Alessandro Weisz

List of Publications by Year
in descending order

165	7,966	53794	58581
papers	citations	45 h-index	82 g-index

172	172	172	10287
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The pleiotropic roles of circular and long noncoding RNAs in cutaneous melanoma. Molecular Oncology, 2022, 16, 565-593.	4.6	11
2	ABCA1, TCF7, NFATC1, PRKCZ, and PDGFA DNA methylation as potential epigenetic-sensitive targets in acute coronary syndrome via network analysis. Epigenetics, 2022, 17, 547-563.	2.7	9
3	Maternal pre-pregnancy overweight and neonatal gut bacterial colonization are associated with cognitive development and gut microbiota composition in pre-school-age offspring. Brain, Behavior, and Immunity, 2022, 100, 311-320.	4.1	32
4	Nasopharyngeal virome analysis of COVIDâ€19 patients during three different waves in Campania region of Italy. Journal of Medical Virology, 2022, , .	5.0	9
5	Rapid and sensitive detection of SARS-CoV-2 variants in nasopharyngeal swabs and wastewaters. Diagnostic Microbiology and Infectious Disease, 2022, 102, 115632.	1.8	6
6	NGS analysis of nasopharyngeal microbiota in SARS-CoV-2 positive patients during the first year of the pandemic in the Campania Region of Italy. Microbial Pathogenesis, 2022, 165, 105506.	2.9	12
7	Histone Methyltransferase DOT1L as a Promising Epigenetic Target for Treatment of Solid Tumors. Frontiers in Genetics, 2022, 13, 864612.	2.3	22
8	Correction: Analysis of miRNA profiles identified miR-196a as a crucial mediator of aberrant PI3K/AKT signaling in lung cancer cells. Oncotarget, 2022, 13, 755-755.	1.8	0
9	Identification of functional pathways and molecular signatures in neuroendocrine neoplasms by multi-omics analysis. Journal of Translational Medicine, 2022, 20, .	4.4	14
10	WIND (Workflow for pIRNAs aNd beyonD): a strategy for in-depth analysis of small RNA-seq data. F1000Research, 2021, 10, 1.	1.6	5
11	A ZFYVE19 gene mutation associated with neonatal cholestasis and cilia dysfunction: case report with a novel pathogenic variant. Orphanet Journal of Rare Diseases, 2021, 16, 179.	2.7	11
12	Higher Integrin Alpha 3 Beta1 Expression in Papillary Thyroid Cancer Is Associated with Worst Outcome. Cancers, 2021, 13, 2937.	3.7	10
13	WIND (Workflow for pIRNAs aNd beyonD): a strategy for in-depth analysis of small RNA-seq data. F1000Research, 2021, 10, 1.	1.6	22
14	HOME-BIO (sHOtgun MEtagenomic analysis of BIOlogical entities): a specific and comprehensive pipeline for metagenomic shotgun sequencing data analysis. BMC Bioinformatics, 2021, 22, 106.	2.6	9
15	Whole-genome sequencing of Pseudomonas sp. TAE6080, a strain capable of inhibiting Staphylococcus epidermidis biofilm. Marine Genomics, 2021, 60, 100887.	1.1	3
16	Circulating SARS-CoV-2 variants in Italy, October 2020â€“March 2021. Virology Journal, 2021, 18, 168.	3.4	36
17	In vitro CSC-derived cardiomyocytes exhibit the typical microRNA-mRNA blueprint of endogenous cardiomyocytes. Communications Biology, 2021, 4, 1146.	4.4	15
18	DNMT3A epigenetically regulates key microRNAs involved in epithelial-to-mesenchymal transition in prostate cancer. Carcinogenesis, 2021, 42, 1449-1460.	2.8	10

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19	Small-bowel carcinomas associated with celiac disease: transcriptomic profiling shows predominance of microsatellite instability-immune and mesenchymal subtypes. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2020, 476, 711-723.	2.8	13
20	Interaction Proteomics Identifies ERbeta Association with Chromatin Repressive Complexes to Inhibit Cholesterol Biosynthesis and Exert An Oncosuppressive Role in Triple-negative Breast Cancer. Molecular and Cellular Proteomics, 2020, 19, 245-260.	3.8	18
