

Xiaoping Miao

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283
papers

15,350
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56
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114
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293
ext. papers

18,206
ext. citations

6.9
avg, IF

6.54
L-index

#	Paper	IF	Citations
283	Neurologic Manifestations of Hospitalized Patients With Coronavirus Disease 2019 in Wuhan, China. <i>JAMA Neurology</i> , 2020 , 77, 683-690	17.2	3641
282	Hirschsprung disease, associated syndromes and genetics: a review. <i>Journal of Medical Genetics</i> , 2008 , 45, 1-14	5.8	622
281	Clinical characteristics and risk factors associated with COVID-19 disease severity in patients with cancer in Wuhan, China: a multicentre, retrospective, cohort study. <i>Lancet Oncology</i> , 2020 , 21, 893-903	21.7	282
280	Skin damage among health care workers managing coronavirus disease-2019. <i>Journal of the American Academy of Dermatology</i> , 2020 , 82, 1215-1216	4.5	252
279	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2011 , 43, 679-84	36.3	233
278	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. <i>Nature Genetics</i> , 2011 , 43, 1215-8	36.3	215
277	Differential contributions of rare and common, coding and noncoding Ret mutations to multifactorial Hirschsprung disease liability. <i>American Journal of Human Genetics</i> , 2010 , 87, 60-74	11	198
276	Genome-wide association analyses of esophageal squamous cell carcinoma in Chinese identify multiple susceptibility loci and gene-environment interactions. <i>Nature Genetics</i> , 2012 , 44, 1090-7	36.3	196
275	Identification of genetic variants in base excision repair pathway and their associations with risk of esophageal squamous cell carcinoma. <i>Cancer Research</i> , 2004 , 64, 4378-84	10.1	196
274	Genetic polymorphisms of interleukin (IL)-1B, IL-1RN, IL-8, IL-10 and tumor necrosis factor {alpha} and risk of gastric cancer in a Chinese population. <i>Carcinogenesis</i> , 2005 , 26, 631-6	4.6	190
273	Pancreatic cancer risk variant in LINC00673 creates a miR-1231 binding site and interferes with PTPN11 degradation. <i>Nature Genetics</i> , 2016 , 48, 747-57	36.3	187
272	Identification of Functional Genetic Variants in Cyclooxygenase-2 and Their Association With Risk of Esophageal Cancer. <i>Gastroenterology</i> , 2005 , 129, 565-576	13.3	181
271	Polymorphisms of death pathway genes FAS and FASL in esophageal squamous-cell carcinoma. <i>Journal of the National Cancer Institute</i> , 2004 , 96, 1030-6	9.7	172
270	lncRNASNP: a database of SNPs in lncRNAs and their potential functions in human and mouse. <i>Nucleic Acids Research</i> , 2015 , 43, D181-6	20.1	165
269	Functional haplotypes in the promoter of matrix metalloproteinase-2 predict risk of the occurrence and metastasis of esophageal cancer. <i>Cancer Research</i> , 2004 , 64, 7622-8	10.1	162
268	Meta-analysis on night shift work and risk of metabolic syndrome. <i>Obesity Reviews</i> , 2014 , 15, 709-20	10.6	161
267	Cohort Profile: the Dongfeng-Tongji cohort study of retired workers. <i>International Journal of Epidemiology</i> , 2013 , 42, 731-40	7.8	155

