

Jill Everland Larsen

List of Publications by Year in descending order

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Version: 2024-02-01

46
papers

3,426
citations

201674

27
h-index

265206

42
g-index

47
all docs

47
docs citations

47
times ranked

6761
citing authors

#	ARTICLE	IF	CITATIONS
1	Effect of KRAS Oncogene Substitutions on Protein Behavior: Implications for Signaling and Clinical Outcome. <i>Journal of the National Cancer Institute</i> , 2012, 104, 228-239.	6.3	424
2	Aldehyde Dehydrogenase Activity Selects for Lung Adenocarcinoma Stem Cells Dependent on Notch Signaling. <i>Cancer Research</i> , 2010, 70, 9937-9948.	0.9	357
3	ZEB1 drives epithelial-to-mesenchymal transition in lung cancer. <i>Journal of Clinical Investigation</i> , 2016, 126, 3219-3235.	8.2	256
4	ASCL1 is a lineage oncogene providing therapeutic targets for high-grade neuroendocrine lung cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 14788-14793.	7.1	205
5	Molecular Biology of Lung Cancer: Clinical Implications. <i>Clinics in Chest Medicine</i> , 2011, 32, 703-740.	2.1	194
6	Human Lung Epithelial Cells Progressed to Malignancy through Specific Oncogenic Manipulations. <i>Molecular Cancer Research</i> , 2013, 11, 638-650.	3.4	192
7	Knockdown of Oncogenic KRAS in Non-Small Cell Lung Cancers Suppresses Tumor Growth and Sensitizes Tumor Cells to Targeted Therapy. <i>Molecular Cancer Therapeutics</i> , 2011, 10, 336-346.	4.1	151
8	Systematic Identification of Molecular Subtype-Selective Vulnerabilities in Non-Small-Cell Lung Cancer. <i>Cell</i> , 2013, 155, 552-566.	28.9	151
9	Exosomes derived from mesenchymal non-small cell lung cancer cells promote chemoresistance. <i>International Journal of Cancer</i> , 2017, 141, 614-620.	5.1	117
10	Gene Expression Signature Predicts Recurrence in Lung Adenocarcinoma. <i>Clinical Cancer Research</i> , 2007, 13, 2946-2954.	7.0	107
11	Pax gene diversity in the basal cnidarian <i>Acropora millepora</i> (Cnidaria, Anthozoa): Implications for the evolution of the Pax gene family. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 4475-4480.	7.1	102
12	MicroRNA-218 Is Deleted and Downregulated in Lung Squamous Cell Carcinoma. <i>PLoS ONE</i> , 2010, 5, e12560.	2.5	100
13	Expression profiling defines a recurrence signature in lung squamous cell carcinoma. <i>Carcinogenesis</i> , 2006, 28, 760-766.	2.8	98
14	Targeted Therapies for Lung Cancer. <i>Cancer Journal (Sudbury, Mass)</i> , 2011, 17, 512-527.	2.0	91
15	NeuroD1 regulates survival and migration of neuroendocrine lung carcinomas via signaling molecules TrkB and NCAM. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6524-6529.	7.1	84
16	Meta- and Pooled Analysis of GSTP1 Polymorphism and Lung Cancer: A HuGE-GSEC Review. <i>American Journal of Epidemiology</i> , 2009, 169, 802-814.	3.4	73
17	CYP1A1 Ile462Val and MPO G-463A interact to increase risk of adenocarcinoma but not squamous cell carcinoma of the lung. <i>Carcinogenesis</i> , 2006, 27, 525-532.	2.8	66
18	SMAC Mimetic (JP1201) Sensitizes Non-Small Cell Lung Cancers to Multiple Chemotherapy Agents in an IAP-Dependent but TNF-Independent Manner. <i>Cancer Research</i> , 2011, 71, 7640-7648.	0.9	55

