

# Jin Eun Choi

## List of Publications by Year in descending order

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49  
papers

865  
citations

471509  
17  
h-index

501196  
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49  
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docs citations

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times ranked

2017  
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#	ARTICLE	IF	CITATIONS
1	Epigenetic readers and lung cancer: the rs2427964C>T variant of the bromodomain and extraterminal domain gene <i>BRD3</i> is associated with poorer survival outcome in NSCLC. <i>Molecular Oncology</i> , 2022, 16, 750-763.	4.6	1
2	Prognostic significance of genetic variants in GLUT1 in stage III non-small cell lung cancer treated with radiotherapy. <i>Thoracic Cancer</i> , 2021, 12, 874-879.	1.9	2
3	Genetic Polymorphisms in Activating Transcription Factor 3 Binding Site and the Prognosis of Early-Stage Non-Small Cell Lung Cancer. <i>Oncology</i> , 2021, 99, 336-344.	1.9	1
4	CD5L as an Extracellular Vesicle-Derived Biomarker for Liquid Biopsy of Lung Cancer. <i>Diagnostics</i> , 2021, 11, 620.	2.6	33
5	Nuclear Pore Glycoprotein 62 Genetic Variant rs9523 is Associated with Clinical Outcomes of Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitors in Lung Adenocarcinoma Patients. <i>Pharmacogenomics and Personalized Medicine</i> , 2021, Volume 14, 1291-1302.	0.7	0
6	Impact of immune checkpoint gene CD155 Ala67Thr and CD226 Gly307Ser polymorphisms on small cell lung cancer clinical outcome. <i>Scientific Reports</i> , 2021, 11, 1794.	3.3	3
7	Prognostic implication of PD-L1 polymorphisms in non-small cell lung cancer treated with radiotherapy. <i>Cancer Medicine</i> , 2021, 10, 8071-8078.	2.8	3
8	Genetic variants in histone modification regions are associated with the prognosis of lung adenocarcinoma. <i>Scientific Reports</i> , 2021, 11, 21520.	3.3	5
9	Polymorphism in ASCL1 target gene DDC is associated with clinical outcomes of small cell lung cancer patients. <i>Thoracic Cancer</i> , 2020, 11, 19-28.	1.9	7
10	Effect of genetic variation in Notch regulator DTX1 on SCLC prognosis compared with the effect on NSCLC prognosis. <i>Thoracic Cancer</i> , 2020, 11, 2698-2703.	1.9	2
11	Expression of key regulatory genes in necroptosis and its effect on the prognosis in non-small cell lung cancer. <i>Journal of Cancer</i> , 2020, 11, 5503-5510.	2.5	32
12	Genetic Variants in One-Carbon Metabolism Pathway Predict Survival Outcomes of Early-Stage Non-Small Cell Lung Cancer. <i>Oncology</i> , 2020, 98, 897-904.	1.9	2
13	Polymorphisms in Glycolysis-Related Genes Are Associated with Clinical Outcomes of Paclitaxel-Cisplatin Chemotherapy in Non-Small Cell Lung Cancer. <i>Oncology</i> , 2020, 98, 468-477.	1.9	5
14	The effect of susceptibility variants, identified in never-smoking female lung cancer cases, on male smokers. <i>Korean Journal of Internal Medicine</i> , 2020, 35, 929-935.	1.7	3
15	Genetic Variant of Notch Regulator DTX1 Predicts Survival After Lung Cancer Surgery. <i>Annals of Surgical Oncology</i> , 2019, 26, 3756-3764.	1.5	4
16	Glucose transporter 3 gene variant is associated with survival outcome of patients with non-small cell lung cancer after surgical resection. <i>Gene</i> , 2019, 703, 58-64.	2.2	5
17	<i>TSC2</i> genetic variant and prognosis in non-small cell lung cancer after curative surgery. <i>Thoracic Cancer</i> , 2019, 10, 335-340.	1.9	5
18	Regulatory variants in cancer-related pathway genes predict survival of patients with surgically resected non-small cell lung cancer. <i>Gene</i> , 2018, 646, 56-63.	2.2	3