266	The role of P53 and MDM2 polymorphisms in the risk of esophageal squamous cell carcinoma. <i>Cancer Research</i> , 2005 , 65, 9582-7	10.1	154
265	Genome-wide association study identifies NRG1 as a susceptibility locus for Hirschsprung@ disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 2694-9	11.5	146
264	Clinical Characteristics of and Medical Interventions for COVID-19 in Hemodialysis Patients in Wuhan, China. <i>Journal of the American Society of Nephrology: JASN</i> , 2020 , 31, 1387-1397	12.7	144
263	Genome-wide association study identifies five loci associated with susceptibility to pancreatic cancer in Chinese populations. <i>Nature Genetics</i> , 2011 , 44, 62-6	36.3	141
262	Genetic polymorphisms in cell cycle regulatory genes MDM2 and TP53 are associated with susceptibility to lung cancer. <i>Human Mutation</i> , 2006 , 27, 110-7	4.7	126
261	Aurora-A/STK15 T+91A is a general low penetrance cancer susceptibility gene: a meta-analysis of multiple cancer types. <i>Carcinogenesis</i> , 2005 , 26, 1368-73	4.6	124
260	Polymorphisms in DNA base excision repair genes ADPRT and XRCC1 and risk of lung cancer. <i>Cancer Research</i> , 2005 , 65, 722-6	10.1	124
259	Clinical characteristics of 54 medical staff with COVID-19: A retrospective study in a single center in Wuhan, China. <i>Journal of Medical Virology</i> , 2020 , 92, 807-813	19.7	114
258	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2014 , 46, 1001-1006	36.3	112
257	Functional polymorphisms in cell death pathway genes FAS and FASL contribute to risk of lung cancer. <i>Journal of Medical Genetics</i> , 2005 , 42, 479-84	5.8	110
256	Polymorphisms of DNA repair genes XRCC1 and XPD and their associations with risk of esophageal squamous cell carcinoma in a Chinese population. <i>International Journal of Cancer</i> , 2002 , 100, 600-5	7.5	109
255	A novel T-77C polymorphism in DNA repair gene XRCC1 contributes to diminished promoter activity and increased risk of non-small cell lung cancer. <i>Oncogene</i> , 2006 , 25, 3613-20	9.2	103
254	PancanQTL: systematic identification of cis-eQTLs and trans-eQTLs in 33 cancer types. <i>Nucleic Acids Research</i> , 2018 , 46, D971-D976	20.1	100
253	FASL -844C polymorphism is associated with increased activation-induced T cell death and risk of cervical cancer. <i>Journal of Experimental Medicine</i> , 2005 , 202, 967-74	16.6	95
252	Identification of functional genetic variants in cyclooxygenase-2 and their association with risk of esophageal cancer. <i>Gastroenterology</i> , 2005 , 129, 565-76	13.3	94
251	Sequence variations in the DNA repair gene XPD and risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2003 , 105, 669-73	7.5	93
250	Genome-wide association study identifies a susceptibility locus for biliary atresia on 10q24.2. <i>Human Molecular Genetics</i> , 2010 , 19, 2917-25	5.6	90
249	Substantial reduction in risk of breast cancer associated with genetic polymorphisms in the promoters of the matrix metalloproteinase-2 and tissue inhibitor of metalloproteinase-2 genes. <i>Carcinogenesis</i> , 2004 , 25, 399-404	4.6	87

