

# Zhengdong Zhang

## List of Publications by Year in descending order

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Version: 2024-02-01

158  
papers

4,462  
citations

117625

34  
h-index

149698

56  
g-index

160  
all docs

160  
docs citations

160  
times ranked

6851  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Genetic variants in splicing factor genes and susceptibility to bladder cancer. <i>Gene</i> , 2022, 809, 146022.  | 2.2  | 1         |
| 2  | Fine Particulate Matter Induces Childhood Asthma Attacks via Extracellular Vesicleâ€Packaged Letâ€7iâ€5pâ€Mediated Modulation of the MAPK Signaling Pathway. <i>Advanced Science</i> , 2022, 9, e2102460.                         | 11.2 | 21        |
| 3  | Global internet search trends related to gastrointestinal symptoms predict regional COVID-19 outbreaks. <i>Journal of Infection</i> , 2022, 84, 56-63.  | 3.3  | 8         |
| 4  | Genome-Wide Association Analyses Identify <i>CATSPERE</i> as a Mediator of Colorectal Cancer Susceptibility and Progression. <i>Cancer Research</i> , 2022, 82, 986-997.  | 0.9  | 3         |
| 5  | High-density lipoprotein, low-density lipoprotein and triglyceride levels and upper gastrointestinal cancers risk: a trans-ancestry Mendelian randomization study. <i>European Journal of Clinical Nutrition</i> , 2022, , .          | 2.9  | 4         |
| 6  | Genetic variants in the Hedgehog signaling pathway genes are associated with gastric cancer risk in a Chinese Han population. <i>Journal of Biomedical Research</i> , 2022, 36, 22.   | 1.6  | 0         |
| 7  | Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3â€eIF3h interaction. <i>Molecular Cancer</i> , 2022, 21, 49.  | 19.2 | 72        |
| 8  | Genetic variants in choline metabolism pathway are associated with the risk of bladder cancer in the Chinese population. <i>Archives of Toxicology</i> , 2022, , 1.   | 4.2  | 1         |
| 9  | Evaluation of genetic variants in nucleosome remodeling and deacetylase (NuRD) complex subunits encoding genes and gastric cancer susceptibility. <i>Archives of Toxicology</i> , 2022, 96, 1739-1749.                                | 4.2  | 2         |
| 10 | Association between circulating vitamin E and ten common cancers: evidence from large-scale Mendelian randomization analysis and a longitudinal cohort study. <i>BMC Medicine</i> , 2022, 20, 168.                                    | 5.5  | 23        |
| 11 | Fine particulate matter induces METTL3-mediated m6A modification of BIRC5 mRNA in bladder cancer. <i>Journal of Hazardous Materials</i> , 2022, 437, 129310.  | 12.4 | 19        |
| 12 | Metabolomics identifying biomarkers of PM2.5 exposure for vulnerable population: based on a prospective cohort study. <i>Environmental Science and Pollution Research</i> , 2021, 28, 14586-14596.                                    | 5.3  | 16        |
| 13 | Systematic evaluation of the effects of genetic variants on PIWI-interacting RNA expression across 33 cancer types. <i>Nucleic Acids Research</i> , 2021, 49, 90-97.  | 14.5 | 22        |
| 14 | A prospective study of the associations among fine particulate matter, genetic variants, and the risk of colorectal cancer. <i>Environment International</i> , 2021, 147, 106309.   | 10.0 | 14        |
| 15 | Genetic variants in N6-methyladenosine are associated with bladder cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021, 95, 299-309.  | 4.2  | 18        |
| 16 | Relationship between particulate matter exposure and female breast cancer incidence and mortality: a systematic review and meta-analysis. <i>International Archives of Occupational and Environmental Health</i> , 2021, 94, 191-201. | 2.3  | 19        |
| 17 | Identification of low-frequency variants of UGT1A3 associated with bladder cancer risk by next-generation sequencing. <i>Oncogene</i> , 2021, 40, 2382-2394.  | 5.9  | 8         |
| 18 | Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of IQCB1 expression. <i>Archives of Toxicology</i> , 2021, 95, 2053-2063.         | 4.2  | 5         |