21	Identification of Antiestrogenâ€Bound Estrogen Receptor Î± Interactomes in Hormoneâ€Responsive Human Breast Cancer Cell Nuclei. Proteomics, 2020, 20, 2000135.	2.2	4
22	Global View of Candidate Therapeutic Target Genes in Hormone-Responsive Breast Cancer. International Journal of Molecular Sciences, 2020, 21, 4068.	4.1	13
23	An Overview of Candidate Therapeutic Target Genes in Ovarian Cancer. Cancers, 2020, 12, 1470.	3.7	20
24	Insights into the Role of Estrogen Receptor Î² in Triple-Negative Breast Cancer. Cancers, 2020, 12, 1477.	3.7	33
25	Atrial myxomas arise from multipotent cardiac stem cells. European Heart Journal, 2020, 41, 4332-4345.	2.2	51
26	Small Non-Coding RNA Profiling Identifies miR-181a-5p as a Mediator of Estrogen Receptor Beta-Induced Inhibition of Cholesterol Biosynthesis in Triple-Negative Breast Cancer. Cells, 2020, 9, 874.	4.1	25
27	Loss of Spry1 reduces growth of BRAFV600-mutant cutaneous melanoma and improves response to targeted therapy. Cell Death and Disease, 2020, 11, 392.	6.3	14
28	Abstract 1794: Loss of Spry1 reduces growth of BRAFV600-mutant cutaneous melanoma and improves response to targeted therapy. , 2020, , .		0
29	Molecular and Functional Characterization of the Somatic PIWIL1/piRNA Pathway in Colorectal Cancer Cells. Cells, 2019, 8, 1390.	4.1	16
30	The Histone Methyltransferase DOT1L Is a Functional Component of Estrogen Receptor Alpha Signaling in Ovarian Cancer Cells. Cancers, 2019, 11, 1720.	3.7	24
31	The RNA-mediated estrogen receptor Î± interactome of hormone-dependent human breast cancer cell nuclei. Scientific Data, 2019, 6, 173.	5.3	18
32	Summary of the International Conference on Onco-Nephrology: an emerging field in medicine. Kidney International, 2019, 96, 555-567.	5.2	47
33	DNA methylation dynamic of bone marrow hematopoietic stem cells after allogeneic transplantation. Stem Cell Research and Therapy, 2019, 10, 138.	5.5	12
34	Inhibition of histone methyltransferase DOT1L silences ERÎ± gene and blocks proliferation of antiestrogen-resistant breast cancer cells. Science Advances, 2019, 5, eaav5590.	10.3	70
35	Fetal cardiac growth is associated with in utero gut colonization. Nutrition, Metabolism and Cardiovascular Diseases, 2019, 29, 170-176.	2.6	10
36	Quantitative mapping of RNA-mediated nuclear estrogen receptor Î² interactome in human breast cancer cells. Scientific Data, 2018, 5, 180031.	5.3	22

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37	Identification of a novel truncating mutation in PALB2 gene by a multigene sequencing panel for mutational screening of breast cancer riskâ€associated and related genes. Journal of Clinical Laboratory Analysis, 2018, 32, e22418.	2.1	5
38	Splicing of platelet resident pre-mRNAs upon activation by physiological stimuli results in functionally relevant proteome modifications. Scientific Reports, 2018, 8, 498.	3.3	65
39	A Large Set of miRNAs Is Dysregulated from the Earliest Steps of Human Hepatocellular Carcinoma Development. American Journal of Pathology, 2018, 188, 785-794.	3.8	15
40	TNF-alpha and metalloproteases as key players in melanoma cells aggressiveness. Journal of Experimental and Clinical Cancer Research, 2018, 37, 326.	8.6	73
41	Identification of long nonâ€coding RNA expression patterns useful for molecularâ€based classification of typeï¿½I endometrial cancers. Oncology Reports, 2018, 41, 1209-1217.	2.6	4
42	miRNA Regulation of the Hyperproliferative Phenotype of Vascular Smooth Muscle Cells in Diabetes. Diabetes, 2018, 67, 2554-2568.	0.6	53