248	The expression levels of plasma microRNAs in atrial fibrillation patients. <i>PLoS ONE</i> , 2012 , 7, e44906	3.7	81
247	A genome-wide association study identifies two new cervical cancer susceptibility loci at 4q12 and 17q12. <i>Nature Genetics</i> , 2013 , 45, 918-22	36.3	80
246	DNA repair gene XPC genotypes/haplotypes and risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2005 , 115, 478-83	7.5	80
245	Functional variants in cell death pathway genes and risk of pancreatic cancer. <i>Clinical Cancer Research</i> , 2008 , 14, 3230-6	12.9	79
244	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, 6616-33	5.6	77
243	Functional STK15 Phe31Ile polymorphism is associated with the occurrence and advanced disease status of esophageal squamous cell carcinoma. <i>Cancer Research</i> , 2004 , 64, 2680-3	10.1	76
242	Urinary levels of bisphenol A, F and S and markers of oxidative stress among healthy adult men: Variability and association analysis. <i>Environment International</i> , 2019 , 123, 301-309	12.9	72
241	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. <i>Human Molecular Genetics</i> , 2015 , 24, 1791-800	5.6	71
240	ELABELA and an ELABELA Fragment Protect against AKI. <i>Journal of the American Society of Nephrology: JASN</i> , 2017 , 28, 2694-2707	12.7	69
239	Functional haplotypes in the promoter of matrix metalloproteinase-2 and lung cancer susceptibility. <i>Carcinogenesis</i> , 2005 , 26, 1117-21	4.6	69
238	Identification of a HOXD13 mutation in a VACTERL patient. <i>American Journal of Medical Genetics, Part A</i> , 2008 , 146A, 3181-5	2.5	68
237	Genetic variations in the TGF β signaling pathway, smoking and risk of colorectal cancer in a Chinese population. <i>Carcinogenesis</i> , 2013 , 34, 936-42	4.6	67
236	A common polymorphism in the 3'UTR of cyclooxygenase 2/prostaglandin synthase 2 gene and risk of lung cancer in a Chinese population. <i>Lung Cancer</i> , 2005 , 48, 11-7	5.9	66
235	A functional polymorphism in the matrix metalloproteinase-2 gene promoter (-1306C/T) is associated with risk of development but not metastasis of gastric cardia adenocarcinoma. <i>Cancer Research</i> , 2003 , 63, 3987-90	10.1	66
234	Interaction of P53 Arg72Pro and MDM2 T309G polymorphisms and their associations with risk of gastric cardia cancer. <i>Carcinogenesis</i> , 2007 , 28, 1996-2001	4.6	64
233	Reelin gene variants and risk of autism spectrum disorders: an integrated meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014 , 165B, 192-200	3.5	63
232	Exome-wide analyses identify low-frequency variant in CYP26B1 and additional coding variants associated with esophageal squamous cell carcinoma. <i>Nature Genetics</i> , 2018 , 50, 338-343	36.3	62
231	A functional polymorphism in lnc-LAMC2-1:1 confers risk of colorectal cancer by affecting miRNA binding. <i>Carcinogenesis</i> , 2016 , 37, 443-51	4.6	60

230	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, 928-34	11	59
229	Dietary legume consumption reduces risk of colorectal cancer: evidence from a meta-analysis of cohort studies. <i>Scientific Reports</i> , 2015 , 5, 8797	4.9	58
228	Genetic variants in the folate pathway and the risk of neural tube defects: a meta-analysis of the published literature. <i>PLoS ONE</i> , 2013 , 8, e59570	3.7	57
227	Evaluation of genetic susceptibility loci for chronic hepatitis B in Chinese: two independent case-control studies. <i>PLoS ONE</i> , 2011 , 6, e17608	3.7	55
226	A functional polymorphism located at transcription factor binding sites, rs6695837 near LAMC1 gene, confers risk of colorectal cancer in Chinese populations. <i>Carcinogenesis</i> , 2017 , 38, 177-183	4.6	54
225	Non-linear dose-response relationship between cigarette smoking and pancreatic cancer risk: evidence from a meta-analysis of 42 observational studies. <i>European Journal of Cancer</i> , 2014 , 50, 193-203	7.5	53
224	Common genetic variants regulating ADD3 gene expression alter biliary atresia risk. <i>Journal of Hepatology</i> , 2013 , 59, 1285-91	13.4	53
223	Adenosine diphosphate ribosyl transferase and x-ray repair cross-complementing 1 polymorphisms in gastric cardia cancer. <i>Gastroenterology</i> , 2006 , 131, 420-7	13.3	52
222	Functional Phe31Ile polymorphism in Aurora A and risk of breast carcinoma. <i>Carcinogenesis</i> , 2004 , 25, 2225-30	4.6	52
221	Potential Influence of Menstrual Status and Sex Hormones on Female Severe Acute Respiratory Syndrome Coronavirus 2 Infection: A Cross-sectional Multicenter Study in Wuhan, China. <i>Clinical Infectious Diseases</i> , 2021 , 72, e240-e248	11.6	52
220	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , 2017 , 66, 581-587	19.2	51
219	Exposure to the Chinese Famine in Childhood Increases Type 2 Diabetes Risk in Adults. <i>Journal of Nutrition</i> , 2016 , 146, 2289-2295	4.1	51
218	A genome wide association study of genetic loci that influence tumour biomarkers cancer antigen 19-9, carcinoembryonic antigen and fletoprotein and their associations with cancer risk. <i>Gut</i> , 2014 , 63, 143-51	19.2	51
217	Polymorphisms of XRCC1 genes and risk of nasopharyngeal carcinoma in the Cantonese population. <i>BMC Cancer</i> , 2006 , 6, 167	4.8	51
216	Association between serum uric acid and the metabolic syndrome among a middle- and old-age Chinese population. <i>European Journal of Epidemiology</i> , 2013 , 28, 669-76	12.1	50
215	Genome-wide interrogation identifies YAP1 variants associated with survival of small-cell lung cancer patients. <i>Cancer Research</i> , 2010 , 70, 9721-9	10.1	49
214	Inactivation of DNA repair gene O6-methylguanine-DNA methyltransferase by promoter hypermethylation and its relation to p53 mutations in esophageal squamous cell carcinoma. <i>Carcinogenesis</i> , 2003 , 24, 1039-44	4.6	49
213	A genome wide association study identifies common variants associated with lipid levels in the Chinese population. <i>PLoS ONE</i> , 2013 , 8, e82420	3.7	48