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|----|--|------|-----------|
| 19 | Effect of PM2.5 exposure on circulating fibrinogen and IL-6 levels: A systematic review and meta-analysis. <i>Chemosphere</i> , 2021, 271, 129565.   | 8.2  | 23        |
| 20 | Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021, 95, 2523-2532.                                  | 4.2  | 4         |
| 21 | Integrative omics provide biological and clinical insights into acute respiratory distress syndrome. <i>Intensive Care Medicine</i> , 2021, 47, 761-771.   | 8.2  | 19        |
| 22 | The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021, 14, 93.  | 17.0 | 31        |
| 23 | CoSMed: a user-friendly web server to estimate 5-year survival probability of left-sided and right-sided colorectal cancer patients using molecular data. <i>Bioinformatics</i> , 2021, 38, 278-281. | 4.1  | 5         |
| 24 | Identification of common genetic variants associated with serum concentrations of p, p̂²-DDE in non-occupational populations in eastern China. <i>Environment International</i> , 2021, 152, 106507. | 10.0 | 3         |
| 25 | METTL3 regulates PM2.5-induced cell injury by targeting OSGIN1 in human airway epithelial cells. <i>Journal of Hazardous Materials</i> , 2021, 415, 125573.  | 12.4 | 32        |
| 26 | Genetic variants in m6A regulators are associated with gastric cancer risk. <i>Archives of Toxicology</i> , 2021, 95, 1081-1088.   | 4.2  | 18        |
| 27 | Functional variants of RPS6KB1 and PIK3R1 in the autophagy pathway genes and risk of bladder cancer. <i>Archives of Toxicology</i> , 2021, , 1.  | 4.2  | 4         |
| 28 | RPTOR methylation in the peripheral blood and breast cancer in the Chinese population. <i>Genes and Genomics</i> , 2021, , 1.  | 1.4  | 3         |
| 29 | Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020, 41, 8-17.   | 2.8  | 38        |
| 30 | Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. <i>Gut</i> , 2020, 69, 641-651.                      | 12.1 | 36        |
| 31 | Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020, 94, 325-333.   | 4.2  | 3         |
| 32 | Remote modulation of lncRNA <i>GCLET</i> by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. <i>Science Advances</i> , 2020, 6, eaay5525.                                  | 10.3 | 23        |
| 33 | Genetic variants in Ras/Raf/MEK/ERK pathway are associated with gastric cancer risk in Chinese Han population. <i>Archives of Toxicology</i> , 2020, 94, 2683-2690.                                  | 4.2  | 4         |
| 34 | Genetic Variant in Long Non-Coding RNA H19 Modulates Its Expression and Predicts Renal Cell Carcinoma Susceptibility and Mortality. <i>Frontiers in Oncology</i> , 2020, 10, 785.                    | 2.8  | 10        |
| 35 | MUC1 is associated with TFF2 methylation in gastric cancer. <i>Clinical Epigenetics</i> , 2020, 12, 37.  | 4.1  | 14        |
| 36 | Genetic variations in Hippo pathway genes influence bladder cancer risk in a Chinese population. <i>Archives of Toxicology</i> , 2020, 94, 785-794.  | 4.2  | 11        |