43	Uncoupling effects of estrogen receptor Î± on LKB1/AMPK interaction upon adiponectin exposure in breast cancer. FASEB Journal, 2018, 32, 4343-4355.	0.5	43
44	Relations of gut liver axis components and gut microbiota in obese children with fatty liver: A pilot study. Clinics and Research in Hepatology and Gastroenterology, 2018, 42, 387-390.	1.5	11
45	miR-196a Is Able to Restore the Aggressive Phenotype of Annexin A1 Knock-Out in Pancreatic Cancer Cells by CRISPR/Cas9 Genome Editing. International Journal of Molecular Sciences, 2018, 19, 1967.	4.1	27
46	Small non-coding RNA landscape is modified by GPAT2 silencing in MDA-MB-231 cells. Oncotarget, 2018, 9, 28141-28154.	1.8	5
47	Abstract 1839: Suppression of Spry1 sensitizes cutaneous melanoma to BRAF-targeted therapy. , 2018, , .		0
48	Infantile spasms in early-onset Niemannâ€“Pick disease with a novel compound heterozygous mutations in SMPD1 gene. European Journal of Molecular and Clinical Medicine, 2017, 2, 155.	0.1	1
49	iSmaRT: a toolkit for a comprehensive analysis of small RNA-Seq data. Bioinformatics, 2017, 33, 938-940.	4.1	21
50	Exploiting a new strategy to induce immunogenic cell death to improve dendritic cell-based vaccines for lymphoma immunotherapy. OncoImmunology, 2017, 6, e1356964.	4.6	42
51	Design and expression of peptides with antimicrobial activity againstSalmonellatyphimurium. Cellular Microbiology, 2017, 19, e12645.	2.1	5
52	Specific gene expression signatures induced by the multiple oncogenic alterations that occur within the PTEN/PI3K/AKT pathway in lung cancer. PLoS ONE, 2017, 12, e0178865.	2.5	49
53	The nuclear receptor ERÎ² engages AGO2 in regulation of gene transcription, RNA splicing and RISC loading. Genome Biology, 2017, 18, 189.	8.8	63
54	Analysis of miRNA profiles identified miR-196a as a crucial mediator of aberrant PI3K/AKT signaling in lung cancer cells. Oncotarget, 2017, 8, 19172-19191.	1.8	32

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55	DNA methylation variations are required for epithelial-to-mesenchymal transition induced by cancer-associated fibroblasts in prostate cancer cells. Oncogene, 2017, 36, 5551-5566.	5.9	88
56	Abstract 1196: Epigenetic drugs modulate long noncoding RNAs expression in BRAF inhibitor-resistant melanoma. , 2017, , .		0
57	Abstract 1467: A large set of miRNAs is deregulated since the earliest steps of human HCC. , 2017, , .		0
58	IBTK Differently Modulates Gene Expression and RNA Splicing in HeLa and K562 Cells. International Journal of Molecular Sciences, 2016, 17, 1848.	4.1	11
59	PDGFR-alpha inhibits melanoma growth via CXCL10/IP-10: a multi-omics approach. Oncotarget, 2016, 7, 77257-77275.	1.8	22
60	Large-scale profiling of signalling pathways reveals an asthma specific signature in bronchial smooth muscle cells. Oncotarget, 2016, 7, 25150-25161.	1.8	32
61	Pesticide toxicogenomics across scales: in vitro transcriptome predicts mechanisms and outcomes of exposure in vivo. Scientific Reports, 2016, 6, 38131.	3.3	20
62	The RNA-Binding Protein SYNCRIP Is a Component of the Hepatocyte Exosomal Machinery Controlling MicroRNA Sorting. Cell Reports, 2016, 17, 799-808.	6.4	438
63	Gut microbiota composition and products contribute to gutâ€“liver axis dysfunction in pediatric obesity related NAFLD, with distinct metabolomic signature. Digestive and Liver Disease, 2016, 48, e260.	0.9	0
64	Kleefstra-variant syndrome with heterozygous mutations in EHMT1 and KCNQ2 genes: a case report. Neurological Sciences, 2016, 37, 829-831.	1.9	4
65	Characterization of two de novo KCNT1 mutations in children with malignant migrating partial seizures in infancy. Molecular and Cellular Neurosciences, 2016, 72, 54-63.	2.2	77
