

Pilar Blancafort

List of Publications by Year in descending order

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Version: 2024-02-01

64
papers

3,261
citations

136950

32
h-index

149698

56
g-index

65
all docs

65
docs citations

65
times ranked

4196
citing authors

#	ARTICLE	IF	CITATIONS
1	Breastmilk Is a Novel Source of Stem Cells with Multilineage Differentiation Potential. <i>Stem Cells</i> , 2012, 30, 2164-2174.	3.2	215
2	Epigenetic reprogramming of cancer cells via targeted DNA methylation. <i>Epigenetics</i> , 2012, 7, 350-360.	2.7	189
3	Development of Zinc Finger Domains for Recognition of the 5â€²-CNN-3â€² Family DNA Sequences and Their Use in the Construction of Artificial Transcription Factors. <i>Journal of Biological Chemistry</i> , 2005, 280, 35588-35597.	3.4	166
4	Systemic Delivery of Modified mRNA Encoding Herpes Simplex Virus 1 Thymidine Kinase for Targeted Cancer Gene Therapy. <i>Molecular Therapy</i> , 2013, 21, 358-367.	8.2	164
5	Designing Transcription Factor Architectures for Drug Discovery. <i>Molecular Pharmacology</i> , 2004, 66, 1361-1371.	2.3	162
6	Evaluation of a Modular Strategy for the Construction of Novel Polydactyl Zinc Finger DNA-Binding Proteins. <i>Biochemistry</i> , 2003, 42, 2137-2148.	2.5	161
7	Targeted silencing of the oncogenic transcription factor SOX2 in breast cancer. <i>Nucleic Acids Research</i> , 2012, 40, 6725-6740.	14.5	138
8	Scanning the human genome with combinatorial transcription factor libraries. <i>Nature Biotechnology</i> , 2003, 21, 269-274.	17.5	120
9	Synthetically controlling dendrimer flexibility improves delivery of large plasmid DNA. <i>Chemical Science</i> , 2017, 8, 2923-2930.	7.4	101
10	Re-activation of a dormant tumor suppressor gene maspin by designed transcription factors. <i>Oncogene</i> , 2007, 26, 2791-2798.	5.9	87
11	Honeybee venom and melittin suppress growth factor receptor activation in HER2-enriched and triple-negative breast cancer. <i>Npj Precision Oncology</i> , 2020, 4, 24.	5.4	86
12	Generation of tumor-initiating cells by exogenous delivery of OCT4transcription factor. <i>Breast Cancer Research</i> , 2011, 13, R94.	5.0	81
13	Gene expression in breastmilk cells is associated with maternal and infant characteristics. <i>Scientific Reports</i> , 2015, 5, 12933.	3.3	77
14	Novel role of Engrailed 1 as a prosurvival transcription factor in basal-like breast cancer and engineering of interference peptides block its oncogenic function. <i>Oncogene</i> , 2014, 33, 4767-4777.	5.9	76
15	Zinc Fingers, TALEs, and CRISPR Systems: A Comparison of Tools for Epigenome Editing. <i>Methods in Molecular Biology</i> , 2018, 1767, 19-63.	0.9	73
16	Stable oncogenic silencing in vivo by programmable and targeted de novo DNA methylation in breast cancer. <i>Oncogene</i> , 2015, 34, 5427-5435.	5.9	71
17	Activating PTEN Tumor Suppressor Expression with the CRISPR/dCas9 System. <i>Molecular Therapy - Nucleic Acids</i> , 2019, 14, 287-300.	5.1	68
18	Epigenome Engineering in Cancer: Fairytale or a Realistic Path to the Clinic?. <i>Frontiers in Oncology</i> , 2015, 5, 22.	2.8	63