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19	Functional intronic variant of <i>SLC5A10</i> affects <i>DRG2</i> expression and survival outcomes of early-stage non-small cell lung cancer. <i>Cancer Science</i> , 2018, 109, 3902-3909.	3.9	22
20	An expression quantitative trait locus variant for <i>LKB1</i> gene predicts the clinical outcomes of chemotherapy in patients with non-small cell lung cancer. <i>Cancer Genetics</i> , 2018, 228-229, 73-82.	0.4	3
21	Intronic variant of <i>EGFR</i> is associated with GBAS expression and survival outcome of early-stage non-small cell lung cancer. <i>Thoracic Cancer</i> , 2018, 9, 916-923.	1.9	9
22	Glucose Transporter 1 Gene Variants Predict the Prognosis of Patients with Early-Stage Non-small Cell Lung Cancer. <i>Annals of Surgical Oncology</i> , 2018, 25, 3396-3403.	1.5	8
23	Effects of polymorphisms identified in genome-wide association studies of never-smoking females on the prognosis of non-small cell lung cancer. <i>Cancer Genetics</i> , 2017, 212-213, 8-12.	0.4	7
24	Development of a prognosis prediction model incorporating genetic polymorphism with pathologic stage in stage I non-small cell lung cancer: <i>A</i> multicenter study. <i>Thoracic Cancer</i> , 2017, 8, 251-259.	1.9	3
25	Association between polymorphisms in microRNA target sites and survival in early-stage non-small cell lung cancer. <i>Thoracic Cancer</i> , 2017, 8, 682-686.	1.9	6
26	Polymorphisms in Epithelial-Mesenchymal Transition-Related Genes and the Prognosis of Surgically Treated Non-small Cell Lung Cancer. <i>Annals of Surgical Oncology</i> , 2017, 24, 3386-3395.	1.5	5
27	Functional polymorphisms in <i>PD-L1</i> gene are associated with the prognosis of patients with early stage non-small cell lung cancer. <i>Gene</i> , 2017, 599, 28-35.	2.2	47
28	Transcriptome analysis of non-small cell lung cancer and genetically matched adjacent normal tissues identifies novel prognostic marker genes. <i>Genes and Genomics</i> , 2017, 39, 277-284.	1.4	6
29	Polymorphisms in mitotic checkpoint-related genes can influence survival outcomes of early-stage non-small cell lung cancer. <i>Oncotarget</i> , 2017, 8, 61777-61785.	1.8	7
30	Genetic Variants in the Wnt Signaling Pathway Are Not Associated with Survival Outcome of Non-Small Cell Lung Cancer in a Korean Population. <i>Journal of Korean Medical Science</i> , 2016, 31, 463.	2.5	1
31	The Different Effect of <i>VEGF</i> Polymorphisms on the Prognosis of Non-Small Cell Lung Cancer according to Tumor Histology. <i>Journal of Korean Medical Science</i> , 2016, 31, 1735.	2.5	6
32	<i>PD-L1</i> polymorphism can predict clinical outcomes of non-small cell lung cancer patients treated with first-line paclitaxel-cisplatin chemotherapy. <i>Scientific Reports</i> , 2016, 6, 25952.	3.3	36
33	Polymorphisms in cancer-related pathway genes and lung cancer. <i>European Respiratory Journal</i> , 2016, 48, 1184-1191.	6.7	5
34	Genetic polymorphisms in glycolytic pathway are associated with the prognosis of patients with early stage non-small cell lung cancer. <i>Scientific Reports</i> , 2016, 6, 35603.	3.3	31
35	<i>TERT</i> Polymorphism rs2853669 Influences on Lung Cancer Risk in the Korean Population. <i>Journal of Korean Medical Science</i> , 2015, 30, 1423.	2.5	23
36	A Panel of Genetic Polymorphism for the Prediction of Prognosis in Patients with Early Stage Non-Small Cell Lung Cancer after Surgical Resection. <i>PLoS ONE</i> , 2015, 10, e0140216.	2.5	11

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37	Putative functional variants of XRCC1 identified by RegulomeDB were not associated with lung cancer risk in a Korean population. <i>Cancer Genetics</i> , 2015, 208, 19-24.	0.4	33
38	Association between GWAS-Identified Genetic Variations and Disease Prognosis for Patients with Colorectal Cancer. <i>PLoS ONE</i> , 2015, 10, e0119649.	2.5	20
39	<i>RACK1</i> is a candidate gene associated with the prognosis of patients with early stage non-small cell lung cancer. <i>Oncotarget</i> , 2015, 6, 4451-4466.	1.8	15
40	Functional intronic ERCC1 polymorphism from regulomeDB can predict survival in lung cancer after surgery. <i>Oncotarget</i> , 2015, 6, 24522-24532.	1.8	24
41	Replication of the results of genome-wide and candidate gene association studies on telomere length in a Korean population. <i>Korean Journal of Internal Medicine</i> , 2015, 30, 719-726.	1.7	24
42	Predictive Efficacy of Low Burden EGFR Mutation Detected by Next-Generation Sequencing on Response to EGFR Tyrosine Kinase Inhibitors in Non-Small-Cell Lung Carcinoma. <i>PLoS ONE</i> , 2013, 8, e81975.	2.5	18
43	MicroRNA Expression Profiles in Korean Non-Small Cell Lung Cancer. <i>Tuberculosis and Respiratory Diseases</i> , 2009, 67, 413.	1.8	13
44	Comprehensive assessment of P21 polymorphisms and lung cancer risk. <i>Journal of Human Genetics</i> , 2008, 53, 87-95.	2.3	22
45	Telomere length and the risk of lung cancer. <i>Cancer Science</i> , 2008, 99, 1385-1389.	3.9	177
46	Aberrant methylation of ADAMTS1 in non-small cell lung cancer. <i>Cancer Genetics and Cytogenetics</i> , 2008, 187, 80-84.	1.0	31
47	Polymorphisms in the survivin gene and the risk of lung cancer. <i>Lung Cancer</i> , 2008, 60, 31-39.	2.0	98
48	Polymorphisms in the SERPINA1 Gene and the Risk of Chronic Obstructive Pulmonary Disease in a Korean Population. <i>Tuberculosis and Respiratory Diseases</i> , 2008, 65, 285.	1.8	0
49	Polymorphisms in the epidermal growth factor receptor gene and the risk of primary lung cancer: a case-control study. <i>BMC Cancer</i> , 2007, 7, 199.	2.6	38