## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Can Muscle Building Supplements Increase Testicular Cancer Risk?. Frontiers in Nutrition, 2022, 9, 778426.	1.6	0
2	ZNF117 regulates glioblastoma stem cell differentiation towards oligodendroglial lineage. Nature Communications, 2022, 13, 2196.	5.8	9
3	Serum polychlorinated biphenyl (PCB) levels and risk of testicular germ cell tumors: A population-based case-control study in Connecticut and Massachusetts. Environmental Pollution, 2021, 273, 116458.	3.7	7
4	Hyperthermia Selectively Destabilizes Oncogenic Fusion Proteins. Blood Cancer Discovery, 2021, 2, 388-401.	2.6	26
5	Dataset of testicular germ cell tumors (TGCT) risk associated with serum polychlorinated biphenyl (PCB) by age at diagnosis and histologic types. Data in Brief, 2021, 36, 107014.	0.5	4
6	Co-distribution of Light At Night (LAN) and COVID-19 incidence in the United States. BMC Public Health, 2021, 21, 1509.	1.2	6
7	Valerian and valeric acid inhibit growth of breast cancer cells possibly by mediating epigenetic modifications. Scientific Reports, 2021, 11, 2519.	1.6	24
8	Valeric Acid Suppresses Liver Cancer Development by Acting as a Novel HDAC Inhibitor. Molecular Therapy - Oncolytics, 2020, 19, 8-18.	2.0	39
9	LRRC31 inhibits DNA repair and sensitizes breast cancer brain metastasis to radiation therapy. Nature Cell Biology, 2020, 22, 1276-1285.	4.6	39
10	Long noncoding RNA FLRL2 alleviated nonalcoholic fatty liver disease through Arntlâ€ <del>S</del> irt1 pathway. FASEB Journal, 2019, 33, 11411-11419.	0.2	28
11	Carcinogenicity of night shift work. Lancet Oncology, The, 2019, 20, 1058-1059.	5.1	219
12	Identification of Cancer Stem Cell Molecular Markers and Effects of hsa-miR-21-3p on Stemness in Esophageal Squamous Cell Carcinoma. Cancers, 2019, 11, 518.	1.7	25
13	Circadian genes and risk of prostate cancer in the prostate cancer prevention trial. Molecular Carcinogenesis, 2018, 57, 462-466.	1.3	15
14	piRNA-8041 is downregulated in human glioblastoma and suppresses tumor growth <i>in vitro</i> and <i>in vivo</i> . Oncotarget, 2018, 9, 37616-37626.	0.8	43
15	Combining Human Epigenetics and Sleep Studies in Caenorhabditis elegans: A Cross-Species Approach for Finding Conserved Genes Regulating Sleep. Sleep, 2017, 40, .	0.6	15
16	Transcriptome-wide piRNA profiling in human brains of Alzheimer's disease. Neurobiology of Aging, 2017, 57, 170-177.	1.5	71
17	Association Between Negative Age Stereotypes and Accelerated Cellular Aging: Evidence from Two Cohorts of Older Adults. Journal of the American Geriatrics Society, 2016, 64, e228-e230.	1.3	12
18	PIWI-Interacting RNAs in Gliomagenesis: Evidence from Post-GWAS and Functional Analyses. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1073-1080.	1.1	32

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19	Green tea consumption and risk of cardiovascular and ischemic related diseases: A meta-analysis. International Journal of Cardiology, 2016, 202, 967-974.	0.8	105
20	PIWI-interacting RNA 021285 is involved in breast tumorigenesis possibly by remodeling the cancer epigenome. Carcinogenesis, 2015, 36, 1094-1102.	1.3	122
21	The circadian gene CRY2 is associated with breast cancer aggressiveness possibly via epigenomic modifications. Tumor Biology, 2015, 36, 3533-3539.	0.8	24
22	Electric light, particularly at night, disrupts human circadian rhythmicity: is that a problem?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140120.	1.8	119