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19	Waking up dormant tumor suppressor genes with zinc fingers, TALEs and the CRISPR/dCas9 system. <i>Oncotarget</i> , 2016, 7, 60535-60554.	1.8	61
20	Precision medicine by designer interference peptides: applications in oncology and molecular therapeutics. <i>Oncogene</i> , 2020, 39, 1167-1184.	5.9	61
21	Reprogramming epigenetic silencing: artificial transcription factors synergize with chromatin remodeling drugs to reactivate the tumor suppressor <i> mammary serine protease inhibitor</i>. <i>Molecular Cancer Therapeutics</i> , 2008, 7, 1080-1090.	4.1	58
22	Analysis of an artificial zinc finger epigenetic modulator: widespread binding but limited regulation. <i>Nucleic Acids Research</i> , 2014, 42, 10856-10868.	14.5	56
23	Tumor penetrating peptides inhibiting MYC as a potent targeted therapeutic strategy for triple-negative breast cancers. <i>Oncogene</i> , 2019, 38, 140-150.	5.9	55
24	Hallmarks of cancer: The CRISPR generation. <i>European Journal of Cancer</i> , 2018, 93, 10-18.	2.8	54
25	In Vivo Selection of Combinatorial Libraries and Designed Affinity Maturation of Polydactyl Zinc Finger Transcription Factors for ICAM-1 Provides New Insights into Gene Regulation. <i>Journal of Molecular Biology</i> , 2004, 341, 635-649.	4.2	49
26	Cisplatin Induces Differentiation of Breast Cancer Cells. <i>Frontiers in Oncology</i> , 2013, 3, 134.	2.8	48
27	Suppression of Breast Tumor Growth and Metastasis by an Engineered Transcription Factor. <i>PLoS ONE</i> , 2011, 6, e24595.	2.5	45
28	Genetic reprogramming of tumor cells by zinc finger transcription factors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 11716-11721.	7.1	44
29	Rab GTPases: Emerging Oncogenes and Tumor Suppressive Regulators for the Editing of Survival Pathways in Cancer. <i>Cancers</i> , 2020, 12, 259.	3.7	43
30	Reactivation of <i>MASPIN</i> in non-small cell lung carcinoma (NSCLC) cells by artificial transcription factors (ATFs). <i>Epigenetics</i> , 2011, 6, 224-235.	2.7	42
31	Targeting Serous Epithelial Ovarian Cancer with Designer Zinc Finger Transcription Factors. <i>Journal of Biological Chemistry</i> , 2012, 287, 29873-29886.	3.4	38
32	Tumour suppression by targeted intravenous non-viral CRISPRa using dendritic polymers. <i>Chemical Science</i> , 2019, 10, 7718-7727.	7.4	37
33	Triple-hit therapeutic approach for triple negative breast cancers using docetaxel nanoparticles, EN1-iPeps and RGD peptides. <i>Nanomedicine: Nanotechnology, Biology, and Medicine</i> , 2019, 20, 102003.	3.3	36
34	Epigenome engineering: new technologies for precision medicine. <i>Nucleic Acids Research</i> , 2020, 48, 12453-12482.	14.5	34
35	Re-expression of Selected Epigenetically Silenced Candidate Tumor Suppressor Genes in Cervical Cancer by TET2-directed Demethylation. <i>Molecular Therapy</i> , 2016, 24, 536-547.	8.2	33
36	Writing and Rewriting the Epigenetic Code of Cancer Cells: From Engineered Proteins to Small Molecules. <i>Molecular Pharmacology</i> , 2013, 83, 563-576.	2.3	30

