

Lung cancer susceptibility locus at 5p15.33

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Citation Report

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Phase I Metabolic Genes and Risk of Lung Cancer: Multiple Polymorphisms and mRNA Expression. PLoS ONE, 2009, 4, e5652. | 1.1 | 91 |
| 2 | A Spectrum of Severe Familial Liver Disorders Associate with Telomerase Mutations. PLoS ONE, 2009, 4, e7926. | 1.1 | 201 |
| 3 | <i>TGFBR1</i> Haplotypes and Risk of Non-Small-Cell Lung Cancer. Cancer Research, 2009, 69, 7046-7052. | 0.4 | 24 |
| 4 | Common genetic variants on 5p15.33 contribute to risk of lung adenocarcinoma in a Chinese population. Carcinogenesis, 2009, 30, 987-990. | 1.3 | 72 |
| 5 | The TERT-CLPTM1L lung cancer susceptibility variant associates with higher DNA adduct formation in the lung. Carcinogenesis, 2009, 30, 1368-1371. | 1.3 | 95 |
| 6 | Lung Cancer in Never Smokers: A Call to Action: Fig. 1.. Clinical Cancer Research, 2009, 15, 5622-5625. | 3.2 | 33 |
| 7 | Telomeres and marrow failure. Hematology American Society of Hematology Education Program, 2009, 2009, 338-343. | 0.9 | 58 |
| 8 | Natural Variation within the Neuronal Nicotinic Acetylcholine Receptor Cluster on Human Chromosome 15q24: Influence on Heritable Autonomic Traits in Twin Pairs. Journal of Pharmacology and Experimental Therapeutics, 2009, 331, 419-428. | 1.3 | 8 |
| 9 | Apolipoprotein E/C1 Locus Variants Modify Renal Cell Carcinoma Risk. Cancer Research, 2009, 69, 8001-8008. | 0.4 | 31 |
| 10 | Common genetic variants on 8q24 contribute to susceptibility to bladder cancer in a Chinese population. Carcinogenesis, 2009, 30, 991-996. | 1.3 | 50 |
| 11 | Polymorphisms in Telomere Maintenance Genes and Risk of Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 2773-2781. | 1.1 | 54 |
| 12 | EUELC project: a multi-centre, multipurpose study to investigate early stage NSCLC, and to establish a biobank for ongoing collaboration. European Respiratory Journal, 2009, 34, 1477-1486. | 3.1 | 15 |
| 13 | Identification of <i>Las2</i> , a Major Modifier Gene Affecting the Pas1 Mouse Lung Tumor Susceptibility Locus. Cancer Research, 2009, 69, 6290-6298. | 0.4 | 6 |
| 14 | Deciphering the Impact of Common Genetic Variation on Lung Cancer Risk: A Genome-Wide Association Study. Cancer Research, 2009, 69, 6633-6641. | 0.4 | 206 |
| 15 | Invited Commentary: Genes, Environment, and Hybrid Vigor. American Journal of Epidemiology, 2009, 170, 703-707. | 1.6 | 5 |
| 16 | Human genetic variations: Beacons on the pathways to successful ageing. Mechanisms of Ageing and Development, 2009, 130, 553-563. | 2.2 | 23 |
| 17 | Association and interaction analysis of variants in <i>CHRNA5/CHRNA3/CHRNA4</i> gene cluster with nicotine dependence in African and European Americans. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2010, 153B, 745-756. | 1.1 | 53 |
| 18 | Body mass index and mortality from lung cancer in smokers and nonsmokers: A nationally representative prospective study of 220,000 men in China. International Journal of Cancer, 2009, 125, 2136-2143. | 2.3 | 54 |

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 19 | The search for genetic polymorphisms in the homocysteine/folate pathway that contribute to the etiology of human neural tube defects. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2009, 85, 285-294. | 1.6 | 74 |
| 20 | Lung cancer biology: a genetic and genomic perspective. <i>Clinical and Translational Oncology</i> , 2009, 11, 263-269. | 1.2 | 14 |
| 21 | Telomerase deficiency and cancer susceptibility syndromes. <i>Clinical and Translational Oncology</i> , 2009, 11, 711-714. | 1.2 | 11 |
| 22 | Telomeres and disease. <i>EMBO Journal</i> , 2009, 28, 2532-2540. | 3.5 | 99 |
| 23 | Variants in the CDKN2B and RTEL1 regions are associated with high-grade glioma susceptibility. <i>Nature Genetics</i> , 2009, 41, 905-908. | 9.4 | 456 |
| 24 | Genetic Susceptibility to Gastrointestinal Cancer: Minireview of the Genomewide Studies. <i>Annals of Surgical Oncology</i> , 2009, 16, 1783-1788. | 0.7 | 1 |
| 25 | A Genome-wide Association Study of Lung Cancer Identifies a Region of Chromosome 5p15 Associated with Risk for Adenocarcinoma. <i>American Journal of Human Genetics</i> , 2009, 85, 679-691. | 2.6 | 489 |
| 26 | Cancer diagnosis in first-degree relatives and non-small cell lung cancer risk: Results from a multi-centre case-control study in Europe. <i>European Journal of Cancer</i> , 2009, 45, 3047-3053. | 1.3 | 17 |
| 27 | Telomere Diseases. <i>New England Journal of Medicine</i> , 2009, 361, 2353-2365. | 13.9 | 723 |
| 28 | Stem cells and lung cancer: future therapeutic targets?. <i>Expert Opinion on Biological Therapy</i> , 2009, 9, 1127-1141. | 1.4 | 16 |
| 29 | Acetylcholine Receptor Pathway and Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2009, 4, 943-946. | 0.5 | 37 |
| 30 | Assessment of cumulative evidence for the association between glutathione S-transferase polymorphisms and lung cancer: application of the Venice interim guidelines. <i>Pharmacogenetics and Genomics</i> , 2010, 20, 586-597. | 0.7 | 33 |
| 31 | Deregulation of the telomerase reverse transcriptase (TERT) gene by chromosomal translocations in B-cell malignancies. <i>Blood</i> , 2010, 116, 1317-1320. | 0.6 | 44 |
| 32 | Recent progress in dyskeratosis congenita. <i>International Journal of Hematology</i> , 2010, 92, 419-424. | 0.7 | 34 |
| 33 | Family history and lung cancer risk: international multicentre case-control study in Eastern and Central Europe and meta-analyses. <i>Cancer Causes and Control</i> , 2010, 21, 1091-1104. | 0.8 | 81 |
| 34 | The association of telomere length and genetic variation in telomere biology genes. <i>Human Mutation</i> , 2010, 31, 1050-1058. | 1.1 | 93 |
| 35 | Role of shelterin in cancer and aging. <i>Aging Cell</i> , 2010, 9, 653-666. | 3.0 | 127 |
| 36 | Genetic variants cis-regulating Xrn2 expression contribute to the risk of spontaneous lung tumor. <i>Oncogene</i> , 2010, 29, 1041-1049. | 2.6 | 26 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 37 | Homeobox gene IRX1 is a tumor suppressor gene in gastric carcinoma. <i>Oncogene</i> , 2010, 29, 3908-3920. | 2.6 | 78 |
| 38 | 2009 Nobel Prize in Physiology or Medicine: telomeres and telomerase. <i>Oncogene</i> , 2010, 29, 1561-1565. | 2.6 | 38 |
| 39 | A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. <i>Nature Genetics</i> , 2010, 42, 224-228. | 9.4 | 539 |
| 40 | Variants near DMRT1, TERT and ATF7IP are associated with testicular germ cell cancer. <i>Nature Genetics</i> , 2010, 42, 604-607. | 9.4 | 320 |
| 41 | Variation in TP63 is associated with lung adenocarcinoma susceptibility in Japanese and Korean populations. <i>Nature Genetics</i> , 2010, 42, 893-896. | 9.4 | 165 |
| 42 | Architecture of inherited susceptibility to common cancer. <i>Nature Reviews Cancer</i> , 2010, 10, 353-361. | 12.8 | 183 |
| 43 | Tobacco, Genetic Susceptibility and Lung cancer. <i>Tobacco Use Insights</i> , 2010, 3, TUI.S2819. | 0.7 | 2 |
| 44 | Genetic Variations in Telomere Maintenance, with Implications on Tissue Renewal Capacity and Chronic Disease Pathologies. <i>Current Pharmacogenomics and Personalized Medicine</i> , 2010, 8, 7-24. | 0.2 | 17 |
| 45 | Replication of Lung Cancer Susceptibility Loci at Chromosomes 15q25, 5p15, and 6p21: A Pooled Analysis From the International Lung Cancer Consortium. <i>Journal of the National Cancer Institute</i> , 2010, 102, 959-971. | 3.0 | 174 |
| 46 | Search for Cancer Risk Factors with Microarray-Based Genome-Wide Association Studies. <i>Technology in Cancer Research and Treatment</i> , 2010, 9, 107-121. | 0.8 | 7 |
| 47 | Biomarkers in cancer epidemiology: an integrative approach. <i>Carcinogenesis</i> , 2010, 31, 121-126. | 1.3 | 39 |
| 48 | A Second Genetic Variant on Chromosome 15q24-25.1 Associates with Lung Cancer. <i>Cancer Research</i> , 2010, 70, 3128-3135. | 0.4 | 5 |
| 49 | A Rigorous and Comprehensive Validation: Common Genetic Variations and Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 240-244. | 1.1 | 37 |
| 50 | A Compendium of Genome-Wide Associations for Cancer: Critical Synopsis and Reappraisal. <i>Journal of the National Cancer Institute</i> , 2010, 102, 846-858. | 3.0 | 68 |
| 51 | Genetic Basis for Susceptibility to Lung Cancer. <i>Advances in Cancer Research</i> , 2010, 109, 51-72. | 1.9 | 65 |
| 52 | Cumulative Effect of Multiple Loci on Genetic Susceptibility to Familial Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 517-524. | 1.1 | 24 |
| 53 | International Lung Cancer Consortium: Coordinated association study of 10 potential lung cancer susceptibility variants. <i>Carcinogenesis</i> , 2010, 31, 625-633. | 1.3 | 56 |
| 54 | A genome-wide association study reveals susceptibility variants for non-small cell lung cancer in the Korean population. <i>Human Molecular Genetics</i> , 2010, 19, 4948-4954. | 1.4 | 78 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 55 | Variation at the <i>TERT</i> locus and predisposition for cancer. <i>Expert Reviews in Molecular Medicine</i> , 2010, 12, e16. | 1.6 | 101 |
| 56 | Role of 5p15.33 (<i>TERT-CLPTM1L</i>), 6p21.33 and 15q25.1 (<i>CHRNA5-CHRNA3</i>) variation and lung cancer risk in never-smokers. <i>Carcinogenesis</i> , 2010, 31, 234-238. | 1.3 | 97 |
| 57 | Individuals susceptible to lung adenocarcinoma defined by combined HLA-DQA1 and <i>TERT</i> genotypes. <i>Carcinogenesis</i> , 2010, 31, 834-841. | 1.3 | 44 |
| 58 | Genome-wide association studies in cancer—current and future directions. <i>Carcinogenesis</i> , 2010, 31, 111-120. | 1.3 | 100 |
| 59 | Genetic variations in <i>TERT-CLPTM1L</i> genes and risk of squamous cell carcinoma of the head and neck. <i>Carcinogenesis</i> , 2010, 31, 1977-1981. | 1.3 | 41 |
| 60 | The evolving discipline of molecular epidemiology of cancer. <i>Carcinogenesis</i> , 2010, 31, 127-134. | 1.3 | 42 |
| 61 | Evaluation of Candidate Stromal Epithelial Cross-Talk Genes Identifies Association between Risk of Serous Ovarian Cancer and <i>TERT</i> , a Cancer Susceptibility “Hot-Spot”. <i>PLoS Genetics</i> , 2010, 6, e1001016. | 1.5 | 48 |
