2015, 568, 81-84.	2.2	31
17	The current status and perspectives regarding the clinical implication of intracellular calcium in breast cancer. Journal of Cellular Physiology, 2018, 233, 5623-5641.	4.1	31
18	Role of Pullulan in preparation of ceria nanoparticles and investigation of their biological activities. Journal of Molecular Structure, 2018, 1157, 127-131.	3 <b>.</b> 6	30

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19	The 9p21 Locus and its Potential Role in Atherosclerosis Susceptibility; Molecular Mechanisms and Clinical Implications. Current Pharmaceutical Design, 2016, 22, 5730-5737.	1.9	30
20	The effect of ABCA1gene polymorphisms on ischaemic stroke risk and relationship with lipid profile. BMC Medical Genetics, 2007, 8, 30.	2.1	27
21	Polymorphisms in non-coding RNAs and risk of colorectal cancer: A systematic review and meta-analysis. Critical Reviews in Oncology/Hematology, 2018, 132, 100-110.	4.4	25
22	Association of high level of hs-CRP with in-stent restenosis: A case-control study. Cardiovascular Revascularization Medicine, 2019, 20, 583-587.	0.8	25
23	Association between serum cytokine concentrations and the presence of hypertriglyceridemia. Clinical Biochemistry, 2016, 49, 750-755.	1.9	24
24	Association of heat shock protein70-2 (HSP70-2) gene polymorphism with coronary artery disease in an Iranian population. Gene, 2014, 550, 180-184.	2.2	21
25	Association of a Vascular Endothelial Growth Factor genetic variant with Serum VEGF level in subjects with Metabolic Syndrome. Gene, 2017, 598, 27-31.	2.2	20
26	A method for improving the efficiency of DNA extraction from clotted blood samples. Journal of Clinical Laboratory Analysis, 2019, 33, e22892.	2.1	20
27	Paraoxonase 1 (PON1) and stroke; the dilemma of genetic variation. Clinical Biochemistry, 2017, 50, 1298-1305.	1.9	19
28	Downregulation of Caspase 8 in a group of Iranian breast cancer patients – A pilot study. Journal of the Egyptian National Cancer Institute, 2017, 29, 191-195.	1.5	19
29	The prognostic and therapeutic values of long noncoding RNA PANDAR in colorectal cancer. Journal of Cellular Physiology, 2019, 234, 1230-1236.	4.1	19
30	The Association Between Oral Lichen Planus and Hepatitis C Virus Infection; A Report From Northeast of Iran. Jundishapur Journal of Microbiology, 2015, 8, e16741.	0.5	18
31	Association of heat shock protein70-2 ( <i>HSP70-2</i> ) gene polymorphism with obesity. Annals of Human Biology, 2016, 43, 542-546.	1.0	17
32	Common KRAS and NRAS gene mutations in sporadic colorectal cancer in Northeastern Iranian patients. Current Problems in Cancer, 2018, 42, 572-581.	2.0	17
33	The effect of curcumin (Curcuma longa L.) on circulating levels of adiponectin in patients with metabolic syndrome. Comparative Clinical Pathology, 2017, 26, 17-23.	0.7	16
34	Molecular aspects of hypercholesterolemia treatment: current perspectives and hopes. Annals of Medicine, 2018, 50, 303-311.	3.8	15
35	Next-generation sequencing and its application in diagnosis of retinitis pigmentosa. Ophthalmic Genetics, 2019, 40, 393-402.	1.2	15
36	Association between serum cell adhesion molecules with hs-CRP, uric acid and VEGF genetic polymorphisms in subjects with metabolic syndrome. Molecular Biology Reports, 2020, 47, 867-875.	2.3	15

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37	GATA3 somatic mutations are associated with clinicopathological features and expression profile in TCGA breast cancer patients. Scientific Reports, 2021, 11, 1679.	3.3	15
38	Prognostic Factors Associating with Pro-oxidant-antioxidant Balance; Neutrophils to Lymphocytes Ratio, Vitamin D, Heat Shock Protein 27, and Red Cell Distribution Width. Archives of Medical Research, 2020, 51, 261-267.	3.3	13
39	Personalised medicine in hypercholesterolaemia: the role of pharmacogenetics in statin therapy. Annals of Medicine, 2020, 52, 462-470.	3.8	12
40	The association between a Fatty Acid Binding Protein 1 (FABP1) gene polymorphism and serum lipid abnormalities in the MASHAD cohort study. Prostaglandins Leukotrienes and Essential Fatty Acids, 2021, 172, 102324.	2.2	12