66	Phenytoin neurotoxicity in a child carrying new STXBP1 and CYP2C9 gene mutations. Seizure: the Journal of the British Epilepsy Association, 2016, 34, 26-28.	2.0	7
67	Small RNA profiling reveals deregulated phosphatase and tensin homolog (PTEN)/phosphoinositide 3-kinase (PI3K)/Akt pathway in bronchial smooth muscle cells from asthmatic patients. Journal of Allergy and Clinical Immunology, 2016, 137, 58-67.	2.9	30
68	Specific patterns of PIWI-interacting small noncoding RNA expression in dysplastic liver nodules and hepatocellular carcinoma. Oncotarget, 2016, 7, 54650-54661.	1.8	63
69	Leptin as a mediator of tumor-stromal interactions promotes breast cancer stem cell activity. Oncotarget, 2016, 7, 1262-1275.	1.8	74
70	Small non-coding RNA deregulation in endometrial carcinogenesis. Oncotarget, 2015, 6, 4677-4691.	1.8	49
71	Identification of cytoplasmic proteins interacting with unliganded estrogen receptor Î± and Î² in human breast cancer cells. Proteomics, 2015, 15, 1801-1807.	2.2	17
72	Identification of Genes Selectively Regulated in Human Hepatoma Cells by Treatment With Dyslipidemic Sera and PUFAs. Journal of Cellular Physiology, 2015, 230, 2059-2066.	4.1	7

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73	Activating stimuli induce platelet microRNA modulation and proteome reorganisation. Thrombosis and Haemostasis, 2015, 114, 96-108.	3.4	40
74	Estrogen receptor beta impacts hormone-induced alternative mRNA splicing in breast cancer cells. BMC Genomics, 2015, 16, 367.	2.8	28
75	The Akt1/IL-6/STAT3 pathway regulates growth of lung tumor initiating cells. Oncotarget, 2015, 6, 42667-42686.	1.8	43
76	Effect of low frequency (LF) electric fields on gene expression of a bone human cell line. Electromagnetic Biology and Medicine, 2014, 33, 289-295.	1.4	6
77	Lack of pathogenic mutations in six patients with MMPSI. Epilepsy Research, 2014, 108, 340-344.	1.6	40
78	Identification of H ferritin-dependent and independent genes in K562 differentiating cells by targeted gene silencing and expression profiling. Gene, 2014, 535, 327-335.	2.2	15
79	Post-transcriptional Regulation of Human Breast Cancer Cell Proteome by Unliganded Estrogen Receptor Î² via microRNAs. Molecular and Cellular Proteomics, 2014, 13, 1076-1090.	3.8	33
80	Timed regulation of P-element-induced wimpy testis-interacting RNA expression during rat liver regeneration. Hepatology, 2014, 60, 798-806.	7.3	48
81	Single-Cell States in the Estrogen Response of Breast Cancer Cell Lines. PLoS ONE, 2014, 9, e88485.	2.5	4
82	Global Transcriptome Profiles of Italian Mediterranean Buffalo Embryos with Normal and Retarded Growth. PLoS ONE, 2014, 9, e90027.	2.5	14
83	RNA sequencing identifies specific PIWI-interacting small non-coding RNA expression patterns in breast cancer. Oncotarget, 2014, 5, 9901-9910.	1.8	145
84	Molecular Mechanisms of Selective Estrogen Receptor Modulator Activity in Human Breast Cancer Cells: Identification of Novel Nuclear Cofactors of Antiestrogenâ€“ERÎ± Complexes by Interaction Proteomics. Journal of Proteome Research, 2013, 12, 421-431.	3.7	32
85	iMir: An integrated pipeline for high-throughput analysis of small non-coding RNA data obtained by smallRNA-Seq. BMC Bioinformatics, 2013, 14, 362.	2.6	62
86	New Insights on Estrogen Receptor Actions in Hormone-Responsive Breast Cancer Cells by Interaction Proteomics. , 2013, , 149-174.		1
87	Direct regulation of microRNA biogenesis and expression by estrogen receptor beta in hormone-responsive breast cancer. Oncogene, 2012, 31, 4196-4206.	5.9	87
88	Signaling Networks Associated with AKT Activation in Non-Small Cell Lung Cancer (NSCLC): New Insights on the Role of Phosphatydil-Inositol-3 kinase. PLoS ONE, 2012, 7, e30427.	2.5	119