| 62 | Genome-wide association study for colorectal cancer identifies risk polymorphisms in German familial cases and implicates MAPK signalling pathways in disease susceptibility. <i>Carcinogenesis</i> , 2010, 31, 1612-1619. | 1.3 | 57 |
| 63 | The 5p15.33 Locus Is Associated with Risk of Lung Adenocarcinoma in Never-Smoking Females in Asia. <i>PLoS Genetics</i> , 2010, 6, e1001051. | 1.5 | 168 |
| 64 | Association between a 15q25 gene variant, smoking quantity and tobacco-related cancers among 17 000 individuals. <i>International Journal of Epidemiology</i> , 2010, 39, 563-577. | 0.9 | 125 |
| 65 | The pursuit of genome-wide association studies: where are we now?. <i>Journal of Human Genetics</i> , 2010, 55, 195-206. | 1.1 | 191 |
| 66 | Genome-wide Association Studies of Cancer Predisposition. <i>Hematology/Oncology Clinics of North America</i> , 2010, 24, 973-996. | 0.9 | 34 |
| 67 | Multiple Genetic Variants in Telomere Pathway Genes and Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 219-228. | 1.1 | 47 |
| 68 | Genome-Wide Association Studies of Cancer. <i>Journal of Clinical Oncology</i> , 2010, 28, 4255-4267. | 0.8 | 159 |
| 69 | Genetic variants and risk of lung cancer in never smokers: a genome-wide association study. <i>Lancet Oncology</i> , The, 2010, 11, 321-330. | 5.1 | 218 |
| 71 | ABO blood group and other genetic variants associated with pancreatic cancer. <i>Genome Medicine</i> , 2010, 2, 39. | 3.6 | 4 |
| 72 | Recent progress in genetic variants associated with cancer and their implications in diagnostics development. <i>Expert Review of Molecular Diagnostics</i> , 2010, 10, 699-703. | 1.5 | 21 |
| 73 | The vitamin D axis in the lung: a key role for vitamin D-binding protein. <i>Thorax</i> , 2010, 65, 456-462. | 2.7 | 166 |

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 74 | A genome-wide association study identifies two new lung cancer susceptibility loci at 13q12.12 and 22q12.2 in Han Chinese. <i>Nature Genetics</i> , 2011, 43, 792-796. | 9.4 | 340 |
| 75 | A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , 2011, 43, 1210-1214. | 9.4 | 279 |
| 76 | Lung Cancer Risk Prediction: Prostate, Lung, Colorectal and Ovarian Cancer Screening Trial Models and Validation. <i>Journal of the National Cancer Institute</i> , 2011, 103, 1058-1068. | 3.0 | 259 |
| 77 | Genetics of lung-cancer susceptibility. <i>Lancet Oncology</i> , The, 2011, 12, 399-408. | 5.1 | 191 |
| 78 | Chronic Obstructive Pulmonary Disease: A Complex Comorbidity of Lung Cancer. <i>Journal of Comorbidity</i> , 2011, 1, 45-50. | 3.9 | 5 |
| 79 | Single nucleotide polymorphisms as susceptibility, prognostic, and therapeutic markers of nonsmall cell lung cancer. <i>Lung Cancer: Targets and Therapy</i> , 2011, 3, 1. | 1.3 | 20 |
| 80 | A Genome-Wide Association Study of Upper Aerodigestive Tract Cancers Conducted within the INHANCE Consortium. <i>PLoS Genetics</i> , 2011, 7, e1001333. | 1.5 | 158 |
| 81 | Most Lung and Colon Cancer Susceptibility Genes Are Pair-Wise Linked in Mice, Humans and Rats. <i>PLoS ONE</i> , 2011, 6, e14727. | 1.1 | 18 |
| 82 | Functional Polymorphisms in the TERT Promoter Are Associated with Risk of Serous Epithelial Ovarian and Breast Cancers. <i>PLoS ONE</i> , 2011, 6, e24987. | 1.1 | 48 |
| 83 | Smoking Related Cancers and Loci at Chromosomes 15q25, 5p15, 6p22.1 and 6p21.33 in the Polish Population. <i>PLoS ONE</i> , 2011, 6, e25057. | 1.1 | 37 |
| 84 | Screening for Lung Cancer: For Patients at Increased Risk for Lung Cancer, It Works. <i>Annals of Internal Medicine</i> , 2011, 155, 540. | 2.0 | 23 |
| 85 | Contribution of the TP53, OGG1, CHRNA3, and HLA-DQA1 Genes to the Risk for Lung Squamous Cell Carcinoma. <i>Journal of Thoracic Oncology</i> , 2011, 6, 813-817. | 0.5 | 55 |
| 86 | Absence of association of a single-nucleotide polymorphism in the TERT-CLPTM1L locus with age-related phenotypes in a large multicohort study: the HALCyon programme. <i>Aging Cell</i> , 2011, 10, 520-532. | 3.0 | 8 |
| 87 | The telomerase activator TA65 elongates short telomeres and increases health span of adult/old mice without increasing cancer incidence. <i>Aging Cell</i> , 2011, 10, 604-621. | 3.0 | 259 |
| 88 | Analysis of Potential Genomic Confounding in Genetic Association Studies and an Online Genomic Confounding Browser (GCB). <i>Annals of Human Genetics</i> , 2011, 75, 723-731. | 0.3 | 2 |
| 89 | Telomeric and extra-telomeric roles for telomerase and the telomere-binding proteins. <i>Nature Reviews Cancer</i> , 2011, 11, 161-176. | 12.8 | 443 |
| 90 | The potential utility of telomere-related markers for cancer diagnosis. <i>Journal of Cellular and Molecular Medicine</i> , 2011, 15, 1227-1238. | 1.6 | 43 |
| 91 | A stratified genetic risk assessment for testicular cancer. <i>Journal of Developmental and Physical Disabilities</i> , 2011, 34, e98-102. | 3.6 | 25 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 92 | Genome-wide association studies provide new insights into the genetic basis of testicular germ cell tumour. <i>Journal of Developmental and Physical Disabilities</i> , 2011, 34, e86-96; discussion e96-7. | 3.6 | 56 |
| 93 | An analysis of single nucleotide polymorphisms of 125 DNA repair genes in the Texas genome-wide association study of lung cancer with a replication for the XRCC4 SNPs. <i>DNA Repair</i> , 2011, 10, 398-407. | 1.3 | 26 |
| 95 | Genes as instruments for studying risk behavior effects: an application to maternal smoking and orofacial clefts. <i>Health Services and Outcomes Research Methodology</i> , 2011, 11, 54-78. | 0.8 | 36 |
| 96 | Cancer susceptibility variants and the risk of adult glioma in a US case-control study. <i>Journal of Neuro-Oncology</i> , 2011, 104, 535-542. | 1.4 | 77 |
| 97 | Current status of genome-wide association studies in cancer. <i>Human Genetics</i> , 2011, 130, 59-78. | 1.8 | 160 |
| 98 | Significance testing in ridge regression for genetic data. <i>BMC Bioinformatics</i> , 2011, 12, 372. | 1.2 | 84 |
| 99 | Constitutional telomerase mutations are genetic risk factors for cirrhosis. <i>Hepatology</i> , 2011, 53, 1600-1607. | 3.6 | 145 |
| 100 | Lung Cancer in Never Smokers. <i>Seminars in Respiratory and Critical Care Medicine</i> , 2011, 32, 010-021. | 0.8 | 31 |
| 101 | Common pathogenic mechanisms and pathways in the development of COPD and lung cancer. <i>Expert Opinion on Therapeutic Targets</i> , 2011, 15, 439-456. | 1.5 | 77 |
| 102 | Genetic variations on chromosomes 5p15 and 15q25 and bladder cancer risk: findings from the Los Angeles-Shanghai bladder case-control study. <i>Carcinogenesis</i> , 2011, 32, 197-202. | 1.3 | 52 |
| 103 | Genome-Wide Association Study of Survival in Non-Small Cell Lung Cancer Patients Receiving Platinum-Based Chemotherapy. <i>Journal of the National Cancer Institute</i> , 2011, 103, 817-825. | 3.0 | 81 |
| 104 | Variation in <i>TP63</i> is Associated with Lung Adenocarcinoma in the UK Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1453-1462. | 1.1 | 25 |
| 105 | Genetic Variation in an miRNA-1827 Binding Site in <i>MYCL1</i> Alters Susceptibility to Small-Cell Lung Cancer. <i>Cancer Research</i> , 2011, 71, 5175-5181. | 0.4 | 73 |
| 106 | Genome-Wide Significant Association Between a Sequence Variant at 15q15.2 and Lung Cancer Risk. <i>Cancer Research</i> , 2011, 71, 1356-1361. | 0.4 | 26 |
| 107 | Novel genetic variants in the chromosome 5p15.33 region associate with lung cancer risk. <i>Carcinogenesis</i> , 2011, 32, 1493-1499. | 1.3 | 59 |
| 108 | Association of the CHRNA5-A3-B4 Gene Cluster With Heaviness of Smoking: A Meta-Analysis. <i>Nicotine and Tobacco Research</i> , 2011, 13, 1167-1175. | 1.4 | 106 |
| 109 | Single-nucleotide polymorphisms (5p15.33, 15q25.1, 6p22.1, 6q27 and 7p15.3) and lung cancer survival in the European Prospective Investigation into Cancer and Nutrition (EPIC). <i>Mutagenesis</i> , 2011, 26, 657-666. | 1.0 | 20 |
| 110 | Genome-wide association studies for detecting cancer susceptibility. <i>British Medical Bulletin</i> , 2011, 97, 27-46. | 2.7 | 22 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 111 | Association of a novel functional promoter variant (rs2075533 C>T) in the apoptosis gene TNFSF 8 with risk of lung cancer—a finding from Texas lung cancer genome-wide association study. <i>Carcinogenesis</i> , 2011, 32, 507-515. | 1.3 | 15 |
| 112 | A common genetic variant of 5p15.33 is associated with risk for prostate cancer in the Chinese population. <i>Genetics and Molecular Research</i> , 2012, 11, 1349-1356. | 0.3 | 2 |
| 113 | Research Opportunities for Cancer Associated with Indoor Air Pollution from Solid-Fuel Combustion. <i>Environmental Health Perspectives</i> , 2012, 120, 1495-1498. | 2.8 | 32 |
| 114 | Worldwide genetic structure in 37 genes important in telomere biology. <i>Heredity</i> , 2012, 108, 124-133. | 1.2 | 11 |
| 115 | Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012, 21, 4980-4995. | 1.4 | 196 |
| 116 | Association between a Genome-Wide Association Study-Identified Locus and the Risk of Lung Cancer in Japanese Population. <i>Journal of Thoracic Oncology</i> , 2012, 7, 790-798. | 0.5 | 37 |
| 117 | Defining a Gene Promoter Methylation Signature in Sputum for Lung Cancer Risk Assessment. <i>Clinical Cancer Research</i> , 2012, 18, 3387-3395. | 3.2 | 96 |
| 118 | Class A Scavenger Receptor Deficiency Exacerbates Lung Tumorigenesis by Cultivating a Procarcinogenic Microenvironment in Humans and Mice. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2012, 186, 763-772. | 2.5 | 23 |
| 119 | Genetic Variants Associated with the Risk of Chronic Obstructive Pulmonary Disease with and without Lung Cancer. <i>Cancer Prevention Research</i> , 2012, 5, 365-373. | 0.7 | 26 |
| 120 | Inherited Variation at Chromosome 12p13.33, Including <i>RAD52</i> , Influences the Risk of Squamous Cell Lung Carcinoma. <i>Cancer Discovery</i> , 2012, 2, 131-139. | 7.7 | 54 |
| 121 | Association analyses identify multiple new lung cancer susceptibility loci and their interactions with smoking in the Chinese population. <i>Nature Genetics</i> , 2012, 44, 895-899. | 9.4 | 129 |
| 122 | The TERT variant rs2736100 is associated with colorectal cancer risk. <i>British Journal of Cancer</i> , 2012, 107, 1001-1008. | 2.9 | 50 |
