List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	27.8	3,823
2	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	21.4	1,818
3	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	21.4	377
4	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. Gastroenterology, 2018, 154, 2152-2164.e19.	1.3	226
5	Association of Aspirin and NSAID Use With Risk of Colorectal Cancer According to Genetic Variants. JAMA - Journal of the American Medical Association, 2015, 313, 1133.	7.4	171
6	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	12.8	169
7	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
8	A Model to Determine Colorectal Cancer Risk Using Common Genetic Susceptibility Loci. Gastroenterology, 2015, 148, 1330-1339.e14.	1.3	129
9	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	6.3	129
10	Genetic determinants of telomere length and risk of common cancers: a Mendelian randomization study. Human Molecular Genetics, 2015, 24, 5356-5366.	2.9	128
11	Mendelian randomization study of adiposity-related traits and risk of breast, ovarian, prostate, lung and colorectal cancer. International Journal of Epidemiology, 2016, 45, 896-908.	1.9	124
12	Estimating the heritability of colorectal cancer. Human Molecular Genetics, 2014, 23, 3898-3905.	2.9	114
13	Trend and disease burden of bacillary dysentery in China (1991-2000). Bulletin of the World Health Organization, 2006, 84, 561-568.	3.3	107
14	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. Cancer Research, 2016, 76, 5103-5114.	0.9	100
15	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. Gastroenterology, 2016, 150, 1633-1645.	1.3	97
16	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. Molecular Psychiatry, 2020, 25, 2392-2409.	7.9	83
17	Genome-Wide Diet-Gene Interaction Analyses for Risk of Colorectal Cancer. PLoS Genetics, 2014, 10, e1004228.	3.5	81
18	Fine Mapping and Identification of BMI Loci in African Americans. American Journal of Human Genetics, 2013, 93, 661-671.	6.2	77

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19	Maternal ethnicity and preâ€eclampsia in New York City, 1995–2003. Paediatric and Perinatal Epidemiology, 2012, 26, 45-52.	1.7	76
20	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and AlcoholÂUse. Biological Psychiatry, 2019, 85, 946-955.	1.3	69
21	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1024-1031.	2.5	67
22	MicroRNA-1246 promotes growth and metastasis of colorectal cancer cells involving CCNG2 reduction. Molecular Medicine Reports, 2016, 13, 273-280.	2.4	65
23	Mendelian randomization study of height and risk of colorectal cancer. International Journal of Epidemiology, 2015, 44, 662-672.	1.9	55
24	A Pooled Analysis of Smoking and Colorectal Cancer: Timing of Exposure and Interactions with Environmental Factors. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1974-1985.	2.5	54
25	Gene–Environment Interaction Involving Recently Identified Colorectal Cancer Susceptibility Loci. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1824-1833.	2.5	48
26	Trends and disease burden of enteric fever in Guangxi province, China, 1994–2004. Bulletin of the World Health Organization, 2010, 88, 689-696.	3.3	40
27	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. PLoS Genetics, 2016, 12, e1006296.	3.5	38
28	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. Human Genetics, 2017, 136, 771-800.	3.8	31
29	Circular RNA RBPMS inhibits bladder cancer progression via miR-330-3p/RAI2 regulation. Molecular Therapy - Nucleic Acids, 2021, 23, 872-886.	5.1	30
30	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. Carcinogenesis, 2015, 36, 999-1007.	2.8	28
31	Combined effect of modifiable and non-modifiable risk factors for colorectal cancer risk in a pooled analysis of 11 population-based studies. BMJ Open Gastroenterology, 2019, 6, e000339.	2.7	28
32	Prostate health index significantly reduced unnecessary prostate biopsies in patients with PSA 2-10 ng/mL and PSA >10 ng/mL: Results from a Multicenter Study in China. Prostate, 2017, 77, 1221	-1 2 29.	26
33	F‑box proteins involved in cancer‑associated drug resistance (Review). Oncology Letters, 2018, 15, 8891-8900.	1.8	20
34	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. British Journal of Cancer, 2016, 114, 221-229.	6.4	18
35	Esophageal squamous cell carcinoma cell proliferation induced by exposure to low concentration of cigarette smoke extract is mediated via targeting miR-101-3p/COX-2 pathway. Oncology Reports, 2016, 35, 463-471.	2.6	16
36	Overexpression of Fâ€box only protein 31 predicts poor prognosis and deregulates p38î±â€•and JNKâ€mediated apoptosis in esophageal squamous cell carcinoma. International Journal of Cancer, 2018, 142, 145-155.	5.1	15