89	Effects of Oestrogen on MicroRNA Expression in Hormone-Responsive Breast Cancer Cells. Hormones and Cancer, 2012, 3, 65-78.	4.9	51
90	Comparative analysis of nuclear estrogen receptor alpha and beta interactomes in breast cancer cells. Molecular BioSystems, 2011, 7, 667-676.	2.9	39

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91	Expression of c-jun is not mandatory for mouse hepatocyte proliferation induced by two nuclear receptor ligands: TCPOBOP and T3. Journal of Hepatology, 2011, 55, 1069-1078.	3.7	8
92	Global analysis of estrogen receptor beta binding to breast cancer cell genome reveals an extensive interplay with estrogen receptor alpha for target gene regulation. BMC Genomics, 2011, 12, 36.	2.8	140
93	Identification of proteins associated with ligandâ€activated estrogen receptor Î± in human breast cancer cell nuclei by tandem affinity purification and nano LCâ€MS/MS. Proteomics, 2011, 11, 172-179.	2.2	35
94	A large set of estrogen receptor Î²â€interacting proteins identified by tandem affinity purification in hormoneâ€responsive human breast cancer cell nuclei. Proteomics, 2011, 11, 159-165.	2.2	36
95	Specific inhibition of NF-Y subunits triggers different cell proliferation defects. Nucleic Acids Research, 2011, 39, 5356-5368.	14.5	73
96	Molecular bases of copper and iron deficiency-associated dyslipidemia: a microarray analysis of the rat intestinal transcriptome. Genes and Nutrition, 2010, 5, 1-8.	2.5	28
97	The p63 target HBP1 is required for skin differentiation and stratification. Cell Death and Differentiation, 2010, 17, 1896-1907.	11.2	19
98	C/EBPÎ´ Gene Targets in Human Keratinocytes. PLoS ONE, 2010, 5, e13789.	2.5	11
99	The Reverse Transcription Inhibitor Abacavir Shows Anticancer Activity in Prostate Cancer Cell Lines. PLoS ONE, 2010, 5, e14221.	2.5	48
100	Identification of a Hormone-regulated Dynamic Nuclear Actin Network Associated with Estrogen Receptor Î± in Human Breast Cancer Cell Nuclei. Molecular and Cellular Proteomics, 2010, 9, 1352-1367.	3.8	59
101	Epigenetic alteration of microRNAs in DNMT3B-mutated patients of ICF syndrome. Epigenetics, 2010, 5, 427-443.	2.7	31
102	Estrogen Receptor Î± Controls a Gene Network in Luminal-Like Breast Cancer Cells Comprising Multiple Transcription Factors and MicroRNAs. American Journal of Pathology, 2010, 176, 2113-2130.	3.8	151
103	Pharmacogenomics and analogues of the antitumour agent N⁶â€isopentenyladenosine. International Journal of Cancer, 2009, 124, 2179-2185.	5.1	25
104	Identification of new genes associated with breast cancer progression by gene expression analysis of predefined sets of neoplastic tissues. International Journal of Cancer, 2008, 123, 1327-1338.	5.1	79
105	Quantitative expression profiling of highly degraded RNA from formalin-fixed, paraffin-embedded breast tumor biopsies by oligonucleotide microarrays. Laboratory Investigation, 2008, 88, 430-440.	3.7	76
106	Time-course analysis of genome-wide gene expression data from hormone-responsive human breast cancer cells. BMC Bioinformatics, 2008, 9, S12.	2.6	27
107	Influence of estrogens and antiestrogens on the expression of selected hormone-responsive genes. Maturitas, 2007, 57, 50-55.	2.4	13
108	Comparative gene expression profiling reveals partially overlapping but distinct genomic actions of different antiestrogens in human breast cancer cells. Journal of Cellular Biochemistry, 2006, 98, 1163-1184.	2.6	43

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109	Tumor-selective action of HDAC inhibitors involves TRAIL induction in acute myeloid leukemia cells. Nature Medicine, 2005, 11, 77-84.	30.7	567