212	Longer Sleep Duration and Midday Napping Are Associated with a Higher Risk of CHD Incidence in Middle-Aged and Older Chinese: the Dongfeng-Tongji Cohort Study. <i>Sleep</i> , 2016 , 39, 645-52	1.1	47
211	A Rare Missense Variant in TCF7L2 Associates with Colorectal Cancer Risk by Interacting with a GWAS-Identified Regulatory Variant in the MYC Enhancer. <i>Cancer Research</i> , 2018 , 78, 5164-5172	10.1	46
210	The effect of laparoscopic excision vs open excision in children with choledochal cyst: a midterm follow-up study. <i>Journal of Pediatric Surgery</i> , 2011 , 46, 662-665	2.6	46
209	Genome-wide copy number analysis uncovers a new HSCR gene: NRG3. <i>PLoS Genetics</i> , 2012 , 8, e10026876		46
208	Mutations in the NRG1 gene are associated with Hirschsprung disease. <i>Human Genetics</i> , 2012 , 131, 67-766.3		45
207	Reduced RET expression in gut tissue of individuals carrying risk alleles of Hirschsprung disease. <i>Human Molecular Genetics</i> , 2010 , 19, 1461-7	5.6	45
206	Genetic polymorphisms in methylenetetrahydrofolate reductase and thymidylate synthase and risk of pancreatic cancer. <i>Clinical Gastroenterology and Hepatology</i> , 2005 , 3, 743-51	6.9	45
205	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016 , 7, 11478	17.4	45
204	Implications of endocrine gland-derived vascular endothelial growth factor/prokineticin-1 signaling in human neuroblastoma progression. <i>Clinical Cancer Research</i> , 2007 , 13, 868-75	12.9	44
203	Genetic polymorphism in myeloperoxidase but not GSTM1 is associated with risk of lung squamous cell carcinoma in a Chinese population. <i>International Journal of Cancer</i> , 2002 , 102, 275-9	7.5	44
202	A low-frequency variant in SMAD7 modulates TGF- β signaling and confers risk for colorectal cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2017 , 56, 1798-1807	5	43
201	A functional polymorphism in the SULT1A1 gene (G638A) is associated with risk of lung cancer in relation to tobacco smoking. <i>Carcinogenesis</i> , 2004 , 25, 773-8	4.6	43
200	Genetic variations in TERT-CLPTM1L locus are associated with risk of lung cancer in Chinese population. <i>Molecular Carcinogenesis</i> , 2013 , 52 Suppl 1, E118-26	5	42
199	Association of candidate genetic variations with gastric cardia adenocarcinoma in Chinese population: a multiple interaction analysis. <i>Carcinogenesis</i> , 2011 , 32, 336-42	4.6	42
198	Dinucleotide polymorphism of p73 gene is associated with a reduced risk of lung cancer in a Chinese population. <i>International Journal of Cancer</i> , 2005 , 114, 455-60	7.5	42
197	Severe acute respiratory syndrome coronavirus 2 detection in the female lower genital tract. <i>American Journal of Obstetrics and Gynecology</i> , 2020 , 223, 131-134	6.4	41
196	A polymorphic MYC response element in KBTBD11 influences colorectal cancer risk, especially in interaction with an MYC-regulated SNP rs6983267. <i>Annals of Oncology</i> , 2018 , 29, 632-639	10.3	41
195	Allium vegetables and garlic supplements do not reduce risk of colorectal cancer, based on meta-analysis of prospective studies. <i>Clinical Gastroenterology and Hepatology</i> , 2014 , 12, 1991-2001.e1-4; quiz e121	6.9	41

