

Crystal N Marconett

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

1,494
citations

361388

20
h-index

361001

35
g-index

45
all docs

45
docs citations

45
times ranked

3665
citing authors

#	ARTICLE	IF	CITATIONS
1	Development of human alveolar epithelial cell models to study distal lung biology and disease. <i>IScience</i> , 2022, 25, 103780.	4.1	15
2	FOXO1 Couples KGF and PI-3K/AKT Signaling to NKX2.1-Regulated Differentiation of Alveolar Epithelial Cells. <i>Cells</i> , 2022, 11, 1122.	4.1	5
3	The Sulfotransferase SULT1C2 Is Epigenetically Activated and Transcriptionally Induced by Tobacco Exposure and Is Associated with Patient Outcome in Lung Adenocarcinoma. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 416.	2.6	3
4	The evolutionarily conserved long non-coding RNA <i>LINC00261</i> drives neuroendocrine prostate cancer proliferation and metastasis via distinct nuclear and cytoplasmic mechanisms. <i>Molecular Oncology</i> , 2021, 15, 1921-1941.	4.6	22
5	Beyond the transcription factor: the under-studied role of epigenomics in lung differentiation. <i>Epigenomics</i> , 2021, 13, 1845-1848.	2.1	0
6	Comprehensive epigenomic profiling of human alveolar epithelial differentiation identifies key epigenetic states and transcription factor co-regulatory networks for maintenance of distal lung identity. <i>BMC Genomics</i> , 2021, 22, 906.	2.8	10
7	Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020, 112, 1223-1232.	2.9	15
8	Integrated Single-Cell RNA-Sequencing Analysis of Aquaporin 5-Expressing Mouse Lung Epithelial Cells Identifies GPRC5A as a Novel Validated Type I Cell Surface Marker. <i>Cells</i> , 2020, 9, 2460.	4.1	8
9	TENET 2.0: Identification of key transcriptional regulators and enhancers in lung adenocarcinoma. <i>PLoS Genetics</i> , 2020, 16, e1009023.	3.5	20
10	Genome-wide integration of microRNA and transcriptomic profiles of differentiating human alveolar epithelial cells. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2020, 319, L173-L184.	2.9	8
11	PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence. <i>PLoS ONE</i> , 2020, 15, e0243791.	2.5	12
12	Title is missing!. , 2020, 15, e0243791.		0
13	Title is missing!. , 2020, 15, e0243791.		0
14	Title is missing!. , 2020, 15, e0243791.		0
15	Title is missing!. , 2020, 15, e0243791.		0
16	<i>LINC00261</i> Is an Epigenetically Regulated Tumor Suppressor Essential for Activation of the DNA Damage Response. <i>Cancer Research</i> , 2019, 79, 3050-3062.	0.9	75
17	Positional integration of lung adenocarcinoma susceptibility loci with primary human alveolar epithelial cell epigenomes. <i>Epigenomics</i> , 2018, 10, 1167-1187.	2.1	14
18	CLDN18.1 attenuates malignancy and related signaling pathways of lung adenocarcinoma <i>in vivo</i> and <i>in vitro</i> . <i>International Journal of Cancer</i> , 2018, 143, 3169-3180.	5.1	20