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19	PROTOCOLADHERIN 7 Acts through SET and PP2A to Potentiate MAPK Signaling by EGFR and KRAS during Lung Tumorigenesis. <i>Cancer Research</i> , 2017, 77, 187-197.	0.9	55
20	Myeloperoxidase G-463A polymorphism and lung cancer: A HuGE Genetic Susceptibility to Environmental Carcinogens pooled analysis. <i>Genetics in Medicine</i> , 2007, 9, 67-73.	2.4	47
21	Genetic association study of CYP1A1 polymorphisms identifies risk haplotypes in nonsmall cell lung cancer. <i>European Respiratory Journal</i> , 2010, 35, 152-159.	6.7	44
22	Combination Therapy Targeting BCL6 and Phospho-STAT3 Defeats Intratumor Heterogeneity in a Subset of Non-Small Cell Lung Cancers. <i>Cancer Research</i> , 2017, 77, 3070-3081.	0.9	36
23	Expression profiling identifies genes involved in emphysema severity. <i>Respiratory Research</i> , 2009, 10, 81.	3.6	35
24	Systematic siRNA Screen Unmasks NSCLC Growth Dependence by Palmitoyltransferase DHHC5. <i>Molecular Cancer Research</i> , 2015, 13, 784-794.	3.4	35
25	Cancer-Specific Production of N-Acetylaspartate via NAT8L Overexpression in Non-Small Cell Lung Cancer and Its Potential as a Circulating Biomarker. <i>Cancer Prevention Research</i> , 2016, 9, 43-52.	1.5	33
26	Risk of non-small cell lung cancer and the cytochrome P4501A1 Ile462Val polymorphism. <i>Cancer Causes and Control</i> , 2005, 16, 579-585.	1.8	31
27	Ras regulates kinesin 13 family members to control cell migration pathways in transformed human bronchial epithelial cells. <i>Oncogene</i> , 2014, 33, 5457-5466.	5.9	29
28	Array-Comparative Genomic Hybridization Reveals Loss of SOCS6 Is Associated with Poor Prognosis in Primary Lung Squamous Cell Carcinoma. <i>PLoS ONE</i> , 2012, 7, e30398.	2.5	28
29	Genes and Gene Ontologies Common to Airflow Obstruction and Emphysema in the Lungs of Patients with COPD. <i>PLoS ONE</i> , 2011, 6, e17442.	2.5	26
30	ADAM28: A potential oncogene involved in asbestos-related lung adenocarcinomas. <i>Genes Chromosomes and Cancer</i> , 2010, 49, 688-698.	2.8	24
31	Epithelial-mesenchymal transition increases tumor sensitivity to COX-2 inhibition by apicoxib. <i>Carcinogenesis</i> , 2012, 33, 1639-1646.	2.8	24
32	NeuroD1 regulation of migration accompanies the differential sensitivity of neuroendocrine carcinomas to TrkB inhibition. <i>Oncogenesis</i> , 2013, 2, e63-e63.	4.9	21
33	An in vivo functional genomics screen of nuclear receptors and their co-regulators identifies FOXA1 as an essential gene in lung tumorigenesis. <i>Neoplasia</i> , 2020, 22, 294-310.	5.3	21
34	Gene expression of lung squamous cell carcinoma reflects mode of lymph node involvement. <i>European Respiratory Journal</i> , 2007, 30, 21-26.	6.7	20
35	Genomic medicine in non-small cell lung cancer: Paving the path to personalized care. <i>Respirology</i> , 2011, 16, 257-263.	2.3	18
36	Combined use of subclinical hydroxyurea and CHK1 inhibitor effectively controls melanoma and lung cancer progression, with reduced normal tissue toxicity compared to gemcitabine. <i>Molecular Oncology</i> , 2019, 13, 1503-1518.	4.6	17

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37	Silencing the Snail-Dependent RNA Splice Regulator ESRP1 Drives Malignant Transformation of Human Pulmonary Epithelial Cells. <i>Cancer Research</i> , 2018, 78, 1986-1999.	0.9	13
38	Nuclear Receptor Expression and Function in Human Lung Cancer Pathogenesis. <i>PLoS ONE</i> , 2015, 10, e0134842.	2.5	12
39	Validation of SCT Methylation as a Hallmark Biomarker for Lung Cancers. <i>Journal of Thoracic Oncology</i> , 2016, 11, 346-360.	1.1	11
40	Identification of a Human Airway Epithelial Cell Subpopulation with Altered Biophysical, Molecular, and Metastatic Properties. <i>Cancer Prevention Research</i> , 2017, 10, 514-524.	1.5	9
41	Loss of miR125a Expression in a Model of K-ras-Dependent Pulmonary Premalignancy. <i>Cancer Prevention Research</i> , 2014, 7, 845-855.	1.5	5
42	Re: Effects of N-(4-Hydroxy-phenyl)retinamide on hTERT Expression in the Bronchial Epithelium of Cigarette Smokers. <i>Journal of the National Cancer Institute</i> , 2002, 94, 949-a-950.	6.3	4
43	Molecular Basis of Lung Carcinogenesis. , 2017, , 447-496.		4
44	Molecular Basis of Lung Cancer. , 2015, , 475-490.e1.		1
45	Abstract 1855: SCT methylation is a potential cancer biomarker for lung cancer. , 2014, , .		0
46	Abstract A22: Differential MYC dependence in NSCLC identified through pharmacological and genetic MYC inhibition. , 2015, , .		0