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|----|---|------|-----------|
| 37 | A transcriptomic study for identifying cardia- and non-cardia-specific gastric cancer prognostic factors using genetic algorithm-based methods. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 9457-9465.  | 3.6  | 3         |
| 38 | Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020, 137, 105543.  | 10.0 | 16        |
| 39 | Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. <i>Gene</i> , 2020, 736, 144447.   | 2.2  | 5         |
| 40 | Multiomics Evaluation of Gastrointestinal and Other Clinical Characteristics of COVID-19. <i>Gastroenterology</i> , 2020, 158, 2298-2301.e7.  | 1.3  | 117       |
| 41 | Alternative splicing related genetic variants contribute to bladder cancer risk. <i>Molecular Carcinogenesis</i> , 2020, 59, 923-929.   | 2.7  | 27        |
| 42 | A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. <i>Journal of Gastroenterology</i> , 2019, 54, 141-148.   | 5.1  | 9         |
| 43 | Vitamin B2 intake reduces the risk for colorectal cancer: a dose-response analysis. <i>European Journal of Nutrition</i> , 2019, 58, 1591-1602.   | 3.9  | 13        |
| 44 | Long non-coding RNA FLJ22763 is involved in the progression and prognosis of gastric cancer. <i>Gene</i> , 2019, 693, 84-91.  | 2.2  | 17        |
| 45 | Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019, 54, 939-949.   | 5.1  | 15        |
| 46 | Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. <i>International Journal of Cancer</i> , 2019, 145, 2661-2669.  | 5.1  | 25        |
| 47 | A genetic variation in the CpG island of pseudogene <i>GBAP1</i> promoter is associated with gastric cancer susceptibility. <i>Cancer</i> , 2019, 125, 2465-2473.   | 4.1  | 25        |
| 48 | The effects of particulate matters on allergic rhinitis in Nanjing, China. <i>Environmental Science and Pollution Research</i> , 2019, 26, 11452-11457.   | 5.3  | 20        |
| 49 | Genome-wide long non-coding RNAs identified a panel of novel plasma biomarkers for gastric cancer diagnosis. <i>Gastric Cancer</i> , 2019, 22, 731-741.   | 5.3  | 37        |
| 50 | Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 2012-2019.   | 3.6  | 9         |
| 51 | Polymorphism rs4787951 in IL-4R contributes to the increased risk of renal cell carcinoma in a Chinese population. <i>Gene</i> , 2019, 685, 242-247.  | 2.2  | 3         |
| 52 | Ambient fine particulate matter (PM2.5) induces oxidative stress and pro-inflammatory response via up-regulating the expression of CYP1A1/1B1 in human bronchial epithelial cells in vitro. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2019, 839, 40-48. | 1.7  | 42        |
| 53 | Association between obesity and bladder cancer recurrence: A meta-analysis. <i>Clinica Chimica Acta</i> , 2018, 480, 41-46.   | 1.1  | 28        |
| 54 | Tagging SNPs in the HOTAIR gene are associated with bladder cancer risk in a Chinese population. <i>Gene</i> , 2018, 664, 22-26.  | 2.2  | 10        |

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|----|--|------|-----------|
| 55 | A Genetic Variant Located in miR-146b Promoter Region Is Associated with Prognosis of Gastric Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 822-828.                      | 2.5  | 15        |
| 56 | Evaluation of vulnerable PM2.5-exposure individuals: a repeated-measure study in an elderly population. <i>Environmental Science and Pollution Research</i> , 2018, 25, 11833-11840.                 | 5.3  | 12        |
| 57 | Hypermethylation of EIF4E promoter is associated with early onset of gastric cancer. <i>Carcinogenesis</i> , 2018, 39, 66-71.  | 2.8  | 15        |
| 58 | Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. <i>Cancer Medicine</i> , 2018, 7, 1194-1200.                                      | 2.8  | 24        |
| 59 | Genetic variants in XDH are associated with prognosis for gastric cancer in a Chinese population. <i>Gene</i> , 2018, 663, 196-202.  | 2.2  | 3         |
| 60 | Probabilistic natural mapping of gene-level tests for genome-wide association studies. <i>Briefings in Bioinformatics</i> , 2018, 19, 545-553.   | 6.5  | 6         |
| 61 | A functional polymorphism in <i>TFF1</i> promoter is associated with the risk and prognosis of gastric cancer. <i>International Journal of Cancer</i> , 2018, 142, 1805-1816.                        | 5.1  | 22        |
| 62 | Exosome-transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression. <i>Molecular Cancer</i> , 2018, 17, 143.   | 19.2 | 217       |
| 63 | Genetic variants in PI3K/Akt/mTOR pathway genes contribute to gastric cancer risk. <i>Gene</i> , 2018, 670, 130-135.   | 2.2  | 14        |
| 64 | Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. <i>Archives of Toxicology</i> , 2018, 92, 1991-1999.                              | 4.2  | 14        |
| 65 | Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018, 92, 2681-2689.   | 4.2  | 30        |
| 66 | LncRNA <i>PCAT1</i> and its genetic variant rs1902432 are associated with prostate cancer risk. <i>Journal of Cancer</i> , 2018, 9, 1414-1420.   | 2.5  | 28        |
| 67 | Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018, 33, 82-87.   | 6.1  | 7         |
| 68 | LncRNA <i>MT1JP</i> functions as a ceRNA in regulating <i>FBXW7</i> through competitively binding to miR-92a-3p in gastric cancer. <i>Molecular Cancer</i> , 2018, 17, 87.                           | 19.2 | 218       |
| 69 | Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. <i>British Journal of Cancer</i> , 2018, 119, 130-132.  | 6.4  | 35        |
| 70 | Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018, 673, 174-180.  | 2.2  | 12        |
| 71 | Genetic Variations in the 3'UTR-untranslated Regions of Genes Involved in the Cell Cycle and Apoptosis Pathways Affect Bladder Cancer Risk. <i>Cancer Genomics and Proteomics</i> , 2018, 15, 67-72. | 2.0  | 5         |
| 72 | Genetic variants, PM2.5 exposure level and global DNA methylation level: A multi-center population-based study in Chinese. <i>Toxicology Letters</i> , 2017, 269, 77-82.                             | 0.8  | 10        |