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19	Claudin-18-mediated YAP activity regulates lung stem and progenitor cell homeostasis and tumorigenesis. <i>Journal of Clinical Investigation</i> , 2018, 128, 970-984.	8.2	115
20	Epigenome-wide analysis of DNA methylation in lung tissue shows concordance with blood studies and identifies tobacco smoke-inducible enhancers. <i>Human Molecular Genetics</i> , 2017, 26, 3014-3027.	2.9	97
21	Cross-Species Transcriptome Profiling Identifies New Alveolar Epithelial Type I Cell-Specific Genes. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2017, 56, 310-321.	2.9	27
22	Intersecting transcriptomic profiling technologies and long non-coding RNA function in lung adenocarcinoma: discovery, mechanisms, and therapeutic applications. <i>Oncotarget</i> , 2017, 8, 81538-81557.	1.8	21
23	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , 2016, 26, ddw414.	2.9	50
24	Pleiotropic Analysis of Lung Cancer and Blood Triglycerides. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw167.	6.3	17
25	The importance of detailed epigenomic profiling of different cell types within organs. <i>Epigenomics</i> , 2016, 8, 817-829.	2.1	22
26	Functional lung adenocarcinoma risk SNPs identified through positional integration with human alveolar epithelial cell epigenomes. <i>Journal of Thoracic Oncology</i> , 2016, 11, S46.	1.1	0
27	The 78-kD Glucose-Regulated Protein Regulates Endoplasmic Reticulum Homeostasis and Distal Epithelial Cell Survival during Lung Development. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2016, 55, 135-149.	2.9	36
28	Combinations of differentiation markers distinguish subpopulations of alveolar epithelial cells in adult lung. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2016, 310, L114-L120.	2.9	43
29	Somatic Genomics and Clinical Features of Lung Adenocarcinoma: A Retrospective Study. <i>PLoS Medicine</i> , 2016, 13, e1002162.	8.4	148
30	Claudin 4 knockout mice: normal physiological phenotype with increased susceptibility to lung injury. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2014, 307, L524-L536.	2.9	70
31	Knockout Mice Reveal Key Roles for Claudin 18 in Alveolar Barrier Properties and Fluid Homeostasis. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2014, 51, 210-222.	2.9	70
32	Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. <i>Nature Communications</i> , 2014, 5, 3365.	12.8	123
33	Transcriptomic profiling of primary alveolar epithelial cell differentiation in human and rat. <i>Genomics Data</i> , 2014, 2, 105-109.	1.3	6
34	Abstract 287: Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. , 2014, , .		0
35	Integrated Transcriptomic and Epigenomic Analysis of Primary Human Lung Epithelial Cell Differentiation. <i>PLoS Genetics</i> , 2013, 9, e1003513.	3.5	46
36	CDKN2A/p16 Inactivation Mechanisms and Their Relationship to Smoke Exposure and Molecular Features in Non-Small-Cell Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2013, 8, 1378-1388.	1.1	71

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37	Indole-3-Carbinol disrupts Estrogen Receptor-alpha dependent expression of Insulin-like Growth Factor-1 Receptor and Insulin Receptor Substrate-1 and proliferation of human breast cancer cells. <i>Molecular and Cellular Endocrinology</i> , 2012, 363, 74-84.	3.2	40
38	HDAC3 And GATA-6/p300 Coordinately Regulate Type I Cell-Specific Aquaporin-5 (Aqp5) Gene Expression During Alveolar Epithelial Cell (AEC) Transdifferentiation. , 2011, , .		0
39	Indole-3-carbinol downregulation of telomerase gene expression requires the inhibition of estrogen receptor-alpha and Sp1 transcription factor interactions within the hTERT promoter and mediates the G1 cell cycle arrest of human breast cancer cells. <i>Carcinogenesis</i> , 2011, 32, 1315-1323.	2.8	55
40	1-Benzyl-indole-3-carbinol is a novel indole-3-carbinol derivative with significantly enhanced potency of anti-proliferative and anti-estrogenic properties in human breast cancer cells. <i>Chemico-Biological Interactions</i> , 2010, 186, 255-266.	4.0	34
41	Indole-3-Carbinol Triggers Aryl Hydrocarbon Receptor-dependent Estrogen Receptor (ER)± Protein Degradation in Breast Cancer Cells Disrupting an ER±-GATA3 Transcriptional Cross-Regulatory Loop. <i>Molecular Biology of the Cell</i> , 2010, 21, 1166-1177.	2.1	67
42	BZL101, a phytochemical extract from the <i>Scutellaria barbata</i> plant, disrupts proliferation of human breast and prostate cancer cells through distinct mechanisms dependent on the cancer cell phenotype. <i>Cancer Biology and Therapy</i> , 2010, 10, 397-405.	3.4	55
43	Artemisinin selectively decreases functional levels of estrogen receptor-alpha and ablates estrogen-induced proliferation in human breast cancer cells. <i>Carcinogenesis</i> , 2008, 29, 2252-2258.	2.8	52