23	Aberrant methylation of miR-34b is associated with long-term shiftwork: a potential mechanism for increased breast cancer susceptibility. Cancer Causes and Control, 2015, 26, 171-178.	0.8	20
24	Epigenome-wide analysis of piRNAs in gene-specific DNA methylation. RNA Biology, 2014, 11, 1301-1312.	1.5	55
25	Targetome profiling and functional genetics implicate miR-618 in lymphomagenesis. Epigenetics, 2014, 9, 730-737.	1.3	37
26	Functional polymorphisms in the <i>NPAS2</i> gene are associated with overall survival in transcatheter arterial chemoembolizationâ€treated hepatocellular carcinoma patients. Cancer Science, 2014, 105, 825-832.	1.7	26
27	Prenatal exposure to organochlorine pesticides and infant birth weight in China. Chemosphere, 2014, 110, 1-7.	4.2	75
28	Functional polymorphisms of circadian negative feedback regulation genes are associated with clinical outcome in hepatocellular carcinoma patients receiving radical resection. Medical Oncology, 2014, 31, 179.	1.2	20
29	Dysregulated methylation at imprinted genes in prostate tumor tissue detected by methylation microarray. BMC Urology, 2013, 13, 37.	0.6	17
30	Advanced sleep schedules affect circadian gene expression in young adults with delayed sleep schedules. Sleep Medicine, 2013, 14, 449-455.	0.8	8
31	Targetome Profiling, Pathway Analysis and Genetic Association Study Implicate miR-202 in Lymphomagenesis. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 327-336.	1.1	35
32	Methylation alterations at imprinted genes detected among longâ€ŧerm shiftworkers. Environmental and Molecular Mutagenesis, 2013, 54, 141-146.	0.9	26
33	Aberrant DNA methylation of miRâ€219 promoter in longâ€ŧerm night shiftworkers. Environmental and Molecular Mutagenesis, 2013, 54, 406-413.	0.9	34
34	Potential cancer-related role of circadian gene TIMELESS suggested by expression profiling and in vitro analyses. BMC Cancer, 2013, 13, 498.	1.1	37
35	A functional polymorphism in <i><scp>PER</scp>3</i> gene is associated with prognosis in hepatocellular carcinoma. Liver International, 2012, 32, 1451-1459.	1.9	44
36	Genetic and epigenetic associations of circadian gene <i>TIMELESS</i> and breast cancer risk. Molecular Carcinogenesis, 2012, 51, 923-929.	1.3	61

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37	Functional polymorphisms of circadian positive feedback regulation genes and clinical outcome of Chinese patients with resected colorectal cancer. Cancer, 2012, 118, 937-946.	2.0	55
38	Epigenetic Impact of Long-Term Shiftwork: Pilot Evidence From Circadian Genes and Whole-Genome Methylation Analysis. Chronobiology International, 2011, 28, 852-861.	0.9	131
39	Genome-wide methylation analysis identifies involvement of TNF-α mediated cancer pathways in prostate cancer. Cancer Letters, 2011, 302, 47-53.	3.2	44
40	Effects of an advanced sleep schedule and morning short wavelength light exposure on circadian phase in young adults with late sleep schedules. Sleep Medicine, 2011, 12, 685-692.	0.8	65
41	Genetic and epigenetic association studies suggest a role of microRNA biogenesis gene exportin-5 (XPO5) in breast tumorigenesis. International Journal of Molecular Epidemiology and Genetics, 2011, 2, 9-18.	0.4	42
42	The circadian gene NPAS2 is a novel prognostic biomarker for breast cancer. Breast Cancer Research and Treatment, 2010, 120, 663-669.	1.1	66
43	Phenotypic effects of the circadian gene Cryptochrome 2 on cancer-related pathways. BMC Cancer, 2010, 10, 110.	1.1	44