194	Integrative expression quantitative trait locus-based analysis of colorectal cancer identified a functional polymorphism regulating SLC22A5 expression. <i>European Journal of Cancer</i> , 2018 , 93, 1-9	7.5	40
193	Perturbation of hoxb5 signaling in vagal neural crests down-regulates ret leading to intestinal hypoganglionosis in mice. <i>Gastroenterology</i> , 2008 , 134, 1104-15	13.3	40
192	Glyceraldehyde-3-phosphate dehydrogenase promotes liver tumorigenesis by modulating phosphoglycerate dehydrogenase. <i>Hepatology</i> , 2017 , 66, 631-645	11.2	39
191	Prevalence and associated risk factors of dyslexic children in a middle-sized city of China: a cross-sectional study. <i>PLoS ONE</i> , 2013 , 8, e56688	3.7	39
190	Interactions between genetic variants in the adiponectin, adiponectin receptor 1 and environmental factors on the risk of colorectal cancer. <i>PLoS ONE</i> , 2011 , 6, e27301	3.7	39
189	Environmental exposure to metals and the risk of hypertension: A cross-sectional study in China. <i>Environmental Pollution</i> , 2018 , 233, 670-678	9.3	38
188	Identification of Genes Associated With Hirschsprung Disease, Based on Whole-Genome Sequence Analysis, and Potential Effects on Enteric Nervous System Development. <i>Gastroenterology</i> , 2018 , 155, 1908-1922.e5	13.3	38
187	Combined effect of genetic polymorphisms in P53, P73, and MDM2 on non-small cell lung cancer survival. <i>Journal of Thoracic Oncology</i> , 2011 , 6, 1793-800	8.9	36
186	Association between polymorphisms of DRD2 and DRD4 and opioid dependence: evidence from the current studies. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011 , 156B, 661-70	3.5	36
185	Bidirectional association between nonalcoholic fatty liver disease and type 2 diabetes in Chinese population: Evidence from the Dongfeng-Tongji cohort study. <i>PLoS ONE</i> , 2017 , 12, e0174291	3.7	36
184	A Rare Variant P507L in TPP1 Interrupts TPP1-TIN2 Interaction, Influences Telomere Length, and Confers Colorectal Cancer Risk in Chinese Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 1029-1035	4	34
183	The SLC4A7 variant rs4973768 is associated with breast cancer risk: evidence from a case-control study and a meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2012 , 136, 847-57	4.4	34
182	Platelet 12-lipoxygenase Arg261Gln polymorphism: functional characterization and association with risk of esophageal squamous cell carcinoma in combination with COX-2 polymorphisms. <i>Pharmacogenetics and Genomics</i> , 2007 , 17, 197-205	1.9	33
181	Significant increase in risk of gastroesophageal cancer is associated with interaction between promoter polymorphisms in thymidylate synthase and serum folate status. <i>Carcinogenesis</i> , 2005 , 26, 1430-5	4.6	33
180	Loss of myeloid-related proteins 8 and myeloid-related proteins 14 expression in human esophageal squamous cell carcinoma correlates with poor differentiation. <i>World Journal of Gastroenterology</i> , 2004 , 10, 1093-7	5.6	33
179	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017 , 152, 2011-2021	13.3	32
178	A functional polymorphism (-1607 1G->2G) in the matrix metalloproteinase-1 promoter is associated with development and progression of lung cancer. <i>Cancer</i> , 2011 , 117, 5172-81	6.4	32
177	AWESOME: a database of SNPs that affect protein post-translational modifications. <i>Nucleic Acids Research</i> , 2019 , 47, D874-D880	20.1	32