41	Association of rs6921438 A <g 2018,="" 667,="" 70-75.<="" concentrations="" endothelial="" factor="" gene,="" growth="" in="" metabolic="" patients="" serum="" syndrome.="" td="" vascular="" with=""><td>2.2</td><td>11</td></g>	2.2	11
42	A novel variant in <i>LPL</i> gene is associated with familial combined hyperlipidemia. BioFactors, 2020, 46, 94-99.	5.4	10
43	Whole exome sequencing and homozygosity mapping reveals genetic defects in consanguineous Iranian families with inherited retinal dystrophies. Scientific Reports, 2020, 10, 19413.	3.3	9
44	Zinc Finger 259 Gene Polymorphism rs964184 is Associated with Serum Triglyceride Levels and Metabolic Syndrome. International Journal of Molecular and Cellular Medicine, 2016, 5, 8-18.	1.1	9
45	Association of <i>tumor necrosis factorâ€</i> ∫î± <i>â€308 G/A</i> gene polymorphism with coronary artery diseases: An evidenceâ€based study. Journal of Clinical Laboratory Analysis, 2018, 32, .	2.1	8
46	A novel mutation in <i>USF1</i> gene is associated with familial combined hyperlipidemia. IUBMB Life, 2020, 72, 616-623.	3.4	8
47	Association of and polymorphisms and their association with breast cancer risk among Iranian population. EXCLI Journal, 2019, 18, 429-438.	0.7	8
48	Haplotype Analysis of Interleukin-10 Gene Promoter Polymorphisms in Chronic Hepatitis C Infection: A Case Control Study. Viral Immunology, 2014, 27, 398-403.	1.3	7
49	Association of genetic polymorphisms of PON1 and CETP with the presence of metabolic syndrome; the effects of genotypes on their serum activity and concentrations. Egyptian Journal of Medical Human Genetics, 2018, 19, 43-48.	1.0	7
50	Significant association of TOX3/LOC643714 locus-rs3803662 and breast cancer risk in a cohort of Iranian population. Molecular Biology Reports, 2019, 46, 805-811.	2.3	7
51	Ki67 Frequency in Breast Cancers without Axillary Lymph Node Involvement and its Relation with Disease-free Survival. Asian Pacific Journal of Cancer Prevention, 2016, 17, 1347-1350.	1.2	7
52	Association of caspase 8 promoter variants and haplotypes with the risk of breast cancer and its molecular profile in an Iranian population: A caseâ€control study. Journal of Cellular Biochemistry, 2019, 120, 16435-16444.	2.6	6
53	The Impact of CASP8 rs10931936 and rs1045485 Polymorphisms as well as the Haplotypes on Breast Cancer Risk: A Case-Control Study. Clinical Breast Cancer, 2019, 19, e563-e577.	2.4	6
54	ESR1 gene variants, haplotypes and diplotypes may influence the risk of breast cancer and mammographic density. Molecular Biology Reports, 2020, 47, 8367-8375.	2.3	6

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55	The impact of CYP19A1 variants and haplotypes on breast cancer risk, clinicopathological features and prognosis. Molecular Genetics & Enomic Medicine, 2021, 9, e1705.	1.2	6
56	Clinical significance of circulating tumor cell related markers in patients with epithelial ovarian cancer before and after adjuvant chemotherapy. Scientific Reports, 2021, 11, 10524.	3.3	6
57	The effects of curcumin and a modified curcumin formulation on serum Cholesteryl Ester Transfer Protein concentrations in patients with metabolic syndrome: A randomized, placebo-controlled clinical trial. Avicenna Journal of Phytomedicine, 2018, 8, 330-337.	0.2	6
58	Association of SMAD7 genetic markers and haplotypes with colorectal cancer risk. BMC Medical Genomics, 2022, $15, 8$ .	1.5	6
59	There is an association between a genetic polymorphism in the ZNF259 gene involved in lipid metabolism and coronary artery disease. Gene, 2019, 704, 80-85.	2.2	5
60	CTLA-4 Gene Haplotypes and the Risk of Chronic Hepatitis C Infection; a Case Control Study. Reports of Biochemistry and Molecular Biology, 2017, 6, 51-58.	1.4	5
61	The interaction between a HSP-70 gene variant with dietary calories in determining serum markers of inflammation and cardiovascular risk. Clinical Nutrition, 2018, 37, 2122-2126.	5.0	4
62	Predictive and prognostic value of LSP1 rs3817198 in sporadic breast cancer in northeastern population of Iran. Experimental and Molecular Pathology, 2020, 116, 104514.	2.1	3
63	Ectopic Expression of miRNA-21 and miRNA-205 in Non-Small Cell Lung Cancer. International Journal of Cancer Management, 2019, In Press, .	0.4	3