| 123 | A Risk Model for Lung Cancer Incidence. <i>Cancer Prevention Research</i> , 2012, 5, 834-846. | 0.7 | 93 |
| 124 | Interaction of Occupational and Personal Risk Factors in Workforce Health and Safety. <i>American Journal of Public Health</i> , 2012, 102, 434-448. | 1.5 | 142 |
| 125 | Chapter 6: Lung Cancer in Never Smokers: Epidemiology and Risk Prediction Models. <i>Risk Analysis</i> , 2012, 32, S69-84. | 1.5 | 73 |
| 126 | Ectopic Expression of Telomerase Safely Increases Health Span and Life Span. <i>Rejuvenation Research</i> , 2012, 15, 435-438. | 0.9 | 12 |
| 127 | Review: a meta-analysis of GWAS and age-associated diseases. <i>Aging Cell</i> , 2012, 11, 727-731. | 3.0 | 168 |
| 128 | Genetic Variants at 6p21.1 and 7p15.3 Are Associated with Risk of Multiple Cancers in Han Chinese. <i>American Journal of Human Genetics</i> , 2012, 91, 928-934. | 2.6 | 76 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 129 | Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335. | 9.4 | 286 |
| 130 | Genetic variation in innate immunity and inflammation pathways associated with lung cancer risk. <i>Cancer</i> , 2012, 118, 5630-5636. | 2.0 | 30 |
| 131 | Tobacco consumption and genetic susceptibility to nasopharyngeal carcinoma (NPC) in Thailand. <i>Cancer Causes and Control</i> , 2012, 23, 1995-2002. | 0.8 | 47 |
| 132 | Lung cancer in never smokers – A review. <i>European Journal of Cancer</i> , 2012, 48, 1299-1311. | 1.3 | 694 |
| 133 | Telomerase Reverse Transcriptase Locus Polymorphisms and Cancer Risk: A Field Synopsis and Meta-Analysis. <i>Journal of the National Cancer Institute</i> , 2012, 104, 840-854. | 3.0 | 119 |
| 134 | Short telomeres result in chromosomal instability in hematopoietic cells and precede malignant evolution in human aplastic anemia. <i>Leukemia</i> , 2012, 26, 700-707. | 3.3 | 95 |
| 135 | Molecular changes in smoking-related lung cancer. <i>Expert Review of Molecular Diagnostics</i> , 2012, 12, 93-106. | 1.5 | 22 |
| 136 | A genome-wide association study identifies two new susceptibility loci for lung adenocarcinoma in the Japanese population. <i>Nature Genetics</i> , 2012, 44, 900-903. | 9.4 | 166 |
| 137 | Prediction of lung cancer risk in a Chinese population using a multifactorial genetic model. <i>BMC Medical Genetics</i> , 2012, 13, 118. | 2.1 | 47 |
| 138 | Comparison of Pathway Analysis Approaches Using Lung Cancer GWAS Data Sets. <i>PLoS ONE</i> , 2012, 7, e31816. | 1.1 | 38 |
| 139 | Integrative Genomic Analysis Reveals Extended Germline Homozygosity with Lung Cancer Risk in the PLCO Cohort. <i>PLoS ONE</i> , 2012, 7, e31975. | 1.1 | 13 |
| 140 | Functional Characterization of CLPTM1L as a Lung Cancer Risk Candidate Gene in the 5p15.33 Locus. <i>PLoS ONE</i> , 2012, 7, e36116. | 1.1 | 89 |
| 141 | TERT-CLPTM1L Polymorphism rs401681 Contributes to Cancers Risk: Evidence from a Meta-Analysis Based on 29 Publications. <i>PLoS ONE</i> , 2012, 7, e50650. | 1.1 | 26 |
| 142 | Comparative Analysis of Methods for Identifying Recurrent Copy Number Alterations in Cancer. <i>PLoS ONE</i> , 2012, 7, e52516. | 1.1 | 11 |
| 143 | Telomere and telomerase in stem cells: relevance in ageing and disease. <i>Frontiers in Bioscience - Scholar</i> , 2012, S4, 16. | 0.8 | 12 |
| 144 | The Investigation of Gene Regulation and Variation in Human Cancers and Other Diseases. , 2012, , . | | 0 |
| 145 | Genetic Polymorphisms Underlying Lung Cancer Susceptibility and Therapeutic Response. <i>Genes and Environment</i> , 2012, 34, 94-100. | 0.9 | 1 |
| 146 | Multiple variants of TERT and CLPTM1L constitute risk factors for lung adenocarcinoma. <i>Genetics and Molecular Research</i> , 2012, 11, 370-378. | 0.3 | 46 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 147 | Association between Lung Cancer Susceptibility Variants Identified by Genome-Wide Association Studies and the Survival of Non-Small Cell Lung Cancer. <i>Journal of Lung Cancer</i> , 2012, 11, 66. | 0.2 | 0 |
| 148 | Telomerase Activity and the Risk of Lung Cancer. <i>Journal of Korean Medical Science</i> , 2012, 27, 141. | 1.1 | 9 |
| 149 | Common genetic variants in <i>TERT</i> contribute to risk of cervical cancer in a Chinese population. <i>Molecular Carcinogenesis</i> , 2012, 51, E118-22. | 1.3 | 28 |
| 150 | Association of genetic variants of human telomerase with colorectal polyps and colorectal cancer risk. <i>Molecular Carcinogenesis</i> , 2012, 51, E176-82. | 1.3 | 34 |
| 151 | Fine-mapping of a region of chromosome 5p15.3 (<i>TERT</i> and <i>CLPTM1L</i>) suggests a novel locus in <i>TERT</i> and a <i>CLPTM1L</i> haplotype are associated with glioma susceptibility in a Chinese population. <i>International Journal of Cancer</i> , 2012, 131, 1569-1576. | 2.3 | 21 |
| 152 | Telomerase gene therapy in adult and old mice delays aging and increases longevity without increasing cancer. <i>EMBO Molecular Medicine</i> , 2012, 4, 691-704. | 3.3 | 403 |
| 153 | Telomere Length and Genetic Variation in Telomere Maintenance Genes in Relation to Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 504-512. | 1.1 | 39 |
| 154 | Genetic variant in TP63 on locus 3q28 is associated with risk of lung adenocarcinoma among never-smoking females in Asia. <i>Human Genetics</i> , 2012, 131, 1197-1203. | 1.8 | 39 |
| 155 | Genetics of leukocyte telomere length and its role in atherosclerosis. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2012, 730, 68-74. | 0.4 | 122 |
| 156 | Replication of results of genome-wide association studies on lung cancer susceptibility loci in a Korean population. <i>Respirology</i> , 2012, 17, 699-706. | 1.3 | 52 |
| 157 | The TERT rs2736100 Polymorphism and Cancer Risk: A Meta-analysis Based on 25 Case-Control Studies. <i>BMC Cancer</i> , 2012, 12, 7. | 1.1 | 27 |
| 158 | Genetic susceptibility to pancreatic cancer. <i>Molecular Carcinogenesis</i> , 2012, 51, 14-24. | 1.3 | 192 |
| 159 | Finding biomarkers for non-small cell lung cancer diagnosis and prognosis. <i>Frontiers in Biology</i> , 2012, 7, 14-23. | 0.7 | 3 |
| 160 | Lack of association between common single nucleotide polymorphisms in the TERT-CLPTM1L locus and breast cancer in women of African ancestry. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 341-345. | 1.1 | 12 |
| 161 | Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , 2012, 131, 217-234. | 1.8 | 183 |
| 162 | The 18p11.22 locus is associated with never smoker non-small cell lung cancer susceptibility in Korean populations. <i>Human Genetics</i> , 2012, 131, 365-372. | 1.8 | 45 |
| 163 | Allelotypes of lung adenocarcinomas featuring ALK fusion demonstrate fewer onco- and suppressor gene changes. <i>BMC Cancer</i> , 2013, 13, 8. | 1.1 | 4 |
| 164 | Hierarchical modeling identifies novel lung cancer susceptibility variants in inflammation pathways among 10,140 cases and 11,012 controls. <i>Human Genetics</i> , 2013, 132, 579-589. | 1.8 | 29 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 165 | Telomerase at the intersection of cancer and aging. <i>Trends in Genetics</i> , 2013, 29, 513-520. | 2.9 | 186 |
| 166 | Rewiring of human lung cell lineage and mitotic networks in lung adenocarcinomas. <i>Nature Communications</i> , 2013, 4, 1701. | 5.8 | 42 |
| 167 | Lung cancer in never smokers: Disease characteristics and risk factors. <i>Critical Reviews in Oncology/Hematology</i> , 2013, 88, 494-503. | 2.0 | 130 |
| 168 | Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384. | 9.4 | 493 |
| 169 | Genetic variant in the telomerase gene modifies cancer risk in Lynch syndrome. <i>European Journal of Human Genetics</i> , 2013, 21, 511-516. | 1.4 | 20 |
| 170 | Validity of the diagnosis of pre-eclampsia in the Medical Birth Registry of Norway. <i>Acta Obstetrica Et Gynecologica Scandinavica</i> , 2013, 92, 943-950. | 1.3 | 82 |
| 171 | TERT polymorphisms modify the risk of acute lymphoblastic leukemia in Chinese children. <i>Carcinogenesis</i> , 2013, 34, 228-235. | 1.3 | 41 |
| 172 | Telomere Dynamics in Mice and Humans. <i>Seminars in Hematology</i> , 2013, 50, 165-174. | 1.8 | 158 |
| 173 | Genetic polymorphisms of TERT and CLPTM1L and risk of lung cancer—A case-control study in a Chinese population. <i>Lung Cancer</i> , 2013, 80, 131-137. | 0.9 | 33 |
| 174 | Telomeres in lung disease. <i>Translational Research</i> , 2013, 162, 343-352. | 2.2 | 32 |
| 175 | Spatial and temporal distributions of lung cancer histopathology in the state of Maine. <i>Lung Cancer</i> , 2013, 82, 55-62. | 0.9 | 13 |
| 176 | Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876. | 9.4 | 179 |
| 177 | Exposing a deadly alliance: Novel insights into the biological links between COPD and lung cancer. <i>Pulmonary Pharmacology and Therapeutics</i> , 2013, 26, 544-554. | 1.1 | 45 |
| 178 | ICD-9 tobacco use codes are effective identifiers of smoking status. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2013, 20, 652-658. | 2.2 | 127 |
| 179 | RGS17: an emerging therapeutic target for lung and prostate cancers. <i>Future Medicinal Chemistry</i> , 2013, 5, 995-1007. | 1.1 | 31 |
| 180 | Epigenetic screen identifies genotype-specific promoter DNA methylation and oncogenic potential of CHRN4. <i>Oncogene</i> , 2013, 32, 3329-3338. | 2.6 | 40 |
| 181 | Genome-Wide Association Study Identifies a Novel Susceptibility Locus at 12q23.1 for Lung Squamous Cell Carcinoma in Han Chinese. <i>PLoS Genetics</i> , 2013, 9, e1003190. | 1.5 | 41 |
| 182 | Regulatory Nexus of Synthesis and Degradation Deciphers Cellular Nrf2 Expression Levels. <i>Molecular and Cellular Biology</i> , 2013, 33, 2402-2412. | 1.1 | 101 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|------|-----------|
| 183 | A Resequencing Analysis of Genomic Loci on Chromosomes 1q32.1, 5p15.33, and 13q22.1 Associated With Pancreatic Cancer Risk. <i>Pancreas</i> , 2013, 42, 209-215. | 0.5 | 5 |
| 184 | Genome-Wide Association Study of Genetic Predictors of Overall Survival for Non-Small Cell Lung Cancer in Never Smokers. <i>Cancer Research</i> , 2013, 73, 4028-4038. | 0.4 | 53 |
| 185 | Fine-mapping of the 5p15.33, 6p22.1-p21.31, and 15q25.1 Regions Identifies Functional and Histology-Specific Lung Cancer Susceptibility Loci in African-Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 251-260. | 1.1 | 36 |