110	Estrogens and Progesterone Promote Persistent CCND1 Gene Activation during G 1 by Inducing Transcriptional Derepression via c- Jun /c- Fos /Estrogen Receptor (Progesterone Receptor) Complex Assembly to a Distal Regulatory Element and Recruitment of Cyclin D1 to Its Own Gene Promoter. Molecular and Cellular Biology, 2004, 24, 7260-7274.	2.3	154
111	A genomic view of estrogen actions in human breast cancer cells by expression profiling of the hormone-responsive transcriptome. Journal of Molecular Endocrinology, 2004, 32, 719-775.	2.5	80
112	Molecular identification of ER?-positive breast cancer cells by the expression profile of an intrinsic set of estrogen regulated genes. Journal of Cellular Physiology, 2004, 200, 440-450.	4.1	44
113	Malignant Mammary Cells Acquire Independence from Extracellular Context for Regulation of Estrogen Receptor Î±. Clinical Cancer Research, 2004, 10, 402s-409s.	7.0	19
114	Identification of genes involved in growth inhibition of breast cancer cells transduced with estrogen receptor. FEBS Letters, 2003, 553, 445-450.	2.8	20
115	Distinct Signaling Pathways Mediate Stimulation of Cell Cycle Progression and Prevention of Apoptotic Cell Death by Estrogen in Rat Pituitary Tumor PR1 Cells. Molecular Biology of the Cell, 2003, 14, 5051-5059.	2.1	15
116	Quantitative real-time RT-PCR analysis of eight novel estrogen-regulated genes in breast cancer. International Journal of Biological Markers, 2003, 18, 123-129.	1.8	16
117	Distinct Nongenomic Signal Transduction Pathways Controlled by 17Î²-Estradiol Regulate DNA Synthesis and Cyclin D1Gene Transcription in HepG2 Cells. Molecular Biology of the Cell, 2002, 13, 3720-3729.	2.1	131
118	Effects of nitric oxide donors on vascular endothelial growth factor gene induction. Biochemical and Biophysical Research Communications, 2002, 296, 976-982.	2.1	45
119	Inducible Nitric Oxide Synthase in a Human Glioblastoma Cell Line. Journal of Neurochemistry, 2002, 64, 85-91.	3.9	53
120	Cell type-specific induction of cyclin D and cyclin-dependent kinase inhibitor p27kip1 expression by estrogen in rat endometrium. Journal of Steroid Biochemistry and Molecular Biology, 2001, 78, 193-199.	2.5	13
121	17Î²-Estradiol inhibits forskolin-induced vascular endothelial growth factor promoter in MCF-7 breast adenocarcinoma cells. Journal of Steroid Biochemistry and Molecular Biology, 2001, 78, 343-349.	2.5	6
122	Inhibition of Human Breast Cancer Cell Growth by Blockade of the Mevalonate-Protein Prenylation Pathway is not Prevented by Overexpression of Cyclin D1. Breast Cancer Research and Treatment, 2001, 67, 23-33.	2.5	4
123	Identification of Hypoxia-inducible Factor 1 Ancillary Sequence and Its Function in Vascular Endothelial Growth Factor Gene Induction by Hypoxia and Nitric Oxide. Journal of Biological Chemistry, 2001, 276, 2292-2298.	3.4	229
124	Insulin Up-Regulates Vascular Endothelial Growth Factor and Stabilizes Its Messengers in Endometrial Adenocarcinoma Cells1. Journal of Clinical Endocrinology and Metabolism, 2001, 86, 363-368.	3.6	33
125	Insulin Up-Regulates Vascular Endothelial Growth Factor and Stabilizes Its Messengers in Endometrial Adenocarcinoma Cells. Journal of Clinical Endocrinology and Metabolism, 2001, 86, 363-368.	3.6	34
126	Estrogens do not modify MAP kinase-dependent nuclear signaling during stimulation of early G(1) progression in human breast cancer cells. Cancer Research, 2001, 61, 6360-6.	0.9	26

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127	Regulation of vascular endothelial growth factor expression by insulin-like growth factor-I in endometrial adenocarcinoma cells. International Journal of Cancer, 2000, 85, 117-123.	5.1	68