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|----|--|------|-----------|
| 73 | Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 2011-2021.   | 1.3  | 58        |
| 74 | KCNMA1 cooperating with PTK2 is a novel tumor suppressor in gastric cancer and is associated with disease outcome. <i>Molecular Cancer</i> , 2017, 16, 46.   | 19.2 | 41        |
| 75 | HOTAIR rs7958904 polymorphism is associated with increased cervical cancer risk in a Chinese population. <i>Scientific Reports</i> , 2017, 7, 3144.  | 3.3  | 32        |
| 76 | The association of rs710886 in lncRNA PCAT1 with bladder cancer risk in a Chinese population. <i>Gene</i> , 2017, 627, 226-232.  | 2.2  | 23        |
| 77 | Mxd1 mediates hypoxia-induced cisplatin resistance in osteosarcoma cells by repression of the PTEN tumor suppressor gene. <i>Molecular Carcinogenesis</i> , 2017, 56, 2234-2244.   | 2.7  | 22        |
| 78 | Evaluation of genome-wide genotyping concordance between tumor tissues and peripheral blood. <i>Genomics</i> , 2017, 109, 108-112.   | 2.9  | 7         |
| 79 | Short-term effects of ambient air pollution and childhood lower respiratory diseases. <i>Scientific Reports</i> , 2017, 7, 4414.   | 3.3  | 41        |
| 80 | A novel mechanism of rs763110 polymorphism contributing to cervical cancer risk by affecting the binding affinity of C/EBP $\beta$ and OCT1 complex to chromatin. <i>International Journal of Cancer</i> , 2017, 140, 756-763. | 5.1  | 17        |
| 81 | Expression and prognostic value of microRNA-26a and microRNA-148a in gastric cancer. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 819-827.  | 2.8  | 35        |
| 82 | Information transduction capacity reduces the uncertainties in annotation-free isoform discovery and quantification. <i>Nucleic Acids Research</i> , 2017, 45, e143-e143.  | 14.5 | 6         |
| 83 | Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017, 8, 1355-1361.  | 2.5  | 12        |
| 84 | Mxi1-0 regulates the growth of human umbilical vein endothelial cells through extracellular signal-regulated kinase 1/2 (ERK1/2) and interleukin-8 (IL-8)-dependent pathways. <i>PLoS ONE</i> , 2017, 12, e0178831.            | 2.5  | 4         |
| 85 | The HOTAIR, PRNCR1 and POLR2E polymorphisms are associated with cancer risk: a meta-analysis. <i>Oncotarget</i> , 2017, 8, 43271-43283.  | 1.8  | 37        |
| 86 | SNORA74B gene silencing inhibits gallbladder cancer cells by inducing PHLPP and suppressing Akt/mTOR signaling. <i>Oncotarget</i> , 2017, 8, 19980-19996.  | 1.8  | 18        |
| 87 | An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017, 8, 37367-37376.   | 1.8  | 42        |
| 88 | XRCC1 mediated the development of cervical cancer through a novel Sp1/Krox-20 switch. <i>Oncotarget</i> , 2017, 8, 86217-86226.  | 1.8  | 12        |
| 89 | Association of genetic variants in lncRNA H19 with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 25470-25477.  | 1.8  | 90        |
| 90 | The association analysis of hOGG1 genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2016, 7, 66061-66068.  | 1.8  | 8         |