| 186 | Inherited predisposition to multiple myeloma. <i>Therapeutic Advances in Hematology</i> , 2013, 4, 291-297. | 1.1 | 13 |
| 187 | EGFR polymorphisms, hormone replacement therapy and lung adenocarcinoma risk: analysis from a genome-wide association study in never-smoking women. <i>Carcinogenesis</i> , 2013, 34, 612-619. | 1.3 | 15 |
| 188 | Genetic susceptibility to lung cancer—light at the end of the tunnel?. <i>Carcinogenesis</i> , 2013, 34, 487-502. | 1.3 | 82 |
| 189 | Gene expression changes with age in skin, adipose tissue, blood and brain. <i>Genome Biology</i> , 2013, 14, R75. | 13.9 | 263 |
| 190 | Significant association of 5p15.33 (<i>TERT</i> / <i>CLPTM1L</i> genes) with lung cancer in Chinese Han population. <i>Experimental Lung Research</i> , 2013, 39, 91-98. | 0.5 | 23 |
| 191 | Genetic variants at 5p15 are associated with risk and early onset of gastric cancer in Chinese populations. <i>Carcinogenesis</i> , 2013, 34, 2539-2542. | 1.3 | 13 |
| 192 | Genetic variations in <i>TERT</i> / <i>CLPTM1L</i> locus are associated with risk of lung cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2013, 52, 118-126. | 1.3 | 44 |
| 193 | Turning of COGS moves forward findings for hormonally mediated cancers. <i>Nature Genetics</i> , 2013, 45, 345-348. | 9.4 | 69 |
| 194 | Ethnic differences in non-small-cell lung cancer treatment: the Asian perspective. <i>Lung Cancer Management</i> , 2013, 2, 309-315. | 1.5 | 2 |
| 195 | Genome-wide association studies of cancer predisposition. , 2013, , 10-20. | | 1 |
| 196 | Genome-Wide Association Study of Lung Cancer in Korean Non-Smoking Women. <i>Journal of Korean Medical Science</i> , 2013, 28, 840. | 1.1 | 17 |
| 197 | Short Communication The genetic variant rs401681C/T is associated with the risk of non-small cell lung cancer in a Chinese mainland population. <i>Genetics and Molecular Research</i> , 2013, 12, 67-73. | 0.3 | 15 |
| 198 | Getting More Out of Biomedical Documents with GATE's Full Lifecycle Open Source Text Analytics. <i>PLoS Computational Biology</i> , 2013, 9, e1002854. | 1.5 | 261 |
| 199 | Pathway Analysis for Genome-Wide Association Study of Lung Cancer in Han Chinese Population. <i>PLoS ONE</i> , 2013, 8, e57763. | 1.1 | 9 |
| 200 | Replication Study in Chinese Population and Meta-Analysis Supports Association of the 5p15.33 Locus with Lung Cancer. <i>PLoS ONE</i> , 2013, 8, e62485. | 1.1 | 17 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 201 | Association between the Telomerase Reverse Transcriptase (TERT) rs2736098 Polymorphism and Cancer Risk: Evidence from a Case-Control Study of Non-Small-Cell Lung Cancer and a Meta-Analysis. PLoS ONE, 2013, 8, e76372. | 1.1 | 32 |
| 202 | Genetic Variations in TERT-CLPTM1L Genes and Risk of Lung Cancer in Chinese Women Nonsmokers. PLoS ONE, 2013, 8, e64988. | 1.1 | 23 |
| 203 | Environmental Exposures and Cancer. , 2013, , 647-666. | | 0 |
| 204 | Epidemiological and clinical aspects of lung cancer. , 0, , 945-1003. | | 1 |
| 205 | Decreased risk of developing lung cancer in subjects carrying the CLPTM1L rs401681 (G>A) polymorphism: evidence from a meta-analysis. Genetics and Molecular Research, 2014, 13, 1373-1382. | 0.3 | 5 |
| 206 | Quantitative Assessment of the Influence of TP63 Gene Polymorphisms and Lung Cancer Risk: Evidence Based on 93,751 Subjects. PLoS ONE, 2014, 9, e87004. | 1.1 | 11 |
| 207 | The GG Genotype of Telomerase Reverse Transcriptase at Genetic Locus rs2736100 Is Associated with Human Atherosclerosis Risk in the Han Chinese Population. PLoS ONE, 2014, 9, e85719. | 1.1 | 7 |
| 208 | Role of Nicotine Dependence on the Relationship between Variants in the Nicotinic Receptor Genes and Risk of Lung Adenocarcinoma. PLoS ONE, 2014, 9, e107268. | 1.1 | 20 |
| 209 | Association of GWAS-Identified Lung Cancer Susceptibility Loci with Survival Length in Patients with Small-Cell Lung Cancer Treated with Platinum-Based Chemotherapy. PLoS ONE, 2014, 9, e113574. | 1.1 | 8 |
| 210 | Overexpression of MEOX2 and TWIST1 Is Associated with H3K27me3 Levels and Determines Lung Cancer Chemoresistance and Prognosis. PLoS ONE, 2014, 9, e114104. | 1.1 | 35 |
| 211 | Susceptibility loci for lung cancer are associated with mRNA levels of nearby genes in the lung. Carcinogenesis, 2014, 35, 2653-2659. | 1.3 | 18 |
| 212 | A genome-wide gene-gene interaction analysis identifies an epistatic gene pair for lung cancer susceptibility in Han Chinese. Carcinogenesis, 2014, 35, 572-577. | 1.3 | 29 |
| 213 | Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. Gut, 2014, 63, 800-807. | 6.1 | 35 |
| 214 | Correlation of CLPTM1L polymorphisms with lung cancer susceptibility and response to cisplatin-based chemotherapy in a Chinese Han population. Tumor Biology, 2014, 35, 12075-12082. | 0.8 | 12 |
| 215 | Combined Analysis with Copy Number Variation Identifies Risk Loci in Lung Cancer. BioMed Research International, 2014, 2014, 1-9. | 0.9 | 4 |
| 216 | Association of Cancer Susceptibility Variants with Risk of Multiple Primary Cancers: The Population Architecture using Genomics and Epidemiology Study. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2568-2578. | 1.1 | 23 |
| 217 | Naturally Occurring Cancers in Dogs: Insights for Translational Genetics and Medicine. ILAR Journal, 2014, 55, 16-45. | 1.8 | 62 |
| 218 | Association Between Interleukin-8 rs251A/T Polymorphism and Risk of Lung Cancer: A Meta-Analysis. Cancer Investigation, 2014, 32, 518-525. | 0.6 | 15 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 219 | Common germline variation at the TERT locus contributes to familial clustering of myeloproliferative neoplasms. <i>American Journal of Hematology</i> , 2014, 89, 1107-1110. | 2.0 | 47 |
| 220 | Genetic variants in DNA repair pathways and risk of upper aerodigestive tract cancers: combined analysis of data from two genome-wide association studies in European populations. <i>Carcinogenesis</i> , 2014, 35, 1523-1527. | 1.3 | 10 |
| 221 | Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633. | 1.4 | 90 |
| 222 | A <i>DRD1</i> Polymorphism Predisposes to Lung Cancer among Those Exposed to Secondhand Smoke during Childhood. <i>Cancer Prevention Research</i> , 2014, 7, 1210-1218. | 0.7 | 25 |
| 223 | Genetic Association of <i>BSF2</i> Polymorphism and Susceptibility to Lung Cancer. <i>Cell Biochemistry and Biophysics</i> , 2014, 70, 1887-1891. | 0.9 | 2 |
| 224 | <i>CRR9p</i> polymorphism as a protective factor for lung cancer. <i>Tumor Biology</i> , 2014, 35, 9557-9562. | 0.8 | 0 |
| 225 | <i>CRR9/CLPTM1L</i> Regulates Cell Survival Signaling and Is Required for Ras Transformation and Lung Tumorigenesis. <i>Cancer Research</i> , 2014, 74, 1116-1127. | 0.4 | 63 |
| 226 | A Functional Polymorphism in <i>CSF1R</i> Gene Is a Novel Susceptibility Marker for Lung Cancer among Never-Smoking Females. <i>Journal of Thoracic Oncology</i> , 2014, 9, 1647-1655. | 0.5 | 19 |
| 227 | Polymorphisms in the TERT gene are associated with lung cancer risk in the Chinese Han population. <i>European Journal of Cancer Prevention</i> , 2014, 23, 497-501. | 0.6 | 38 |
| 228 | Genetic Susceptibility and Risk of Pancreatic Cancer. , 2014, , 169-194. | | 0 |
| 229 | Tag SNPs in complement receptor-1 contribute to the susceptibility to non-small cell lung cancer. <i>Molecular Cancer</i> , 2014, 13, 56. | 7.9 | 14 |
| 230 | Quantitative assessment of the influence of common variations on 6p21 and lung cancer risk. <i>Tumor Biology</i> , 2014, 35, 689-694. | 0.8 | 7 |
| 231 | Association between TERT rs2736100 polymorphism and lung cancer susceptibility: evidence from 22 case-control studies. <i>Tumor Biology</i> , 2014, 35, 4435-4442. | 0.8 | 9 |
| 232 | Common Genetic Variants in <i>53BP1</i> Associated with Nonsmall-cell Lung Cancer Risk in Han Chinese. <i>Archives of Medical Research</i> , 2014, 45, 84-89. | 1.5 | 7 |
| 233 | Design, Analysis, and Interpretation of Genome-Wide Association Scans. <i>Statistics in the Health Sciences</i> , 2014, , . | 0.2 | 16 |
| 234 | TERT rs2736100 polymorphism contributes to lung cancer risk: a meta-analysis including 49,869 cases and 73,464 controls. <i>Tumor Biology</i> , 2014, 35, 5569-5574. | 0.8 | 12 |
| 235 | Cleft lip and palate transmembrane protein 1 rs31489 polymorphism is associated with lung cancer risk: a meta-analysis. <i>Tumor Biology</i> , 2014, 35, 5583-5588. | 0.8 | 2 |
| 236 | Lung Cancer Screening: Review and Performance Comparison Under Different Risk Scenarios. <i>Lung</i> , 2014, 192, 55-63. | 1.4 | 36 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 237 | Genetic variants in telomerase reverse transcriptase (TERT) and telomerase-associated protein 1 (TEP1) and the risk of male infertility. <i>Gene</i> , 2014, 534, 139-143. | 1.0 | 30 |
| 238 | CLPTM1L Promotes Growth and Enhances Aneuploidy in Pancreatic Cancer Cells. <i>Cancer Research</i> , 2014, 74, 2785-2795. | 0.4 | 48 |
| 239 | Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. <i>Nature Communications</i> , 2014, 5, 3365. | 5.8 | 123 |
| 240 | Genetic variability in the regulation of gene expression in ten regions of the human brain. <i>Nature Neuroscience</i> , 2014, 17, 1418-1428. | 7.1 | 620 |
| 241 | Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000. | 9.4 | 294 |
| 242 | A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. <i>Carcinogenesis</i> , 2014, 35, 1528-1535. | 1.3 | 47 |
| 243 | Genetic polymorphisms of TERT and CLPTM1L, cooking oil fume exposure, and risk of lung cancer: a caseâ€“control study in a Chinese non-smoking female population. <i>Medical Oncology</i> , 2014, 31, 114. | 1.2 | 33 |
| 244 | Lung cancer risk and genetic variants in East Asians: a meta-analysis. <i>Tumor Biology</i> , 2014, 35, 5173-5179. | 0.8 | 6 |
| 245 | Increased lung cancer risk associated with the TERT rs2736100 polymorphism: an updated meta-analysis. <i>Tumor Biology</i> , 2014, 35, 5763-5769. | 0.8 | 13 |
| 246 | Quantitative assessment of common genetic variants on chromosome 5p15 and lung cancer risk. <i>Tumor Biology</i> , 2014, 35, 6055-6063. | 0.8 | 7 |
| 247 | Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014, 46, 736-741. | 9.4 | 360 |