128	Hypoxia response element of the human vascular endothelial growth factor gene mediates transcriptional regulation by nitric oxide: control of hypoxia-inducible factor-1 activity by nitric oxide. Blood, 2000, 95, 189-197.	1.4	395
129	The antiestrogen ICI 182,780 inhibits proliferation of human breast cancer cells by interfering with multiple, sequential estrogen-regulated processes required for cell cycle completion. Molecular and Cellular Endocrinology, 2000, 165, 199-209.	3.2	21
130	Hypoxia response element of the human vascular endothelial growth factor gene mediates transcriptional regulation by nitric oxide: control of hypoxia-inducible factor-1 activity by nitric oxide. Blood, 2000, 95, 189-197.	1.4	11
131	Regulation of vascular endothelial growth factor expression by insulin-like growth factor-I in endometrial adenocarcinoma cells. International Journal of Cancer, 2000, 85, 117.	5.1	5
132	ERE environment- and cell type-specific transcriptional effects of estrogen in normal endometrial cells. Molecular and Cellular Endocrinology, 1998, 139, 153-160.	3.2	2
133	Constitutive overexpression of cyclin D1 does not prevent inhibition of hormone-responsive human breast cancer cell growth by antiestrogens. Cancer Research, 1998, 58, 871-6.	0.9	25
134	Estrogen Induces Early and Timed Activation of Cyclin-Dependent Kinases 4, 5, and 6 and Increases Cyclin Messenger Ribonucleic Acid Expression in Rat Uterus¹. Endocrinology, 1997, 138, 978-984.	2.8	78
135	Inhibition of inducible nitric oxide synthase gene expression by glucocorticoid-induced protein(s) in lipopolysaccharide-stimulated J774 cells. European Journal of Pharmacology, 1997, 339, 87-95.	3.5	13
136	Estrogen Induces Early and Timed Activation of Cyclin-Dependent Kinases 4, 5, and 6 and Increases Cyclin Messenger Ribonucleic Acid Expression in Rat Uterus. Endocrinology, 1997, 138, 978-984.	2.8	19
137	Stimulation of Human Breast Cancer MCF-7 Cells with Estrogen Prevents Cell Cycle Arrest by HMG-CoA Reductase Inhibitors. Biochemical and Biophysical Research Communications, 1996, 220, 864-870.	2.1	40
138	Regulation of the mouse inducible-type nitric oxide synthase gene promoter by interferon-Î³, bacterial lipopolysaccharide and NG-monomethyl-l-arginine. Biochemical Journal, 1996, 316, 209-215.	3.7	149
139	17 beta-Estradiol overcomes a G1 block induced by HMG-CoA reductase inhibitors and fosters cell cycle progression without inducing ERK-1 and -2 MAP kinases activation. Oncogene, 1996, 12, 753-63.	5.9	46
140	17beta-Estradiol induces cyclin D1 gene transcription, p36D1-p34cdk4 complex activation and p105Rb phosphorylation during mitogenic stimulation of G(1)-arrested human breast cancer cells. Oncogene, 1996, 12, 2315-24.	5.9	244
141	Implication of nitric oxide synthase in carcinogenesis: analysis of the human inducible nitric oxide synthase gene. Pharmacogenetics and Genomics, 1995, 5, S166-S170.	5.7	10
142	In vivo functional analysis of the mouse estrogen receptor gene promoter: a transgenic mouse model to study tissue-specific and developmental regulation of estrogen receptor gene transcription. Molecular Endocrinology, 1995, 9, 1077-1090.	3.7	18
143	Enhancement of inducible-type NO synthase gene transcription by protein synthesis inhibitors. FEBS Letters, 1994, 338, 326-330.	2.8	16
144	Dual mechanism for the control of inducible-type NO synthase gene expression in macrophages during activation by interferon-gamma and bacterial lipopolysaccharide. Transcriptional and post-transcriptional regulation. Journal of Biological Chemistry, 1994, 269, 8324-33.	3.4	121