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|-----|---|------|-----------|
| 91  | Circulating MicroRNA-26a in Plasma and Its Potential Diagnostic Value in Gastric Cancer. PLoS ONE, 2016, 11, e0151345.  | 2.5  | 34        |
| 92  | Genetic variation in IGF1 predicts renal cell carcinoma susceptibility and prognosis in Chinese population. Scientific Reports, 2016, 6, 39014.   | 3.3  | 6         |
| 93  | Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. Nature Communications, 2016, 7, 11478.  | 12.8 | 73        |
| 94  | Genome-Wide Association Study of Bladder Cancer in a Chinese Cohort Reveals a New Susceptibility Locus at 5q12.3. Cancer Research, 2016, 76, 3277-3284.   | 0.9  | 46        |
| 95  | Genetic variants in lncRNA <i>H19</i> are associated with the risk of bladder cancer in a Chinese population. Mutagenesis, 2016, 31, 531-538.   | 2.6  | 70        |
| 96  | Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. Human Molecular Genetics, 2016, 25, ddw112.                              | 2.9  | 10        |
| 97  | The rs767649 polymorphism in the promoter of miR-155 contributes to the decreased risk for cervical cancer in a Chinese population. Gene, 2016, 595, 109-114.   | 2.2  | 18        |
| 98  | miR-107 regulates tumor progression by targeting NF1 in gastric cancer. Scientific Reports, 2016, 6, 36531.   | 3.3  | 51        |
| 99  | The influence of genetic variants of sorafenib on clinical outcomes and toxic effects in patients with advanced renal cell carcinoma. Scientific Reports, 2016, 6, 20089.                                 | 3.3  | 22        |
| 100 | PSCA rs2294008 polymorphism contributes to the decreased risk for cervical cancer in a Chinese population. Scientific Reports, 2016, 6, 23465.  | 3.3  | 14        |
| 101 | Rare variants in BRCA2 and CHEK2 are associated with the risk of urinary tract cancers. Scientific Reports, 2016, 6, 33542.   | 3.3  | 22        |
| 102 | A functional variant in <i>TP63</i> at 3q28 associated with bladder cancer risk by creating an miR-140 binding site. International Journal of Cancer, 2016, 139, 65-74.                                   | 5.1  | 27        |
| 103 | A functional variant in miR-143 promoter contributes to prostate cancer risk. Archives of Toxicology, 2016, 90, 403-414.  | 4.2  | 43        |
| 104 | Genetic variants in multisynthetase complex genes are associated with DNA damage levels in Chinese populations. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2016, 786, 8-13. | 1.0  | 6         |
| 105 | Functional annotation of colorectal cancer susceptibility loci identifies <i>MLH1</i> rs1800734 associated with MSI patients. Gut, 2016, 65, 1227-1228.   | 12.1 | 11        |
| 106 | Pri-miR-34b/c rs4938723 polymorphism contributes to acute lymphoblastic leukemia susceptibility in Chinese children. Leukemia and Lymphoma, 2016, 57, 1436-1441.  | 1.3  | 31        |
| 107 | Association Between MIF-AS rs755622 and Nephrolithiasis Risk in a Chinese Population. Medical Science Monitor, 2016, 22, 563-568.   | 1.1  | 4         |
| 108 | A genetic study and meta-analysis of the genetic predisposition of prostate cancer in a Chinese population. Oncotarget, 2016, 7, 21393-21403.   | 1.8  | 18        |