44	The Core Circadian Gene <i>Cryptochrome 2</i> Influences Breast Cancer Risk, Possibly by Mediating Hormone Signaling. Cancer Prevention Research, 2010, 3, 539-548.	0.7	90
45	<i>CLOCK</i> in Breast Tumorigenesis: Genetic, Epigenetic, and Transcriptional Profiling Analyses. Cancer Research, 2010, 70, 1459-1468.	0.4	158
46	Clock-Cancer Connection in Non–Hodgkin's Lymphoma: A Genetic Association Study and Pathway Analysis of the Circadian Gene Cryptochrome 2. Cancer Research, 2009, 69, 3605-3613.	0.4	98
47	Testing the Circadian Gene Hypothesis in Prostate Cancer: A Population-Based Case-Control Study. Cancer Research, 2009, 69, 9315-9322.	0.4	137
48	Cancer-related transcriptional targets of the circadian gene NPAS2 identified by genome-wide ChIP-on-chip analysis. Cancer Letters, 2009, 284, 149-156.	3.2	36
49	microRNA miR-196a-2 and Breast Cancer: A Genetic and Epigenetic Association Study and Functional Analysis. Cancer Research, 2009, 69, 5970-5977.	0.4	325
50	Non-synonymous polymorphisms in the circadian gene NPAS2 and breast cancer risk. Breast Cancer Research and Treatment, 2008, 107, 421-425.	1.1	104
51	Correlating observed odds ratios from lung cancer case–control studies to SNP functional scores predicted by bioinformatic tools. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2008, 639, 80-88.	0.4	22
52	A SNP in a <i>let-7</i> microRNA Complementary Site in the <i>KRAS</i> 3′ Untranslated Region Increases Non–Small Cell Lung Cancer Risk. Cancer Research, 2008, 68, 8535-8540.	0.4	609
53	Clock-cancer connection in non-Hodgkin's lymphoma. Medical Hypotheses, 2008, 70, 788-792.	0.8	13
54	The Circadian Gene <i>NPAS2</i> , a Putative Tumor Suppressor, Is Involved in DNA Damage Response. Molecular Cancer Research, 2008, 6, 1461-1468.	1.5	93

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55	Correlation between Circadian Gene Variants and Serum Levels of Sex Steroids and Insulin-like Growth Factor-I. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 3268-3273.	1.1	31
56	Ala394Thr polymorphism in the clock geneNPAS2: A circadian modifier for the risk of non-Hodgkin's lymphoma. International Journal of Cancer, 2007, 120, 432-435.	2.3	100
57	Does "Clock" Matter in Prostate Cancer?. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 3-5.	1.1	47
58	A Putative Exonic Splicing Polymorphism in the BCL6 Gene and the Risk of Non-Hodgkin Lymphoma. Journal of the National Cancer Institute, 2005, 97, 1616-1618.	3.0	33
59	Genotypes and haplotypes of the methyl-CpG-binding domain 2 modify breast cancer risk dependent upon menopausal status. Breast Cancer Research, 2005, 7, R745-52.	2.2	13
60	Period3 structural variation: a circadian biomarker associated with breast cancer in young women. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 268-70.	1.1	78
61	Methyl-CpG-binding domain 2. Cancer, 2004, 100, 1853-1858.	2.0	19
62	An Evolutionary Perspective on Single-Nucleotide Polymorphism Screening in Molecular Cancer Epidemiology. Cancer Research, 2004, 64, 2251-2257.	0.4	100
63	Genetic instability of specific chromosomes associated with a family history of cancer. Cancer Genetics and Cytogenetics, 2002, 136, 73-77.	1.0	8
64	A case-control analysis of lymphocytic chromosome 9 aberrations in lung cancer. International Journal of Cancer, 2002, 102, 536-540.	2.3	7
65	BPDE-induced lymphocytic 3p21.3 aberrations may predict head and neck carcinoma risk. Cancer, 2002, 95, 563,568	2.0	10