176	Susceptibility to gastric cardia adenocarcinoma and genetic polymorphisms in methylenetetrahydrofolate reductase in an at-risk Chinese population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002 , 11, 1454-8	4	32
175	Exposure to the Chinese famine in early life and hypertension prevalence risk in adults. <i>Journal of Hypertension</i> , 2017 , 35, 63-68	1.9	31
174	Potential of Arbidol for Post-exposure Prophylaxis of COVID-19 Transmission: A Preliminary Report of a Retrospective Cohort Study. <i>Current Medical Science</i> , 2020 , 40, 480-485	2.8	31
173	Dietary mushroom intake may reduce the risk of breast cancer: evidence from a meta-analysis of observational studies. <i>PLoS ONE</i> , 2014 , 9, e93437	3.7	31
172	Genetic variant in KIAA0319, but not in DYX1C1, is associated with risk of dyslexia: an integrated meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012 , 159B, 970-6	3.5	31
171	A functional -77T>C polymorphism in XRCC1 is associated with risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011 , 125, 479-87	4.4	31
170	MNX1 (HLXB9) mutations in Currarino patients. <i>Journal of Pediatric Surgery</i> , 2009 , 44, 1892-8	2.6	31
169	Liver intestine-cadherin (CDH17) haplotype is associated with increased risk of hepatocellular carcinoma. <i>Clinical Cancer Research</i> , 2006 , 12, 5248-52	12.9	31
168	Green tea consumption is associated with reduced incident CHD and improved CHD-related biomarkers in the Dongfeng-Tongji cohort. <i>Scientific Reports</i> , 2016 , 6, 24353	4.9	30
167	Assessment of XPD Lys751Gln and XRCC1 T-77C polymorphisms in advanced non-small-cell lung cancer patients treated with platinum-based chemotherapy. <i>Lung Cancer</i> , 2011 , 73, 110-5	5.9	30
166	Common genetic polymorphisms in the 5'flanking region of the SULT1A1 gene: haplotypes and their association with platelet enzymatic activity. <i>Pharmacogenetics and Genomics</i> , 2005 , 15, 465-73	1.9	30
165	Genetic variants in the SWI/SNF complex and smoking collaborate to modify the risk of pancreatic cancer in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015 , 54, 761-8	5	28
164	Different physical activity subtypes and risk of metabolic syndrome in middle-aged and older Chinese people. <i>PLoS ONE</i> , 2013 , 8, e53258	3.7	28
163	Risk prediction of esophageal squamous-cell carcinoma with common genetic variants and lifestyle factors in Chinese population. <i>Carcinogenesis</i> , 2013 , 34, 1782-6	4.6	27
162	Functional polymorphisms in FAS/FASL system increase the risk of neuroblastoma in Chinese population. <i>PLoS ONE</i> , 2013 , 8, e71656	3.7	27
161	Functional evaluation of missense variations in the human MAD1L1 and MAD2L1 genes and their impact on susceptibility to lung cancer. <i>Journal of Medical Genetics</i> , 2010 , 47, 616-22	5.8	27
160	Fine mapping of the NRG1 Hirschsprung@ disease locus. <i>PLoS ONE</i> , 2011 , 6, e16181	3.7	27
159	Systematic Functional Interrogation of Genes in GWAS Loci Identified ATF1 as a Key Driver in Colorectal Cancer Modulated by a Promoter-Enhancer Interaction. <i>American Journal of Human Genetics</i> , 2019 , 105, 29-47	11	26