64	A comparison of dietary intake between personnel of a gas processing company and a sample population of public employees from Mashhad. Clinical Nutrition ESPEN, 2020, 38, 124-128.	1.2	2
65	Evaluating the Association between CCR5delta32 Polymorphism (rs333) and the Risk of Breast Cancer in a Cohort of Iranian Population. Iranian Journal of Public Health, 2021, 50, 583-591.	0.5	2
66	Whole exome sequencing in 17 consanguineous Iranian pedigrees expands the mutational spectrum of inherited retinal dystrophies. Scientific Reports, 2021, 11, 19332.	3.3	2
67	Association of PICK1 and BDNF variations with increased risk of methamphetamine dependence among Iranian population: a case–control study. BMC Medical Genomics, 2021, 14, 27.	1.5	2
68	The Dilemma of TP53 Codon 72 Polymorphism (rs1042522) and Breast Cancer Risk: A Case-Control Study and Meta-Analysis in The Iranian Population. Cell Journal, 2020, 22, 185-192.	0.2	2
69	Haplotype Analysis of Hemochromatosis Gene Polymorphisms in Chronic Hepatitis C Virus Infection: A Case Control Study. Iranian Red Crescent Medical Journal, 2016, 18, e24675.	0.5	2
70	The significant role of a functional polymorphism in the NF-κB1 gene in breast cancer: evidence from an Iranian cohort. Future Oncology, 2021, 17, 4895-4905.	2.4	2
71	ÂEvaluation of serum HBV viral load, transaminases and histological features in chronic HBeAg-negative hepatitis B patients. Gastroenterology and Hepatology From Bed To Bench, 2017, 10, 39-43.	0.6	2
72	Association of Endonuclease G Gene Variants with Cardiovascular Disease Risk Factors. Reports of Biochemistry and Molecular Biology, 2019, 8, 147-152.	1.4	2

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73	Paraoxonase-1 Q192R polymorphism and its association with hs-CRP and fasting blood glucose levels and risk of coronary artery disease. Diabetes and Metabolic Syndrome: Clinical Research and Reviews, 2019, 13, 1053-1057.	3.6	1
74	Association of Interleukin-10 –592 CÂ>ÂA gene polymorphism with coronary artery disease: A case-control study and meta-analysis. Cytokine, 2021, 139, 155403.	3.2	1
75	Design, Construction and Evaluation of 1a/JFH1 HCV Chimera by Replacing the Intergenotypic Variable Region. Hepatitis Monthly, 2016, 16, e38261.	0.2	1
76	The association between genetic polymorphisms of the interleukin-10, tumor necrosis factor-alpha, and annexin A5 gene loci and restenosis after percutaneous coronary angioplasty and stenting. Journal of Research in Medical Sciences, 2019, 24, 68.	0.9	1
77	Genetic Determinants of Premature Menopause in A Mashhad Population Cohort. International Journal of Fertility & Sterility, 2021, 15, 26-33.	0.2	1
78	The relationship between genetic variants associated with primary ovarian insufficiency and lipid profile in women recruited from MASHAD cohort study. BMC Women's Health, 2022, 22, 2.	2.0	1
79	Association of macro-and micro-nutrients dietary intakes with rs2241883 genetic variants of FABP 1 gene in MASHAD study population. Clinical Nutrition ESPEN, 2021, 45, 262-266.	1.2	0
80	"p53 Polymorphism at Codon 72 and Breast Cancerâ€⊷ Letter. Journal of Cancer Prevention, 2017, 22, 55-55.	2.0	0
81	Secondary findings from whole-exome sequencing data in families with familial combined hyperlipidemia (FCHL). Egyptian Journal of Medical Human Genetics, 2021, 22, .	1.0	0
82	Comparison of (IFNG) +874 T/A Single Nucleotide Polymorphism in Hepatitis C Virus Infected Patients and Non-Infected Controls in Mashhad, Iran. Iranian Journal of Pathology, 2017, 12, 248-256.	0.5	0
83	A Novel Splice Site Variant in the LDLRAP1 Gene Causes Familial Hypercholesterolemia. Iranian Biomedical Journal, 2021, 25, 374-9.	0.7	0
84	The association between a variant of the cyclin-dependent kinase inhibitor 2A/B gene and risk of cardiovascular disease. Gene Reports, 2022, 26, 101480.	0.8	0
85	SARS-CoV-2 Liability: The Hidden Mystery Behind Its Presentation in Children. Advances in Experimental Medicine and Biology, 2021, 1353, 225-241.	1.6	0
86	A Genetic Variant in Proline and Serine Rich Coiled-Coil 1 Gene Is Associated with the Risk of Cardiovascular Disease. Reports of Biochemistry and Molecular Biology, 2022, 10, 653-663.	1.4	0
87	2q35-rs13387042 variant and the risk of breast cancer: a case–control study. Molecular Biology Reports, 2022, , 1.	2.3	0