| 248 | A genetic sequence variant (GSV) at susceptibility loci of 5p15.33 (TERT-CLPTM1L) is associated with survival outcome in locally advanced and metastatic non-small-cell lung cancer (NSCLC). <i>Lung Cancer</i> , 2014, 84, 289-294. | 0.9 | 11 |
| 249 | Association between CLPTM1L polymorphisms (rs402710 and rs401681) and lung cancer susceptibility: evidence from 27 caseâ€“control studies. <i>Molecular Genetics and Genomics</i> , 2014, 289, 1001-1012. | 1.0 | 20 |
| 250 | Genetic polymorphisms of TERT and CLPTM1L and risk of lung cancer: a caseâ€“control study in northeast Chinese male population. <i>Medical Oncology</i> , 2014, 31, 18. | 1.2 | 17 |
| 251 | High-resolution detection of recurrent aberrations in lung adenocarcinomas by array comparative genomic hybridization and expression analysis of selective genes by quantitative PCR. <i>International Journal of Oncology</i> , 2014, 44, 2068-2076. | 1.4 | 4 |
| 252 | A Review of Cancer Risk Prediction Models with Genetic Variants. <i>Cancer Informatics</i> , 2014, 13s2, CIN.S13788. | 0.9 | 15 |
| 254 | CLPTM1L Genetic Polymorphisms and Interaction With Smoking and Alcohol Drinking in Lung Cancer Risk. <i>Medicine (United States)</i> , 2014, 93, e289. | 0.4 | 16 |
| 255 | Estrogen Receptor Gene Polymorphisms and Lung Adenocarcinoma Risk in Never-Smoking Women. <i>Journal of Thoracic Oncology</i> , 2015, 10, 1413-1420. | 0.5 | 17 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 256 | Potentially functional polymorphisms in PAK 1 are associated with risk of lung cancer in a Chinese population. <i>Cancer Medicine</i> , 2015, 4, 1781-1787. | 1.3 | 6 |
| 257 | Common variants identified in genome-wide association studies of testicular germ cell tumour: an update, biological insights and clinical application. <i>Andrology</i> , 2015, 3, 34-46. | 1.9 | 46 |
| 258 | The eQTL-missense polymorphisms of APOBEC3H are associated with lung cancer risk in a Han Chinese population. <i>Scientific Reports</i> , 2015, 5, 14969. | 1.6 | 15 |
| 259 | Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369. | 1.6 | 35 |
| 260 | Hard Work Ahead: Fine Mapping and Functional Follow-up of Susceptibility Alleles in Cancer GWAS. <i>Current Epidemiology Reports</i> , 2015, 2, 205-217. | 1.1 | 1 |
| 261 | Genetic variants associated with longer telomere length are associated with increased lung cancer risk among never-smoking women in Asia: a report from the female lung cancer consortium in Asia. <i>International Journal of Cancer</i> , 2015, 137, 311-319. | 2.3 | 72 |
| 262 | An exposure-weighted score test for genetic associations integrating environmental risk factors. <i>Biometrics</i> , 2015, 71, 596-605. | 0.8 | 11 |
| 263 | TERT gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , 2015, 137, 2175-2183. | 2.3 | 57 |
| 264 | F2RL3 methylation, lung cancer incidence and mortality. <i>International Journal of Cancer</i> , 2015, 137, 1739-1748. | 2.3 | 65 |
| 265 | Increased risk of developing lung cancer in Asian patients carrying the TERT rs2736098 G>A polymorphism: evidence from 3,354 cases and 3,518 controls. <i>OncoTargets and Therapy</i> , 2015, 8, 2757. | 1.0 | 5 |
| 266 | Association of genetic polymorphisms in TERT-CLPTM1L with lung cancer in a Chinese population. <i>Genetics and Molecular Research</i> , 2015, 14, 4469-4476. | 0.3 | 15 |
| 267 | The Genetic Polymorphisms of TERT-CLPTM1L and Lung Cancer Susceptibility: A Meta-Analysis. <i>Epidemiology (Sunnyvale, Calif)</i> , 2015, 05, . | 0.3 | 0 |
| 268 | Association of TERT Polymorphisms with Clinical Outcome of Non-Small Cell Lung Cancer Patients. <i>PLoS ONE</i> , 2015, 10, e0129232. | 1.1 | 11 |
| 269 | A Regulatory MDM4 Genetic Variant Locating in the Binding Sequence of Multiple MicroRNAs Contributes to Susceptibility of Small Cell Lung Cancer. <i>PLoS ONE</i> , 2015, 10, e0135647. | 1.1 | 32 |
| 270 | The Association between Telomere Length and Cancer Prognosis: Evidence from a Meta-Analysis. <i>PLoS ONE</i> , 2015, 10, e0133174. | 1.1 | 59 |
| 271 | Potentially Functional Polymorphisms in POU5F1 Gene Are Associated with the Risk of Lung Cancer in Han Chinese. <i>BioMed Research International</i> , 2015, 2015, 1-7. | 0.9 | 3 |
| 272 | Telomere Length and Recurrence Risk after Curative Resection in Patients with Early-Stage Non-Small-Cell Lung Cancer: A Prospective Cohort Study. <i>Journal of Thoracic Oncology</i> , 2015, 10, 302-308. | 0.5 | 17 |
| 273 | Genetic Polymorphism, Telomere Biology and Non-Small Lung Cancer Risk. <i>Journal of Genetics and Genomics</i> , 2015, 42, 549-561. | 1.7 | 26 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 274 | Lung Cancer Risk Prediction Using Common SNPs Located in GWAS-Identified Susceptibility Regions. <i>Journal of Thoracic Oncology</i> , 2015, 10, 1538-1545. | 0.5 | 33 |
| 275 | Genotyping of wild-type cytochrome P450 2A6 and whole-gene deletion using human blood samples and a multiplex real-time polymerase chain reaction method with dual-labeled probes. <i>Clinica Chimica Acta</i> , 2015, 441, 71-74. | 0.5 | 10 |
| 276 | Systematical analyses of variants in CTCF-binding sites identified a novel lung cancer susceptibility locus among Chinese population. <i>Scientific Reports</i> , 2015, 5, 7833. | 1.6 | 16 |
| 277 | Informed Genome-Wide Association Analysis With Family History As a Secondary Phenotype Identifies Novel Loci of Lung Cancer. <i>Genetic Epidemiology</i> , 2015, 39, 197-206. | 0.6 | 11 |
| 278 | Biomarkers for Assessing Risk of Cancer. , 2015, , 317-330.e3. | | 0 |
| 279 | Effect of Occupational Exposures on Lung Cancer Susceptibility: A Study of Gene-Environment Interaction Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 570-579. | 1.1 | 14 |
| 280 | Using covariate-specific disease prevalence information to increase the power of case-control studies. <i>Biometrika</i> , 2015, 102, 169-180. | 1.3 | 36 |
| 281 | Epidemiology and Inherited Predisposition for Sporadic Pancreatic Adenocarcinoma. <i>Hematology/Oncology Clinics of North America</i> , 2015, 29, 619-640. | 0.9 | 30 |
| 282 | TERT Polymorphism rs2736100-C Is Associated with EGFR Mutation-Positive Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 5173-5180. | 3.2 | 47 |
| 283 | Germline mutations causing familial lung cancer. <i>Journal of Human Genetics</i> , 2015, 60, 597-603. | 1.1 | 20 |
| 284 | Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. <i>Human Genetics</i> , 2015, 134, 231-245. | 1.8 | 34 |
| 285 | A Genetic Lung Cancer Susceptibility Test may have a Positive Effect on Smoking Cessation. <i>Journal of Genetic Counseling</i> , 2015, 24, 522-531. | 0.9 | 5 |
| 286 | Polymorphic Variants of Cytochrome P450. <i>Advances in Pharmacology</i> , 2015, 74, 85-111. | 1.2 | 34 |
| 287 | Low-Frequency Coding Variants at 6p21.33 and 20q11.21 Are Associated with Lung Cancer Risk in Chinese Populations. <i>American Journal of Human Genetics</i> , 2015, 96, 832-840. | 2.6 | 41 |
| 289 | Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015, 36, 1314-1326. | 1.3 | 15 |
| 290 | Two Novel Variants on 13q22.1 Are Associated with Risk of Esophageal Squamous Cell Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1774-1780. | 1.1 | 12 |
| 291 | Therapeutic inhibition of TRF1 impairs the growth of p53-deficient KRas ^{G12V} induced lung cancer by induction of telomeric DNA damage. <i>EMBO Molecular Medicine</i> , 2015, 7, 930-949. | 3.3 | 45 |
| 292 | Jackknife-based gene-gene interaction tests for untyped SNPs. <i>BMC Genetics</i> , 2015, 16, 85. | 2.7 | 0 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 294 | Structure and Functions of Telomeres in Organismal Homeostasis and Disease. , 2015, , 247-283. | | 0 |
| 295 | Cross Cancer Genomic Investigation of Inflammation Pathway for Five Common Cancers: Lung, Ovary, Prostate, Breast, and Colorectal Cancer. Journal of the National Cancer Institute, 2015, 107, djv246. | 3.0 | 63 |
| 296 | Deciphering associations for lung cancer risk through imputation and analysis of 12â€™%316 cases and 16â€™%831 controls. European Journal of Human Genetics, 2015, 23, 1723-1728. | 1.4 | 22 |
| 297 | Telomere length, genetic variants and gastric cancer risk in a Chinese population. Carcinogenesis, 2015, 36, 963-970. | 1.3 | 46 |
| 298 | U-shaped association between telomere length and esophageal squamous cell carcinoma risk: a case-control study in Chinese population. Frontiers of Medicine, 2015, 9, 478-486. | 1.5 | 10 |
| 299 | Common genetic variants on 3q28 contribute to non-small cell lung cancer susceptibility: evidence from 10 case-control studies. Molecular Genetics and Genomics, 2015, 290, 573-584. | 1.0 | 4 |
| 300 | TERT polymorphisms rs2853669 and rs7726159 influence on prostate cancer risk in Russian population. Tumor Biology, 2015, 36, 841-847. | 0.8 | 23 |
| 301 | Risk of multiple myeloma is associated with polymorphisms within telomerase genes and telomere length. International Journal of Cancer, 2015, 136, E351-8. | 2.3 | 30 |
| 302 | Pancreatic Cancer Genetics. International Journal of Biological Sciences, 2016, 12, 314-325. | 2.6 | 90 |
| 303 | An Overview of Genetic Polymorphism and Lung Cancer Risk. Advances in Cancer Prevention, 2016, 01, . | 0.2 | 0 |
| 304 | Multiple-level validation identifies <i>PARK2</i> in the development of lung cancer and chronic obstructive pulmonary disease. Oncotarget, 2016, 7, 44211-44223. | 0.8 | 42 |
| 305 | Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. Oncotarget, 2016, 7, 66328-66343. | 0.8 | 88 |
| 306 | Prognostic significance of CLPTM1L expression and its effects on migration and invasion of human lung cancer cells. Cancer Biomarkers, 2016, 16, 445-452. | 0.8 | 9 |
| 307 | Combining Telomerase Reverse Transcriptase Genetic Variant rs2736100 with Epidemiologic Factors in the Prediction of Lung Cancer Susceptibility. Journal of Cancer, 2016, 7, 846-853. | 1.2 | 14 |
| 308 | The association between the TERT rs2736100 AC genotype and reduced risk of upper tract urothelial carcinomas in a Han Chinese population. Oncotarget, 2016, 7, 31972-31979. | 0.8 | 14 |
| 309 | Genomic architecture of lung cancers. Current Opinion in Oncology, 2016, 28, 52-57. | 1.1 | 9 |
| 310 | Association of human telomerase reverse transcriptase gene polymorphisms, serum levels, and telomere length with renal cell carcinoma risk and pathology. Molecular Carcinogenesis, 2016, 55, 1458-1466. | 1.3 | 33 |