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145	Functional antagonism between the estrogen receptor and Fos in the regulation of c-fos protooncogene transcription.. Molecular Endocrinology, 1993, 7, 1472-1483.	3.7	37
146	Functional antagonism between the estrogen receptor and Fos in the regulation of c-fos protooncogene transcription. Molecular Endocrinology, 1993, 7, 1472-1483.	3.7	21
147	Identification of a specific pattern of "immediate-early" gene activation induced by estrogen during mitogenic stimulation of rat uterine cells. Receptor, 1993, 3, 17-30.	0.8	39
148	Estrogen regulation of proto-oncogenes coding for nuclear proteins. Critical Reviews in Oncogenesis, 1993, 4, 361-88.	0.4	47
149	Transcriptional activation of jun and actin genes by estrogen during mitogenic stimulation of rat uterine cells. Journal of Steroid Biochemistry and Molecular Biology, 1992, 41, 523-528.	2.5	47
150	Estrogen induces c-fos expression specifically in the luminal and glandular epithelia of adult rat uterus. Biochemical and Biophysical Research Communications, 1991, 175, 480-485.	2.1	54
151	Inverse relationship between poly (ADP-ribose) polymerase activity and 2?,5?-oligoadenylates core level in estrogen-treated immature rat. Molecular and Cellular Biochemistry, 1990, 99, 33-39.	3.1	7
152	Identification of an estrogen response element upstream of the human c-fosgene that binds the estrogen receptor and the AP-1 transcription factor. Nucleic Acids Research, 1990, 18, 5097-5106.	14.5	293
153	Estrogen Stimulates Transcription of c-junProtooncogene. Molecular Endocrinology, 1990, 4, 1041-1050.	3.7	149
154	Activation of â€˜immediate-earlyâ€™ genes by estrogen is not sufficient to achieve stimulation of DNA synthesis in rat uterus. Biochemical and Biophysical Research Communications, 1990, 171, 287-292.	2.1	33
155	Estrogen Induces Expression of c-fosand c-mycProtooncogenes in Rat Uterus. Molecular Endocrinology, 1988, 2, 816-824.	3.7	221
156	Mineralcorticoid receptor from rat kidney. Interaction with heparin and purification to a CBG-free stage. The Journal of Steroid Biochemistry, 1986, 24, 461-467.	1.1	3
157	Specific binding of estrogen receptor to sites upstream and within the transcribed region of the chicken ovalbumin gene. Biochemical and Biophysical Research Communications, 1986, 139, 396-402.	2.1	15
158	Role of parathyroid hormone in the glucose intolerance of chronic renal failure.. Journal of Clinical Investigation, 1985, 75, 1037-1044.	8.2	126
159	Interaction of rat liver glucocorticoid receptor with heparin. Biochemistry, 1984, 23, 5393-5397.	2.5	11
160	Mechanisms of glucocorticoid hormone action. The Journal of Steroid Biochemistry, 1984, 20, 77-88.	1.1	70
161	Interaction of sodium molybdate with highly purified glucocorticoid receptor. The Journal of Steroid Biochemistry, 1984, 20, 289-293.	1.1	17
162	Estradiol and progesterone receptors in malignant gastrointestinal tumors. Cancer Research, 1984, 44, 4670-4.	0.9	114

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163	Steroid derivatives for electrophilic affinity labelling of glucocorticoid binding sites: Interaction with the glucocorticoid receptor and biological activity. The Journal of Steroid Biochemistry, 1983, 18, 375-382.	1.1	10
164	Competitive binding assay for glucocorticoids. Influence of experimental conditions on measurement of the affinity of competitive steroids for the receptor. The Journal of Steroid Biochemistry, 1982, 16, 515-520.	1.1	7
165	Assay of total estradiol receptor in tissue homogenate and tissue fractions by exchange with sodium thiocyanate at low temperature. Biochemistry, 1981, 20, 686-693.	2.5	53