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37	Genome-Wide Association Study of Serum Selenium Concentrations. Nutrients, 2013, 5, 1706-1718.	4.1	14
38	Roles of F-box proteins in human digestive system tumors (Review). International Journal of Oncology, 2014, 45, 2199-2207.	3.3	12
39	Function and mechanism of F-box proteins in gastric cancer (Review). International Journal of Oncology, 2015, 47, 43-50.	3.3	12
40	Calcipotriol attenuates liver fibrosis through the inhibition of vitamin D receptor-mediated NF-κB signaling pathway. Bioengineered, 2022, 13, 2658-2672.	3.2	12
41	Race-specific genetic risk score is more accurate than nonrace-specific genetic risk score for predicting prostate cancer and high-grade diseases. Asian Journal of Andrology, 2016, 18, 525.	1.6	11
42	Research on optimal immunization strategies for hepatitis B in different endemic areas in China. World Journal of Gastroenterology, 2000, 6, 392.	3.3	11
43	Increased Expression of Lysine-Specific Demethylase 5B (KDM5B) Promotes Tumor Cell Growth in Hep3B Cells and is an Independent Prognostic Factor in Patients with Hepatocellular Carcinoma. Medical Science Monitor, 2018, 24, 7586-7594.	1.1	10
44	No Evidence of Gene–Calcium Interactions from Genome-Wide Analysis of Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2971-2976.	2.5	9
45	Epidemiologic Association between Inflammatory Bowel Diseases and Type 1 Diabetes Mellitus: a Meta-Analysis. Journal of Gastrointestinal and Liver Diseases, 2020, 29, 407-413.	0.9	9
46	The role of F‑box only protein 31 in cancer (Review). Oncology Letters, 2018, 15, 4047-4052.	1.8	7
47	CREB1 acts via the miR‑922/ARID2 axis to enhance malignant behavior of liver cancer cells. Oncology Reports, 2021, 45, .	2.6	7
48	Long-term effectiveness of infancy low-dose hepatitis B vaccine immunization in Zhuang minority area in China. World Journal of Gastroenterology, 1999, 5, 122.	3.3	6
49	Phi-based risk calculators performed better in the prediction of prostate cancer in the Chinese population. Asian Journal of Andrology, 2019, 21, 592.	1.6	5
50	Germline genetic variations in PDZD2 and ITPR2 genes are associated with clear cell renal cell carcinoma in Chinese population. Oncotarget, 2017, 8, 24196-24201.	1.8	5
51	Pocket-creation method improves efficacy of colorectal endoscopic submucosal dissection: a system review and meta-analysis. European Journal of Gastroenterology and Hepatology, 2021, 33, 1241-1246.	1.6	5
52	New insights into the mechanism of F-box proteins in colorectal cancer (Review). Oncology Reports, 2015, 33, 2113-2120.	2.6	4
53	Comprehensive analysis of abnormal expression, prognostic value and oncogenic role of the hub gene FN1 in pancreatic ductal adenocarcinoma <i>via</i> bioinformatic analysis and <i>in vitro</i> experiments. PeerJ, 2021, 9, e12141.	2.0	4
54	11β‑hydroxysteroid dehydrogenase‑1 is associated with the activation of hepatic stellate cells in the development of hepatic fibrosis. Molecular Medicine Reports, 2020, 22, 3191-3200.	2.4	4

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55	Involvement of F-box proteins in esophageal cancer (Review). International Journal of Oncology, 2016, 48, 886-894.	3.3	3
56	A Germline Variant at 8q24 Contributes to the Serum p2PSA Level in a Chinese Prostate Biopsy Cohort. Frontiers in Oncology, 2021, 11, 753920.	2.8	2
57	Commentary: Mendelian randomization, testosterone, and cardiovascular disease. International Journal of Epidemiology, 2015, 44, 621-622.	1.9	1