| 311 | <i>TERT</i> rs2736100 A>C SNP and <i>JAK2</i> 46/1 haplotype significantly contribute to the occurrence of <i>JAK2</i> V617F and <i>CALR</i> mutated myeloproliferative neoplasms â€™ a multicentric study on 529 patients. British Journal of Haematology, 2016, 174, 218-226. | 1.2 | 32 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 312 | Elucidating Genomic Characteristics of Lung Cancer Progression from In Situ to Invasive Adenocarcinoma. <i>Scientific Reports</i> , 2016, 6, 31628. | 1.6 | 36 |
| 313 | Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , 2016, 98, 33-42. | 0.9 | 49 |
| 314 | Converging findings from linkage and association analyses on susceptibility genes for smoking and other addictions. <i>Molecular Psychiatry</i> , 2016, 21, 992-1008. | 4.1 | 33 |
| 315 | Risk factors for lung cancer worldwide. <i>European Respiratory Journal</i> , 2016, 48, 889-902. | 3.1 | 546 |
| 316 | Polymorphisms in cancer-related pathway genes and lung cancer. <i>European Respiratory Journal</i> , 2016, 48, 1184-1191. | 3.1 | 5 |
| 317 | Nicotinic acetylcholine receptors and cancer. <i>Biomedical Reports</i> , 2016, 4, 515-518. | 0.9 | 48 |
| 318 | Systematic Review of Genetic Variation in Chromosome 5p15.33 and Telomere Length as Predictive and Prognostic Biomarkers for Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1537-1549. | 1.1 | 17 |
| 319 | Lung cancer incidence in never smokers: Genetic and gender basis. <i>Gene Reports</i> , 2016, 4, 198-207. | 0.4 | 11 |
| 321 | Genome-wide association analyses identify new susceptibility loci for oral cavity and pharyngeal cancer. <i>Nature Genetics</i> , 2016, 48, 1544-1550. | 9.4 | 164 |
| 322 | A Novel Genetic Variant in Long Non-coding RNA Gene NEXN-AS1 is Associated with Risk of Lung Cancer. <i>Scientific Reports</i> , 2016, 6, 34234. | 1.6 | 48 |
| 323 | Functional evaluation of TERT-CLPTM1L genetic variants associated with susceptibility of papillary thyroid carcinoma. <i>Scientific Reports</i> , 2016, 6, 26037. | 1.6 | 22 |
| 324 | Different dietary patterns and reduction of lung cancer risk: A large case-control study in the U.S.. <i>Scientific Reports</i> , 2016, 6, 26760. | 1.6 | 18 |
| 325 | A systematic comparison of copy number alterations in four types of female cancer. <i>BMC Cancer</i> , 2016, 16, 913. | 1.1 | 13 |
| 326 | Genetic determinants of cytochrome P450 2A6 activity and biomarkers of tobacco smoke exposure in relation to risk of lung cancer development in the Shanghai cohort study. <i>International Journal of Cancer</i> , 2016, 138, 2161-2171. | 2.3 | 38 |
| 327 | Gene mutation discovery research of non-smoking lung cancer patients due to indoor radon exposure. <i>Annals of Occupational and Environmental Medicine</i> , 2016, 28, 13. | 0.3 | 16 |
| 328 | Genetic variant in DNA repair gene <i>GTF2H4</i> is associated with lung cancer risk: a large-scale analysis of six published GWAS datasets in the TRICL consortium. <i>Carcinogenesis</i> , 2016, 37, 888-896. | 1.3 | 15 |
| 329 | Fine mapping of chromosome 5p15.33 based on a targeted deep sequencing and high density genotyping identifies novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2016, 37, 96-105. | 1.3 | 36 |
| 330 | CLPTM1L polymorphism as a protective factor for lung cancer: a case-control study in southern Chinese population. <i>Tumor Biology</i> , 2016, 37, 10533-10538. | 0.8 | 6 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 331 | Exome-wide analysis of rare coding variation identifies novel associations with COPD and airflow limitation in <i>MOCS3</i> , <i>IFIT3</i> and <i>SERPINA12</i> . <i>Thorax</i> , 2016, 71, 501-509. | 2.7 | 22 |
| 332 | Polymorphisms of the centrosomal gene (<i>FGFR1OP</i>) and lung cancer risk: a meta-analysis of 14 463 cases and 44 188 controls. <i>Carcinogenesis</i> , 2016, 37, 280-289. | 1.3 | 7 |
| 333 | Understanding TERT Promoter Mutations: A Common Path to Immortality. <i>Molecular Cancer Research</i> , 2016, 14, 315-323. | 1.5 | 222 |
| 334 | Novel Anti-CRR9/CLPTM1L Antibodies with Antitumorigenic Activity Inhibit Cell Surface Accumulation, PI3K Interaction, and Survival Signaling. <i>Molecular Cancer Therapeutics</i> , 2016, 15, 985-997. | 1.9 | 15 |
| 336 | Co-occurrence of Myeloproliferative Neoplasms and Solid Tumors Is Attributed to a Synergism Between Cytoreductive Therapy and the Common <i>TERT</i> Polymorphism rs2736100. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 98-104. | 1.1 | 21 |
| 337 | Common Pathways in IPF and Lung Cancer. , 2016, , 217-247. | | 0 |
| 338 | Association of the functional BCL-2 rs2279115 genetic variant and small cell lung cancer. <i>Tumor Biology</i> , 2016, 37, 1693-1698. | 0.8 | 8 |
| 339 | Epidemiology of Lung Cancer. , 2016, , 927-939.e5. | | 5 |
| 340 | Leucocyte telomere length, genetic variants at the <i>TERT</i> gene region and risk of pancreatic cancer. <i>Gut</i> , 2017, 66, 1116-1122. | 6.1 | 39 |
| 341 | A polymorphism in miR-1262 regulatory region confers the risk of lung cancer in Chinese population. <i>International Journal of Cancer</i> , 2017, 141, 958-966. | 2.3 | 26 |
| 342 | Genetic variants of PTPN2 are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. <i>Scientific Reports</i> , 2017, 7, 825. | 1.6 | 10 |
| 343 | Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. <i>Nature Communications</i> , 2017, 8, 15034. | 5.8 | 40 |
| 344 | Fine mapping of chromosome 5p15.33 identifies novel lung cancer susceptibility loci in Han Chinese. <i>International Journal of Cancer</i> , 2017, 141, 447-456. | 2.3 | 17 |
| 345 | Functional variants in DCAF4 associated with lung cancer risk in European populations. <i>Carcinogenesis</i> , 2017, 38, 541-551. | 1.3 | 16 |
| 346 | Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. <i>Scientific Reports</i> , 2017, 7, 44634. | 1.6 | 10 |
| 347 | The associations of TERT-CLPTM1L variants and TERT mRNA expression with the prognosis of early stage non-small cell lung cancer. <i>Cancer Gene Therapy</i> , 2017, 24, 20-27. | 2.2 | 14 |
| 348 | An evaluation study of reported pancreatic adenocarcinoma risk-associated SNPs from genome-wide association studies in Chinese population. <i>Pancreatology</i> , 2017, 17, 931-935. | 0.5 | 6 |
| 349 | Genetic polymorphisms and lung cancer risk: Evidence from meta-analyses and genome-wide association studies. <i>Lung Cancer</i> , 2017, 113, 18-29. | 0.9 | 36 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 350 | Pharmacotherapy for smoking cessation: effects by subgroup defined by genetically informed biomarkers. The Cochrane Library, 2018, 2018, CD011823. | 1.5 | 28 |
| 351 | Quantifying the Genetic Correlation between Multiple Cancer Types. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1427-1435. | 1.1 | 48 |
| 352 | A GWAS in uveal melanoma identifies risk polymorphisms in the CLPTM1L locus. Npj Genomic Medicine, 2017, 2, . | 1.7 | 17 |
| 353 | Differences in Toxicological and Pharmacological Responses Mediated by Polymorphic Cytochromes P450 and Related Drug-Metabolizing Enzymes. Chemical Research in Toxicology, 2017, 30, 53-60. | 1.7 | 9 |
| 354 | Susceptibility loci of <i>CNOT6</i> in the general mRNA degradation pathway and lung cancer risk—A reanalysis of eight GWASs. Molecular Carcinogenesis, 2017, 56, 1227-1238. | 1.3 | 10 |
| 355 | Atlas of human diseases influenced by genetic variants with extreme allele frequency differences. Human Genetics, 2017, 136, 39-54. | 1.8 | 15 |
| 356 | A new functional <i>IDH2</i> genetic variant is associated with the risk of lung cancer. Molecular Carcinogenesis, 2017, 56, 1082-1087. | 1.3 | 7 |
| 357 | Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. Carcinogenesis, 2018, 39, 336-346. | 1.3 | 29 |
| 358 | Genome-Wide Association Studies in Glioma. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 418-428. | 1.1 | 34 |
| 359 | A Decade of GWAS Results in Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 363-379. | 1.1 | 162 |
| 360 | Genome-Wide Association Studies of Cancer in Diverse Populations. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 405-417. | 1.1 | 57 |
| 361 | Genomic profiles of lung cancer associated with idiopathic pulmonary fibrosis. Journal of Pathology, 2018, 244, 25-35. | 2.1 | 29 |
| 362 | Integrating expression-related SNPs into genome-wide gene and pathway-based analyses identified novel lung cancer susceptibility genes. International Journal of Cancer, 2018, 142, 1602-1610. | 2.3 | 14 |
| 363 | Identification of Pleiotropic Cancer Susceptibility Variants from Genome-Wide Association Studies Reveals Functional Characteristics. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 75-85. | 1.1 | 25 |
| 364 | Facteurs de risque des cancers bronchiques : tabac, exposition professionnelle et environnementale. Revue Des Maladies Respiratoires Actualites, 2018, 10, 186-191. | 0.0 | 0 |
| 365 | Fine mapping in <i>TERT</i> - <i>CLPTM1L</i> region identified three independent lung cancer susceptibility signals: A large-scale multi-ethnic population study. Molecular Carcinogenesis, 2018, 57, 1289-1299. | 1.3 | 5 |
| 366 | Genomic characterization of individuals presenting extreme phenotypes of high and low risk to develop tobacco-induced lung cancer. Cancer Medicine, 2018, 7, 3474-3483. | 1.3 | 11 |
| 367 | Germline mutations in young non-smoking women with lung adenocarcinoma. Lung Cancer, 2018, 122, 76-82. | 0.9 | 36 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 368 | rs401681 and rs402710 confer lung cancer susceptibility by regulating TERT expression instead of CLPTM1L in East Asian populations. <i>Carcinogenesis</i> , 2018, 39, 1216-1221. | 1.3 | 12 |
| 369 | Telomerase Reverse Transcriptase Polymorphism rs2736100: A Balancing Act between Cancer and Non-Cancer Disease, a Meta-Analysis. <i>Frontiers in Medicine</i> , 2018, 5, 41. | 1.2 | 23 |
| 370 | Telomere Maintenance Mechanisms in Cancer. <i>Genes</i> , 2018, 9, 241. | 1.0 | 91 |
| 371 | The JAK2 GGCC (46/1) Haplotype in Myeloproliferative Neoplasms: Causal or Random?. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1152. | 1.8 | 16 |