158	A solute carrier family 22 member 3 variant rs3088442 G->A associated with coronary heart disease inhibits lipopolysaccharide-induced inflammatory response. <i>Journal of Biological Chemistry</i> , 2015 , 290, 5328-40	5.4	26
157	<i>Helicobacter pylori</i> infection is associated with type 2 diabetes among a middle- and old-age Chinese population. <i>Diabetes/Metabolism Research and Reviews</i> , 2016 , 32, 95-101	7.5	26
156	A genetic variant rs1801274 in FCGR2A as a potential risk marker for Kawasaki disease: a case-control study and meta-analysis. <i>PLoS ONE</i> , 2014 , 9, e103329	3.7	26
155	GSK-3 β polymorphism discriminates bipolar disorder and schizophrenia: a systematic meta-analysis. <i>Molecular Neurobiology</i> , 2013 , 48, 404-11	6.2	26
154	Sleep Duration and Midday Napping with 5-Year Incidence and Reversion of Metabolic Syndrome in Middle-Aged and Older Chinese. <i>Sleep</i> , 2016 , 39, 1911-1918	1.1	26
153	Association between bilirubin and risk of Non-Alcoholic Fatty Liver Disease based on a prospective cohort study. <i>Scientific Reports</i> , 2016 , 6, 31006	4.9	25
152	Exome sequencing identified NRG3 as a novel susceptible gene of Hirschsprung's disease in a Chinese population. <i>Molecular Neurobiology</i> , 2013 , 47, 957-66	6.2	25
151	Polymorphisms of death pathway genes FAS and FASL and risk of nasopharyngeal carcinoma. <i>Molecular Carcinogenesis</i> , 2010 , 49, 944-50	5	25
150	Correlation between genetic variations in Hox clusters and Hirschsprung's disease. <i>Annals of Human Genetics</i> , 2007 , 71, 526-36	2.2	25
149	Pancan-meQTL: a database to systematically evaluate the effects of genetic variants on methylation in human cancer. <i>Nucleic Acids Research</i> , 2019 , 47, D1066-D1072	20.1	25
148	CancerSplicingQTL: a database for genome-wide identification of splicing QTLs in human cancer. <i>Nucleic Acids Research</i> , 2019 , 47, D909-D916	20.1	25
147	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. <i>Nature Communications</i> , 2018 , 9, 3688	17.4	25
146	Epigenomic landscape of 5-hydroxymethylcytosine reveals its transcriptional regulation of lncRNAs in colorectal cancer. <i>British Journal of Cancer</i> , 2017 , 116, 658-668	8.7	24
145	CYP2C19 polymorphism and clinical outcomes among patients of different races treated with clopidogrel: A systematic review and meta-analysis. <i>Journal of Huazhong University of Science and Technology [Medical Sciences]</i> , 2015 , 35, 147-156		24
144	Reducing protein regulator of cytokinesis 1 as a prospective therapy for hepatocellular carcinoma. <i>Cell Death and Disease</i> , 2018 , 9, 534	9.8	24
143	The epidemiology and etiology of influenza-like illness in Chinese children from 2008 to 2010. <i>Journal of Medical Virology</i> , 2012 , 84, 672-8	19.7	24
142	Multi-loci analysis reveals the importance of genetic variations in sensitivity of platinum-based chemotherapy in non-small-cell lung cancer. <i>Molecular Carcinogenesis</i> , 2013 , 52, 923-31	5	24
141	Variations in HSPA1B at 6p21.3 are associated with lung cancer risk and prognosis in Chinese populations. <i>Cancer Research</i> , 2011 , 71, 7576-86	10.1	24

140	N-methyladenosine mRNA methylation of regulates AKT signalling to promote PTEN-deficient pancreatic cancer progression. <i>Gut</i> , 2020 , 69, 2180-2192	19.2	24
139	Systematic confirmation study of GWAS-identified genetic variants for Kawasaki disease in a Chinese population. <i>Scientific Reports</i> , 2015 , 5, 8194	4.9	23
138	Associations of environmental exposure to metals with the risk of hypertension in China. <i>Science of the Total Environment</i> , 2018 , 622-623, 184-191	10.2	23
137	Using different anthropometric indices to assess prediction ability of type 2 diabetes in elderly population: a 5-year prospective study. <i>BMC Geriatrics</i> , 2018 , 18, 218	4.1	23
136	New integrated strategy emphasizing infection source control to curb Schistosomiasis japonica in a marshland area of Hubei Province, China: findings from an eight-year longitudinal survey. <i>PLoS ONE</i> , 2014 , 9, e89779	3.7	22
135	Application of a hybrid model for predicting the incidence of tuberculosis in Hubei, China. <i>PLoS ONE</i> , 2013 , 8, e80969	3.7	22
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