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37	Crambescidin 800, Isolated from the Marine Sponge <i>Monanchora viridis</i> , Induces Cell Cycle Arrest and Apoptosis in Triple-Negative Breast Cancer Cells. <i>Marine Drugs</i> , 2018, 16, 53.	4.6	30
38	Expression of the Pluripotency Transcription Factor OCT4 in the Normal and Aberrant Mammary Gland. <i>Frontiers in Oncology</i> , 2013, 3, 79.	2.8	28
39	Promoter-targeted Phage Display Selections with Preassembled Synthetic Zinc Finger Libraries for Endogenous Gene Regulation. <i>Journal of Molecular Biology</i> , 2004, 340, 599-613.	4.2	26
40	Modulation of drug resistance by artificial transcription factors. <i>Molecular Cancer Therapeutics</i> , 2008, 7, 688-697.	4.1	22
41	Interrogating Genomes with Combinatorial Artificial Transcription Factor Libraries: Asking Zinc Finger Questions. <i>Assay and Drug Development Technologies</i> , 2006, 4, 317-331.	1.2	20
42	Breaking through an epigenetic wall. <i>Epigenetics</i> , 2013, 8, 164-176.	2.7	20
43	Aurantioside C Targets and Induces Apoptosis in Triple Negative Breast Cancer Cells. <i>Marine Drugs</i> , 2018, 16, 361.	4.6	19
44	The oncogene AAMDC links PI3K-AKT-mTOR signaling with metabolic reprogramming in estrogen receptor-positive breast cancer. <i>Nature Communications</i> , 2021, 12, 1920.	12.8	19
45	The recognition of a noncanonical RNA base pair by a zinc finger protein. <i>Chemistry and Biology</i> , 1999, 6, 585-597.	6.0	17
46	Rational Design, Selection and Specificity of Artificial Transcription Factors (ATFs): The Influence of Chromatin in Target Gene Regulation. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2008, 11, 146-158.	1.1	17
47	Sequence-Specific Biosensors Report Drug-Induced Changes in Epigenetic Silencing in Living Cells. <i>DNA and Cell Biology</i> , 2012, 31, S-2-S-10.	1.9	15
48	Reprogramming the anti-tumor immune response via CRISPR genetic and epigenetic editing. <i>Molecular Therapy - Methods and Clinical Development</i> , 2021, 21, 592-606.	4.1	11
49	Long live the stem cell: The use of stem cells isolated from post mortem tissues for translational strategies. <i>International Journal of Biochemistry and Cell Biology</i> , 2014, 56, 74-81.	2.8	10
50	Innovative Precision Gene Editing Tools in Personalized Cancer Medicine. <i>Advanced Science</i> , 2020, 7, 1902552.	11.2	9
51	Remodeling Genomes with Artificial Transcription Factors (ATFs). <i>Methods in Molecular Biology</i> , 2010, 649, 163-182.	0.9	9
52	Atomistic molecular dynamics simulations of bioactive engrailed 1 interference peptides (EN1-iPeps). <i>Oncotarget</i> , 2018, 9, 22383-22397.	1.8	9
53	Non-viral Methodology for Efficient Co-transfection. <i>Methods in Molecular Biology</i> , 2018, 1767, 241-254.	0.9	5
54	SP94-Targeted Nanoparticles Enhance the Efficacy of Sorafenib and Improve Liver Cancer Cell Discrimination. <i>ACS Applied Bio Materials</i> , 2021, 4, 1023-1029.	4.6	5

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55	Design and Characterization of a Cell-Penetrating Peptide Derived from the SOX2 Transcription Factor. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9354.	4.1	4
56	Manipulating the NKG2D Receptor-Ligand Axis Using CRISPR: Novel Technologies for Improved Host Immunity. <i>Frontiers in Immunology</i> , 2021, 12, 712722.	4.8	2
57	Poll-driven integrative expression vectors for yeast. <i>Journal of Biotechnology</i> , 1997, 56, 41-47.	3.8	1
58	A peptide-functionalised dendronised polymer for selective transfection in human liver cancer cells. <i>New Journal of Chemistry</i> , 2021, 45, 19315-19320.	2.8	1
59	Breastmilk Stem Cells: Recent Advances and Future Prospects. , 2015, , 185-195.		1
60	Promoter-targeted Phage Display Selections with Preassembled Synthetic Zinc Finger Libraries for Endogenous Gene Regulation. <i>Journal of Molecular Biology</i> , 2004, 340, 599-599.	4.2	0
61	Engineering Transcription Factors in Breast Cancer Stem Cells. , 2011, , .		0
62	C9a-induced epigenetic silencing of maspin in human claudin-low breast tumor initiating cells. <i>FASEB Journal</i> , 2011, 25, 122.6.	0.5	0
63	Breastmilk and the lactating breast as a tool to elucidate breast cancer. <i>FASEB Journal</i> , 2013, 27, 629.9.	0.5	0
64	The CRISPR road: from bench to bedside on an RNA-guided path. <i>Annals of Translational Medicine</i> , 2015, 3, 174.	1.7	0