| 372 | Weak sharing of genetic association signals in three lung cancer subtypes: evidence at the SNP, gene, regulation, and pathway levels. <i>Genome Medicine</i> , 2018, 10, 16. | 3.6 | 32 |
| 373 | Radon Exposure-induced Genetic Variations in Lung Cancers among Never Smokers. <i>Journal of Korean Medical Science</i> , 2018, 33, e207. | 1.1 | 6 |
| 374 | Multi-Omics Analysis Reveals a HIF Network and Hub Gene EPAS1 Associated with Lung Adenocarcinoma. <i>EBioMedicine</i> , 2018, 32, 93-101. | 2.7 | 35 |
| 375 | A decade in psychiatric GWAS research. <i>Molecular Psychiatry</i> , 2019, 24, 378-389. | 4.1 | 78 |
| 376 | Papillary Thyroid Carcinoma Variants are Characterized by Co-dysregulation of Immune and Cancer Associated Genes. <i>Cancers</i> , 2019, 11, 1179. | 1.7 | 19 |
| 377 | Development of a general logistic model for disease risk prediction using multiple SNPs. <i>FEBS Open Bio</i> , 2019, 9, 2006-2012. | 1.0 | 3 |
| 378 | TERT and JAK2 polymorphisms define genetic predisposition to myeloproliferative neoplasms in Japanese patients. <i>International Journal of Hematology</i> , 2019, 110, 690-698. | 0.7 | 5 |
| 379 | Telomerase and Telomeres in Endometrial Cancer. <i>Frontiers in Oncology</i> , 2019, 9, 344. | 1.3 | 20 |
| 380 | Genetic variant rs10937405 of TP63 and susceptibility to lung cancer risk in north Indian population. <i>Journal of Genetics</i> , 2019, 98, 1. | 0.4 | 2 |
| 381 | Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. <i>Oncotarget</i> , 2019, 10, 1760-1774. | 0.8 | 25 |
| 382 | Implementation planning for lung cancer screening in China. <i>Precision Clinical Medicine</i> , 2019, 2, 13-44. | 1.3 | 28 |
| 383 | Precision oncology of lung cancer: genetic and genomic differences in Chinese population. <i>Npj Precision Oncology</i> , 2019, 3, 14. | 2.3 | 31 |
| 384 | Cross-Cancer Pleiotropic Associations with Lung Cancer Risk in African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 715-723. | 1.1 | 11 |
| 385 | Integrated Somatic and Germline Whole-Exome Sequencing Analysis in Women with Lung Cancer after a Previous Breast Cancer. <i>Cancers</i> , 2019, 11, 441. | 1.7 | 3 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 386 | The TERT locus genotypes of rs2736100-CC/CA and rs2736098-AA predict shorter survival in renal cell carcinoma. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2019, 37, 301.e1-301.e10. | 0.8 | 15 |
| 387 | Prediagnostic Leukocyte Telomere Length and Pancreatic Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1868-1875. | 1.1 | 17 |
| 388 | Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. <i>International Journal of Epidemiology</i> , 2019, 48, 751-766. | 0.9 | 32 |
| 389 | Comprehensive characterization of functional eRNAs in lung adenocarcinoma reveals novel regulators and a prognosis-related molecular subtype. <i>Theranostics</i> , 2020, 10, 11264-11277. | 4.6 | 20 |
| 390 | Epigenetic Inactivation of the Tumor Suppressor IRX1 Occurs Frequently in Lung Adenocarcinoma and Its Silencing Is Associated with Impaired Prognosis. <i>Cancers</i> , 2020, 12, 3528. | 1.7 | 13 |
| 391 | Genetic Determinants of Lung Cancer Prognosis in Never Smokers: A Pooled Analysis in the International Lung Cancer Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1983-1992. | 1.1 | 10 |
| 392 | Association of strong risk of <i>hTERT</i> gene polymorphic variants to malignant glioma and its prognostic implications with respect to different histological types and survival of glioma cases. <i>Journal of Gene Medicine</i> , 2020, 22, e3260. | 1.4 | 5 |
| 393 | Barriers to Low-Dose CT Lung Cancer Screening among Middle-Aged Chinese. <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 7107. | 1.2 | 7 |
| 394 | Pulmonary Nodules—An Epidemic? Work Up and Management, Specific, and Unique Issues in the Elderly. <i>Current Geriatrics Reports</i> , 2020, 9, 107-112. | 1.1 | 0 |
| 395 | The TERT rs2736100 Polymorphism and Susceptibility to Myeloproliferative Neoplasms: A Systematic Review and Meta-Analysis. <i>Genetic Testing and Molecular Biomarkers</i> , 2020, 24, 181-187. | 0.3 | 3 |
| 396 | <i>RAD52</i> variants influence NSCLC risk in the Chinese population in a high altitude area. <i>Therapeutic Advances in Respiratory Disease</i> , 2020, 14, 175346662091819. | 1.0 | 4 |
| 397 | The telomerase gene polymorphisms, but not telomere length, increase susceptibility to primary glomerulonephritis/end stage renal diseases in females. <i>Journal of Translational Medicine</i> , 2020, 18, 184. | 1.8 | 12 |
| 398 | Association of Relative Leucocyte Telomere Length and Gene Single Nucleotide Polymorphisms (TERT,) Tj ETQq0 0 0 rgBT /Overlock 10 T | 1.0 | 5 |
| 399 | Potentially functional genetic variants in <i>PLIN2</i> , <i>SULT2A1</i> and <i>UGT1A9</i> genes of the ketone pathway and survival of nonsmall cell lung cancer. <i>International Journal of Cancer</i> , 2020, 147, 1559-1570. | 2.3 | 8 |
| 400 | TERT-rs33963617 and CLPTM1L-rs77518573 reduce the risk of non-small cell lung cancer in Chinese population. <i>Gene</i> , 2020, 731, 144357. | 1.0 | 2 |
| 401 | Effects of ALOX5, IL6R and SFTPD gene polymorphisms on the risk of lung cancer: A case-control study in China. <i>International Immunopharmacology</i> , 2020, 79, 106155. | 1.7 | 13 |
| 402 | A Likelihood Ratio Test for Gene-Environment Interaction Based on the Trend Effect of Genotype Under an Additive Risk Model Using the Gene-Environment Independence Assumption. <i>American Journal of Epidemiology</i> , 2021, 190, 129-141. | 1.6 | 2 |
| 403 | TERT rs2736100 and TERC rs16847897 genotypes moderate the association between internalizing mental disorders and accelerated telomere length attrition among HIV+ children and adolescents in Uganda. <i>BMC Medical Genomics</i> , 2021, 14, 15. | 0.7 | 5 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 404 | CLPTM1L Is a Novel Putative Oncogene Promoting Tumorigenesis in Oral Squamous Cell Carcinoma. Cell Transplantation, 2021, 30, 096368972110459. | 1.2 | 3 |
| 405 | Polymorphisms in the Gene Encoding Caspase 8 May Predict the Response to First-Line Platinum-Based Chemotherapy in Locally Advanced or Advanced Non-Small-Cell Lung Cancer. Journal of Clinical Medicine, 2021, 10, 1126. | 1.0 | 2 |
| 407 | A Robust Test for Additive Gene-Environment Interaction Under the Trend Effect of Genotype Using an Empirical Bayes-Type Shrinkage Estimator. American Journal of Epidemiology, 2021, 190, 1948-1960. | 1.6 | 0 |
| 408 | Polymorphic tandem DNA repeats activate the human telomerase reverse transcriptase gene. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, . | 3.3 | 6 |
| 409 | International strategy in cancer epidemiology: Japan's involvement in global projects and future role. Global Health & Medicine, 2021, 3, 187-195. | 0.6 | 2 |
| 410 | Locally advanced undifferentiated sarcomatoid carcinoma of the right maxillary sinus with PDCD6-TERT fusion: A rare case report. Oral Oncology, 2021, 124, 105466. | 0.8 | 0 |
| 411 | Familial Melanoma and Susceptibility Genes: A Review of the Most Common Clinical and Dermoscopic Phenotypic Aspect, Associated Malignancies and Practical Tips for Management. Journal of Clinical Medicine, 2021, 10, 3760. | 1.0 | 19 |
| 412 | Analysis of MNS16A VNTR polymorphic sequence variations of the TERT gene and associated risk for development of bladder cancer. Current Urology, 2021, Publish Ahead of Print, 225-230. | 0.4 | 1 |
| 413 | Molecular Basis of Lung Carcinogenesis. , 2017, , 447-496. | | 4 |
| 414 | Genome-wide association study of familial lung cancer. Carcinogenesis, 2018, 39, 1135-1140. | 1.3 | 42 |
| 415 | Cigarette Smoking Associated With Lung Adenocarcinoma In Situ in a Large Case-Control Study (SFBALCS). Journal of Thoracic Oncology, 2012, 7, 1352-1360. | 0.5 | 14 |
| 417 | Potential Genes Associated with the Survival of Lung Adenocarcinoma Were Identified by Methylation. Computational and Mathematical Methods in Medicine, 2020, 2020, 1-13. | 0.7 | 4 |
| 418 | Using Prior Information from the Medical Literature in GWAS of Oral Cancer Identifies Novel Susceptibility Variant on Chromosome 4 - the AdAPT Method. PLoS ONE, 2012, 7, e36888. | 1.1 | 17 |
| 419 | Genome-Wide Search for Gene-Gene Interactions in Colorectal Cancer. PLoS ONE, 2012, 7, e52535. | 1.1 | 35 |
| 420 | CLPTM1L Is Overexpressed in Lung Cancer and Associated with Apoptosis. PLoS ONE, 2012, 7, e52598. | 1.1 | 41 |
| 421 | Longer Telomere Length in Peripheral White Blood Cells Is Associated with Risk of Lung Cancer and the rs2736100 (CLPTM1L-TERT) Polymorphism in a Prospective Cohort Study among Women in China. PLoS ONE, 2013, 8, e59230. | 1.1 | 106 |
| 422 | Pathway-Based Analysis Using Genome-wide Association Data from a Korean Non-Small Cell Lung Cancer Study. PLoS ONE, 2013, 8, e65396. | 1.1 | 22 |
| 423 | The Impact of Polymorphic Variations in the 5p15, 6p12, 6p21 and 15q25 Loci on the Risk and Prognosis of Portuguese Patients with Non-Small Cell Lung Cancer. PLoS ONE, 2013, 8, e72373. | 1.1 | 26 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 424 | The SNP rs402710 in 5p15.33 Is Associated with Lung Cancer Risk: A Replication Study in Chinese Population and a Meta-Analysis. PLoS ONE, 2013, 8, e76252. | 1.1 | 13 |
| 425 | Genetic Variant rs401681 at 5p15.33 Modifies Susceptibility to Lung Cancer but Not Esophageal Squamous Cell Carcinoma. PLoS ONE, 2013, 8, e84277. | 1.1 | 10 |
| 426 | A Two-Dimensional Pooling Strategy for Rare Variant Detection on Next-Generation Sequencing Platforms. PLoS ONE, 2014, 9, e93455. | 1.1 | 18 |
| 427 | TERT-CLPTM1L Rs401681 C>T Polymorphism Was Associated with a Decreased Risk of Esophageal Cancer in a Chinese Population. PLoS ONE, 2014, 9, e100667. | 1.1 | 16 |
| 428 | ePIANNO: ePIgenomics ANNOtation tool. PLoS ONE, 2016, 11, e0148321. | 1.1 | 4 |
| 429 | Risk assessment models for genetic risk predictors of lung cancer using two-stage replication for Asian and European populations. Oncotarget, 2017, 8, 53959-53967. | 0.8 | 11 |
| 430 | RTEL1 polymorphisms are associated with lung cancer risk in the Chinese Han population. Oncotarget, 2016, 7, 70475-70480. | 0.8 | 10 |
| 431 | Association of polymorphisms in the telomere-related gene ACYP2 with lung cancer risk in the Chinese Han population. Oncotarget, 2016, 7, 87473-87478. | 0.8 | 13 |
