Mark D Long

List of Publications by Year in descending order

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471509 454955 1,079 38 17 30 citations h-index g-index papers 41 41 41 1431 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	The Genomic Impact of DNA CpG Methylation on Gene Expression; Relationships in Prostate Cancer. Biomolecules, 2017, 7, 15.	4.0	92
2	MUC1-C regulates lineage plasticity driving progression to neuroendocrine prostate cancer. Nature Communications, 2020, 11, 338.	12.8	87
3	Overcoming primary and acquired resistance to anti-PD-L1 therapy by induction and activation of tumor-residing cDC1s. Nature Communications, 2020, 11, 5415.	12.8	85
4	Serum microRNA expression patterns that predict early treatment failure in prostate cancer patients. Oncotarget, 2014, 5, 824-840.	1.8	52
5	Activation of NF- $\hat{\mathbb{P}}$ B and p300/CBP potentiates cancer chemoimmunotherapy through induction of MHC-I antigen presentation. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	47
6	MUC1-C Activates the BAF (mSWI/SNF) Complex in Prostate Cancer Stem Cells. Cancer Research, 2021, 81, 1111-1122.	0.9	46
7	The miR-96 and RAR \hat{I}^3 signaling axis governs androgen signaling and prostate cancer progression. Oncogene, 2019, 38, 421-444.	5.9	45
8	MUC1-C Activates the NuRD Complex to Drive Dedifferentiation of Triple-Negative Breast Cancer Cells. Cancer Research, 2019, 79, 5711-5722.	0.9	45
9	MUC1-C integrates activation of the IFN- \hat{l}^3 pathway with suppression of the tumor immune microenvironment in triple-negative breast cancer., 2021, 9, e002115.		41
10	MUC1-C activates the PBAF chromatin remodeling complex in integrating redox balance with progression of human prostate cancer stem cells. Oncogene, 2021, 40, 4930-4940.	5.9	41
11	Pan-Cancer Analyses of the Nuclear Receptor Superfamily. Nuclear Receptor Research, 2015, 2, .	2.5	40
12	Pharmacological polyamine catabolism upregulation with methionine salvage pathway inhibition as an effective prostate cancer therapy. Nature Communications, 2020, 11, 52.	12.8	37
13	MUC1-C drives stemness in progression of colitis to colorectal cancer. JCI Insight, 2020, 5, .	5.0	36
14	Vitamin D Receptor and RXR in the Postâ€Genomic Era. Journal of Cellular Physiology, 2015, 230, 758-766.	4.1	35
15	Integration of VDR genome wide binding and GWAS genetic variation data reveals co-occurrence of VDR and NF-κB binding that is linked to immune phenotypes. BMC Genomics, 2017, 18, 132.	2.8	35
16	Inhibition of the aryl hydrocarbon receptor/polyamine biosynthesis axis suppresses multiple myeloma. Journal of Clinical Investigation, 2018, 128, 4682-4696.	8.2	35
17	CD8 ⁺ T cell immunity blocks the metastasis of carcinogen-exposed breast cancer. Science Advances, 2021, 7, .	10.3	24
18	Integrative genomic analysis in K562 chronic myelogenous leukemia cells reveals that proximal NCOR1 binding positively regulates genes that govern erythroid differentiation and Imatinib sensitivity. Nucleic Acids Research, 2015, 43, 7330-7348.	14.5	22

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19	Cooperative behavior of the nuclear receptor superfamily and its deregulation in prostate cancer. Carcinogenesis, 2014, 35, 262-271.	2.8	19
20	Identification of transcription factor co-regulators that drive prostate cancer progression. Scientific Reports, 2020, 10, 20332.	3.3	19
21	Reduced NCOR2 expression accelerates androgen deprivation therapy failure in prostate cancer. Cell Reports, 2021, 37, 110109.	6.4	19
22	MUC1-C represses the RASSF1A tumor suppressor in human carcinoma cells. Oncogene, 2019, 38, 7266-7277.	5.9	17
23	MUC1-C Dictates JUN and BAF-Mediated Chromatin Remodeling at Enhancer Signatures in Cancer Stem Cells. Molecular Cancer Research, 2022, 20, 556-567.	3.4	17
24	MUC1-C integrates type II interferon and chromatin remodeling pathways in immunosuppression of prostate cancer. Oncolmmunology, 2022, 11, 2029298.	4.6	17
25	Integrative genomic approaches to dissect clinically-significant relationships between the VDR cistrome and gene expression in primary colon cancer. Journal of Steroid Biochemistry and Molecular Biology, 2017, 173, 130-138.	2.5	14
26	miRNAs as drivers of TMPRSS2-ERG negative prostate tumors in African American men. Frontiers in Bioscience - Landmark, 2017, 22, 212-229.	3.0	14
27	Generation of cDC-like cells from human induced pluripotent stem cells via Notch signaling. , 2022, 10, e003827.		14
28	Targeting MUC1-C Suppresses Chronic Activation of Cytosolic Nucleotide Receptors and STING in Triple-Negative Breast Cancer. Cancers, 2022, 14, 2580.	3.7	14
29	Dietary folate levels alter the kinetics and molecular mechanism of prostate cancer recurrence in the CWR22 model. Oncotarget, 2017, 8, 103758-103774.	1.8	13
30	Wholeâ€exome sequencing of ovarian cancer families uncovers putative predisposition genes. International Journal of Cancer, 2020, 146, 2147-2155.	5.1	12
31	Induction of cell death in ovarian cancer cells by doxorubicin and oncolytic vaccinia virus is associated with CREB3L1 activation. Molecular Therapy - Oncolytics, 2021, 23, 38-50.	4.4	9
32	Dependence on the MUC1-C Oncoprotein in Classic, Variant, and Non–neuroendocrine Small Cell Lung Cancer. Molecular Cancer Research, 2022, 20, 1379-1390.	3.4	8
33	Positionâ€Scanning Peptide Libraries as Particle Immunogens for Improving CD8 + Tâ€Cell Responses. Advanced Science, 2021, , 2103023.	11.2	5
34	An <i>In Vivo</i> Screen to Identify Short Peptide Mimotopes with Enhanced Antitumor Immunogenicity. Cancer Immunology Research, 2022, 10, 314-326.	3.4	5
35	Dynamic patterns of DNA methylation in the normal prostate epithelial differentiation program are targets of aberrant methylation in prostate cancer. Scientific Reports, 2021, 11, 11405.	3.3	3
36	Concurrent Aspirin Use Is Associated with Improved Outcome in Rectal Cancer Patients Who Undergo Chemoradiation Therapy. Cancers, 2021, 13, 205.	3.7	3

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37	Comparable transforming growth factor beta-mediated immune suppression in ex vivo-expanded natural killer cells from cord blood and peripheral blood: implications for adoptive immunotherapy. Cytotherapy, 2022, 24, 802-817.	0.7	2
38	Whole-Genome Sequencing Identifies PPARGC1A as a Putative Modifier of Cancer Risk in BRCA1/2 Mutation Carriers. Cancers, 2022, 14, 2350.	3.7	1