Pantea Izadi

List of Publications by Year in descending order

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687363 677142 32 525 13 22 h-index citations g-index papers 33 33 33 942 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	A Panel of Plasma miRNAs 199b-3p, 224-5p and Let-7d-3p as Non-Invasive Diagnostic Biomarkers for Endometriosis. Reproductive Sciences, 2021, 28, 991-999.	2.5	17
2	A methylation signature at the CpG island promoter of estrogen receptor beta (ER- \hat{l}^2) in breasts of women may be an early footmark of lack of breastfeeding and nulliparity. Pathology Research and Practice, 2021, 218, 153328.	2.3	2
3	The Role of miRNAs 340-5p, 92a-3p, and 381-3p in Patients with Endometriosis: A Plasma and Mesenchymal Stem-Like Cell Study. BioMed Research International, 2021, 2021, 1-15.	1.9	6
4	Evaluation of Plasma MMP-9 and VEGF-A mRNAs as Non-invasive Diagnostic Biomarkers in Women with Endometriosis. Gene, Cell and Tissue, 2021, 9, .	0.2	1
5	Assessment of and Gene Promoter Methylation in Breast Invasive Ductal Carcinoma and Metastasis. Cell Journal, 2021, 23, 397-405.	0.2	2
6	Resveratrol alleviates non-alcoholic fatty liver disease through epigenetic modification of the Nrf2 signaling pathway. International Journal of Biochemistry and Cell Biology, 2020, 119, 105667.	2.8	88
7	Influence of plasma circulating exosomes obtained from obese women on tumorigenesis and tamoxifen resistance in ⟨scp⟩MCF⟨ scp⟩â€7 cells. IUBMB Life, 2020, 72, 1930-1940.	3.4	11
8	Up-regulation of FTO gene expression was associated with increase in skeletal muscle mass in overweight male adolescents. Archives of Medical Science, 2019, 15, 1133-1137.	0.9	24
9	Changes in FTO and IRX3 gene expression in obese and overweight male adolescents undergoing an intensive lifestyle intervention and the role of FTO genotype in this interaction. Journal of Translational Medicine, 2019, 17, 176.	4.4	30
10	Methylation of progesterone receptor isoform A promoter in normal breast tissue: An epigenetic link between early age at menarche and risk of breast cancer?. Journal of Cellular Biochemistry, 2019, 120, 12393-12401.	2.6	4
11	Interactions between macro-nutrients' intake, FTO and IRX3 gene expression, and FTO genotype in obese and overweight male adolescents. Adipocyte, 2019, 8, 386-391.	2.8	31
12	The Role of FTO Genotype in the Association Between FTO Gene Expression and Anthropometric Measures in Obese and Overweight Adolescent Boys. American Journal of Men's Health, 2019, 13, 155798831880811.	1.6	3
13	Early Parity Epigenetic Footprint of FOXA1 Gene Body in Normal Breast Tissue of Iranian Women. Iranian Biomedical Journal, 2019, 23, 99-106.	0.7	4
14	A link between expression level of long-non-coding RNA ZFAS1 in breast tissue of healthy women and obesity. International Journal of Biological Markers, 2018, 33, 500-506.	1.8	1
15	Expression levels of breast cancer-relatedGAS5andLSINCT5IncRNAs in cancer-free breast tissue: Molecular associations with age at menarche and obesity. Breast Journal, 2018, 24, 876-882.	1.0	17
16	PARP-1Overexpression as an Independent Prognostic Factor in Adult Non-M3 Acute Myeloid Leukemia. Genetic Testing and Molecular Biomarkers, 2018, 22, 343-349.	0.7	18
17	A haplotype of three SNPs in FTO had a strong association with body composition and BMI in Iranian male adolescents. PLoS ONE, 2018, 13, e0195589.	2.5	21
18	A complete linkage disequilibrium in a haplotype of three SNPs in Fat Mass and Obesity associated (FTO) gene was strongly associated with anthropometric indices after controlling for calorie intake and physical activity. BMC Medical Genetics, 2018, 19, 146.	2.1	9

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19	The Association between PARP1 and LIG3 Expression Levels and Chromosomal Translocations in Acute Myeloid Leukemia Patients. Cell Journal, 2018, 20, 204-210.	0.2	8
20	Radioprotective effect of melatonin on expression of Cdkn1a and Rad50 genes in rat peripheral blood. Journal of Cancer Research and Therapeutics, 2018, 14, 1070.	0.9	11
21	Comparison of BRCA1 Expression between Triple-Negative and Luminal Breast Tumors. Iranian Biomedical Journal, 2018, 22, 210-4.	0.7	5
22	Methylomics of breast cancer: Seeking epimarkers in peripheral blood of young subjects. Tumor Biology, 2017, 39, 101042831769504.	1.8	14
23	The High Frequency of PIK3CA Mutations in Iranian Breast Cancer Patients. Cancer Investigation, 2017, 35, 36-42.	1.3	10
24	Epigenetic Changes of the <i>ESR1</i> Gene in Breast Tissue of Healthy Women: A Missing Link with Breast Cancer Risk Factors?. Genetic Testing and Molecular Biomarkers, 2017, 21, 464-470.	0.7	17
25	Modulation of radiation-induced base excision repair pathway gene expression by melatonin. Journal of Medical Physics, 2017, 42, 245.	0.3	31
26	Effects of Maternal Isocaloric Diet Containing Different Amounts of Soy Oil and Extra Virgin Olive Oil on Weight, Serum Glucose, and Lipid Profile of Female Mice Offspring. Iranian Journal of Medical Sciences, 2017, 42, 161-169.	0.4	3
27	Femtomolar level detection of RASSF1A tumor suppressor gene methylation by electrochemical nano-genosensor based on Fe 3 O 4 /TMC/Au nanocomposite and PT-modified electrode. Biosensors and Bioelectronics, 2016, 77, 1095-1103.	10.1	70
28	Extra virgin olive oil in maternal diet increases osteogenic genes expression, but high amounts have deleterious effects on bones in mice offspring at adolescence. Iranian Journal of Basic Medical Sciences, 2016, 19, 1299-1307.	1.0	9
29	DNA methylation as a promising landscape: A simple blood test for breast cancer prediction. Tumor Biology, 2015, 36, 4905-4912.	1.8	30
30	Epigenetic marks in estrogen receptor alpha CpG island correlate with some reproductive risk factors in breast cancer. Molecular Biology Reports, 2014, 41, 7607-7612.	2.3	11
31	Mutation analysis of androgen receptor gene: Multiple uses for a single test. Gene, 2014, 552, 234-238.	2.2	3
32	Association of Poor Prognosis Subtypes of Breast Cancer with Estrogen Receptor Alpha Methylation in Iranian Women. Asian Pacific Journal of Cancer Prevention, 2012, 13, 4113-4117.	1.2	14