| 432 | The TERT promoter mutation incidence is modified by germline TERT rs2736098 and rs2736100 polymorphisms in hepatocellular carcinoma. Oncotarget, 2017, 8, 23120-23129. | 0.8 | 34 |
| 433 | <i>TERT</i> promoter mutations and polymorphisms as prognostic factors in primary glioblastoma. Oncotarget, 2015, 6, 16663-16673. | 0.8 | 100 |
| 434 | Common variations in TERT-CLPTM1L locus are reproducibly associated with the risk of nasopharyngeal carcinoma in Chinese populations. Oncotarget, 2016, 7, 759-770. | 0.8 | 18 |
| 435 | The identification of two regulatory ESCC susceptibility genetic variants in the <i>TERT-CLPTM1L</i> loci. Oncotarget, 2016, 7, 5495-5506. | 0.8 | 19 |
| 436 | Association study of nicotinic acetylcholine receptor genes identifies a novel lung cancer susceptibility locus near CHRNA1 in African-Americans. Oncotarget, 2012, 3, 1428-1438. | 0.8 | 11 |
| 437 | Prevalence and clinical significance of pathogenic germline BRCA1/2 mutations in Chinese non-small cell lung cancer patients. Cancer Biology and Medicine, 2019, 16, 556-564. | 1.4 | 36 |
| 438 | Lung Adenocarcinoma in Never Smokers: Problems of Primary Prevention from Aspects of Susceptible Genes and Carcinogens. Anticancer Research, 2016, 36, 6207-6224. | 0.5 | 30 |
| 439 | Inherited lung cancer: a review. Ecancermedicalsecience, 2020, 14, 1008. | 0.6 | 20 |
| 440 | Genetic susceptibility to lung cancer and co-morbidities. Journal of Thoracic Disease, 2013, 5 Suppl 5, S454-62. | 0.6 | 49 |
| 441 | Different Effects of <i>TERT</i> , <i>TP63</i> , and <i>CYP2A6</i> Polymorphism on Individual Risk of Tobacco-Related Lung Cancer in Male Japanese Smokers. Journal of Cancer Therapy, 2011, 02, 690-696. | 0.1 | 6 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|-----|-----------|
| 442 | Effects of ADH1C, ALDH2, and CYP2A6 Polymorphisms on Individual Risk of Tobacco-Related Lung Cancer in Male Japanese Smokers. <i>Journal of Cancer Therapy</i> , 2013, 04, 29-35. | 0.1 | 5 |
| 443 | A hybrid parametric and empirical likelihood model for evaluating interactions in case-control studies. <i>Statistics and Its Interface</i> , 2016, 9, 147-158. | 0.2 | 1 |
| 444 | The MNS16A VNTR polymorphism of the TERT gene in bladder cancer. <i>Turkish Journal of Urology</i> , 2020, 46, 44-49. | 1.3 | 5 |
| 445 | Acetylcholine receptor pathway in lung cancer: New twists to an old story. <i>World Journal of Clinical Oncology</i> , 2014, 5, 667. | 0.9 | 23 |
| 446 | East meets West: ethnic differences in epidemiology and clinical behaviors of lung cancer between East Asians and Caucasians. <i>Chinese Journal of Cancer</i> , 2011, 30, 287-292. | 4.9 | 205 |
| 447 | Application of Cancer Genomics to Solve Unmet Clinical Needs. <i>Genomics and Informatics</i> , 2013, 11, 174. | 0.4 | 9 |
| 448 | Apoptosis-related single nucleotide polymorphisms and the risk of non-small cell lung cancer in women. <i>Journal of Cancer Therapeutics & Research</i> , 2014, 3, 1. | 1.2 | 31 |
| 449 | Association of TERT rs2736098 Polymorphism with Cancer Risk: a Meta-analysis. <i>Asian Pacific Journal of Cancer Prevention</i> , 2012, 13, 4943-4946. | 0.5 | 11 |
| 450 | TERT rs2736098 Polymorphism and Cancer Risk: Results of a Meta-analysis. <i>Asian Pacific Journal of Cancer Prevention</i> , 2012, 13, 3483-3488. | 0.5 | 18 |
| 451 | TP63 Gene Polymorphisms, Cooking Oil Fume Exposure and Risk of Lung Adenocarcinoma in Chinese Non-smoking Females. <i>Asian Pacific Journal of Cancer Prevention</i> , 2013, 14, 6519-6522. | 0.5 | 12 |
| 452 | No Association of XRCC1 and CLPTM1L Polymorphisms with Non-small Cell Lung Cancer in a Non-Smoking Han Chinese Population. <i>Asian Pacific Journal of Cancer Prevention</i> , 2013, 14, 5171-5174. | 0.5 | 17 |
| 453 | Genetic Variant in CLPTM1L Confers Reduced Risk of Lung Cancer: a Replication Study in Chinese and a Meta-analysis. <i>Asian Pacific Journal of Cancer Prevention</i> , 2014, 15, 9241-9247. | 0.5 | 4 |
| 454 | Genetic Variations in TERT-CLPTM1L Genes and Risk of Lung Cancer in a Chinese Population. <i>Asian Pacific Journal of Cancer Prevention</i> , 2014, 15, 2809-2813. | 0.5 | 19 |
| 455 | Genome-wide analysis of runs of homozygosity identifies new susceptibility regions of lung cancer in Han Chinese. <i>Journal of Biomedical Research</i> , 2013, 27, 208. | 0.7 | 18 |
| 456 | Analysis of Population-Based Genetic Association Studies Applied to Cancer Susceptibility and Prognosis. , 2009, , 149-191. | | 0 |
| 458 | Pharmacogenetics of Lung Cancer. , 2010, , 87-106. | | 0 |
| 460 | Assessing the impact of nicotine dependence genes on the risk of facial clefts: An example of the use of national registry and biobank data. <i>Norsk Epidemiologi</i> , 2012, 21, 241-250. | 0.2 | 5 |
| 461 | A copula-model based semiparametric interaction test under the casecontrol design. <i>Statistica Sinica</i> , 2013, 23, 1505-1521. | 0.2 | 1 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 462 | An Introduction to Association Analysis. Statistics in the Health Sciences, 2014, , 79-133. | 0.2 | 0 |
| 463 | Lung Cancer: Genetic Susceptibility. , 2014, , 231-241. | | 0 |
| 464 | Discovery and Characterization of Cancer Genetic Susceptibility Alleles. , 2014, , 309-321.e3. | | 0 |
| 465 | Genetic Susceptibility to Lung Adenocarcinoma. Genes and Environment, 2014, 36, 160-166. | 0.9 | 0 |
| 466 | Characterization of population-based variation and putative functional elements for the multiple-cancer susceptibility loci at 5p15.33. F1000Research, 2014, 3, 231. | 0.8 | 0 |
| 467 | TERT polymorphisms rs2853669, rs2736100, rs7726159 and rs10069690 and the risk of non-Hodgkinâ€™s lymphoma in ethnical Russians. Telomere and Telomerase, 0, , . | 0.0 | 1 |
| 468 | Lung Cancer Among Asian Americans. , 2016, , 107-136. | | 0 |
| 469 | Genetic Susceptibility to Lung Cancer. Molecular Pathology Library, 2018, , 19-43. | 0.1 | 0 |
| 471 | Lung Cancer: Genetic Susceptibility. , 2020, , 273-286. | | 0 |
| 472 | The effect of susceptibility variants, identified in never-smoking female lung cancer cases, on male smokers. Korean Journal of Internal Medicine, 2020, 35, 929-935. | 0.7 | 3 |
| 473 | The current and future roles of genomics. , 0, , 79-94. | | 0 |
| 476 | Telomere length and variation in telomere biology genes in individuals with osteosarcoma. International Journal of Molecular Epidemiology and Genetics, 2011, 2, 19-29. | 0.4 | 27 |
| 477 | Germ line variation in nucleotide excision repair genes and lung cancer risk in smokers. International Journal of Molecular Epidemiology and Genetics, 2012, 3, 1-17. | 0.4 | 35 |
| 478 | Genetic variation in telomere maintenance genes in relation to ovarian cancer survival. International Journal of Molecular Epidemiology and Genetics, 2012, 3, 252-61. | 0.4 | 2 |
| 480 | Identification of an SCLC susceptibility rs7963551 genetic polymorphism in a previously GWAS-identified 12p13.33 RAD52 lung cancer risk locus in the Chinese population. International Journal of Clinical and Experimental Medicine, 2015, 8, 16528-35. | 1.3 | 6 |
| 482 | Assessment of Telomerase Reverse Transcriptase Single Nucleotide Polymorphism in Sleep Bruxism. Journal of Clinical Medicine, 2022, 11, 525. | 1.0 | 6 |
| 483 | The association of telomere maintenance and TERT expression with susceptibility to human papillomavirus infection in cervical epithelium. Cellular and Molecular Life Sciences, 2022, 79, 110. | 2.4 | 3 |
| 484 | Genetic variant rs10937405 of TP63 and susceptibility to lung cancer risk in north Indian population. Journal of Genetics, 2019, 98, . | 0.4 | 1 |

| # | ARTICLE | IF | CITATIONS |
|-----|---|------|-----------|
| 488 | A Large-Scale Genome-Wide Gene-Gene Interaction Study of Lung Cancer Susceptibility in Europeans With a Trans-Ethnic Validation in Asians. <i>Journal of Thoracic Oncology</i> , 2022, 17, 974-990. | 0.5 | 18 |
| 489 | Genetic Analysis of Lung Cancer and the Germline Impact on Somatic Mutation Burden. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1159-1166. | 3.0 | 8 |
| 491 | Cumulative Evidence for Relationships Between Multiple Variants in the TERT and CLPTM1L Region and Risk of Cancer and Non-Cancer Disease. <i>Frontiers in Oncology</i> , 0, 12, . | 1.3 | 8 |
| 492 | Functional studies of lung cancer GWAS beyond association. <i>Human Molecular Genetics</i> , 2022, 31, R22-R36. | 1.4 | 8 |
| 493 | MicrobiomeGWAS: A Tool for Identifying Host Genetic Variants Associated with Microbiome Composition. <i>Genes</i> , 2022, 13, 1224. | 1.0 | 9 |
| 494 | Cross-ancestry genome-wide meta-analysis of 61,047 cases and 947,237 controls identifies new susceptibility loci contributing to lung cancer. <i>Nature Genetics</i> , 2022, 54, 1167-1177. | 9.4 | 35 |
| 496 | Analyses of rare predisposing variants of lung cancer in 6,004 whole genomes in Chinese. <i>Cancer Cell</i> , 2022, 40, 1223-1239.e6. | 7.7 | 23 |
| 497 | Identification of rs2736099 as a novel cis-regulatory variation for TERT and implications for tumorigenesis and cell proliferation. <i>Journal of Cancer Research and Clinical Oncology</i> , 0, , . | 1.2 | 1 |
| 498 | Genomic landscape of lung adenocarcinomas in different races. <i>Frontiers in Oncology</i> , 0, 12, . | 1.3 | 6 |
| 499 | Etiology of lung cancer: Evidence from epidemiologic studies. <i>Journal of the National Cancer Center</i> , 2022, 2, 216-225. | 3.0 | 2 |
| 501 | Implications of risk conferred by 5p15.33 loci genetic variants; human telomerase reverse transcriptase rs2736098 and rs2736100 in predisposition of bladder cancer. <i>Reports of Practical Oncology and Radiotherapy</i> , 2022, 27, 787-796. | 0.3 | 0 |
| 502 | Liquid Biopsy for Lung Cancer: Up-to-Date and Perspectives for Screening Programs. <i>International Journal of Molecular Sciences</i> , 2023, 24, 2505. | 1.8 | 21 |
| 504 | Common genetic variations in telomere length genes and lung cancer: a Mendelian randomisation study and its novel application in lung tumour transcriptome. <i>ELife</i> , 0, 12, . | 2.8 | 3 |
| 506 | Lung Cancer in Developing Countries. , 2023, , 1-28. | | 0 |
| 513 | Lung cancer in patients who have never smoked " an emerging disease. <i>Nature Reviews Clinical Oncology</i> , 2024, 21, 121-146. | 12.5 | 1 |