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|-----|---|------|-----------|
| 109 | Functional POR A503V is associated with the risk of bladder cancer in a Chinese population. <i>Scientific Reports</i> , 2015, 5, 11751.   | 3.3  | 18        |
| 110 | A novel antisense long noncoding RNA regulates the expression of MDC1 in bladder cancer. <i>Oncotarget</i> , 2015, 6, 484-493.  | 1.8  | 56        |
| 111 | The association analysis of lncRNA <i>HOTAIR</i> genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2015, 6, 31255-31262.  | 1.8  | 95        |
| 112 | The prognostic significance of HOTAIR for predicting clinical outcome in patients with digestive system tumors. <i>Journal of Cancer Research and Clinical Oncology</i> , 2015, 141, 2139-2145.   | 2.5  | 33        |
| 113 | Circulating miR-497 and miR-663b in plasma are potential novel biomarkers for bladder cancer. <i>Scientific Reports</i> , 2015, 5, 10437.   | 3.3  | 105       |
| 114 | Genetic variation in C12orf51 is associated with prognosis of intestinal-type gastric cancer in a Chinese population. <i>Biomedicine and Pharmacotherapy</i> , 2015, 69, 133-138.   | 5.6  | 8         |
| 115 | Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015, 121, 2044-2052.  | 4.1  | 56        |
| 116 | Genetic variants of H2AX gene were associated with P M 2.5 -modulated DNA damage levels in Chinese Han populations. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2015, 778, 41-45.                    | 1.0  | 3         |
| 117 | Personal exposure to PM2.5, genetic variants and DNA damage: A multi-center population-based study in Chinese. <i>Toxicology Letters</i> , 2015, 235, 172-178.  | 0.8  | 34        |
| 118 | Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. <i>Nature Communications</i> , 2015, 6, 8469.  | 12.8 | 51        |
| 119 | Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015, 556, 227-234.  | 2.2  | 66        |
| 120 | Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. <i>Mutagenesis</i> , 2015, 30, 303-310.  | 2.6  | 128       |
| 121 | A miR-29c binding site genetic variant in the 3'-untranslated region of LAMTOR3 gene is associated with gastric cancer risk. <i>Biomedicine and Pharmacotherapy</i> , 2015, 69, 70-75.  | 5.6  | 10        |
| 122 | Identification of novel piRNAs in bladder cancer. <i>Cancer Letters</i> , 2015, 356, 561-567.   | 7.2  | 115       |
| 123 | A Functional Ser326Cys Polymorphism in hOGG1 Is Associated with Noise-Induced Hearing Loss in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e89662.  | 2.5  | 37        |
| 124 | Clinical Significance of POU5F1P1 rs10505477 Polymorphism in Chinese Gastric Cancer Patients Receiving Cisplatin-Based Chemotherapy after Surgical Resection. <i>International Journal of Molecular Sciences</i> , 2014, 15, 12764-12777. | 4.1  | 16        |
| 125 | Associations of NR5A2 Gene Polymorphisms with the Clinicopathological Characteristics and Survival of Gastric Cancer. <i>International Journal of Molecular Sciences</i> , 2014, 15, 22902-22917.   | 4.1  | 8         |
| 126 | Variants in angiogenesis-related genes and the risk of clear cell renal cell carcinoma. <i>Mutagenesis</i> , 2014, 29, 419-425.   | 2.6  | 18        |



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|-----|---|-----|-----------|
| 127 | Genetic variations in microRNAs and the risk and survival of renal cell cancer. <i>Carcinogenesis</i> , 2014, 35, 1629-1635.  | 2.8 | 47        |
| 128 | Hsa-miR-196a2 polymorphism increases the risk of acute lymphoblastic leukemia in Chinese children. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 759, 16-21.                           | 1.0 | 35        |
| 129 | Global gene expression profiling of human bronchial epithelial cells exposed to airborne fine particulate matter collected from Wuhan, China. <i>Toxicology Letters</i> , 2014, 228, 25-33.                                     | 0.8 | 58        |
| 130 | Cumulative effect of genome-wide association study-identified genetic variants for bladder cancer. <i>International Journal of Cancer</i> , 2014, 135, 2653-2660.   | 5.1 | 31        |
| 131 | Genetic variation rs10484761 on 6p21.1 derived from a genome-wide association study is associated with gastric cancer survival in a Chinese population. <i>Gene</i> , 2014, 536, 59-64.   | 2.2 | 15        |
| 132 | Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014, 35, 2723-2730.   | 2.8 | 57        |
| 133 | A common genetic variation in the promoter of miR-107 is associated with gastric adenocarcinoma susceptibility and survival. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 769, 35-41. | 1.0 | 28        |
| 134 | Genetic variants in SMARC genes are associated with DNA damage levels in Chinese population. <i>Toxicology Letters</i> , 2014, 229, 327-332.  | 0.8 | 6         |
| 135 | Effects of TSP-1 -696 C/T polymorphism on bladder cancer susceptibility and clinicopathologic features. <i>Cancer Genetics</i> , 2014, 207, 247-252.  | 0.4 | 5         |
| 136 | Assessing the Effectiveness of Problem-Based Learning of Preventive Medicine Education in China. <i>Scientific Reports</i> , 2014, 4, 5126.   | 3.3 | 25        |
| 137 | Association study between XPC Asp1104His polymorphism and colorectal cancer risk in a Chinese population. <i>Scientific Reports</i> , 2014, 4, 6700.  | 3.3 | 23        |
| 138 | A genetic variant of miR-148a binding site in the SCRNI 3'-UTR is associated with susceptibility and prognosis of gastric cancer. <i>Scientific Reports</i> , 2014, 4, 7080.  | 3.3 | 16        |
| 139 | MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014, 4, 4851.  | 3.3 | 14        |
| 140 | Genetic Polymorphisms in IGF-I and IGFBP-3 Are Associated with Prostate Cancer in the Chinese Population. <i>PLoS ONE</i> , 2014, 9, e85609.  | 2.5 | 18        |
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