

# Gong-Hong Wei

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

46  
papers

2,809  
citations

20  
h-index

53  
g-index

54  
ext. papers

3,496  
ext. citations

10.9  
avg, IF

4.48  
L-index

#	Paper	IF	Citations
46	Human transcription factor protein interaction networks.. <i>Nature Communications</i> , <b>2022</b> , 13, 766	17.4	6
45	An enhancer variant at 16q22.1 predisposes to hepatocellular carcinoma via regulating PRMT7 expression.. <i>Nature Communications</i> , <b>2022</b> , 13, 1232	17.4	0
44	VHL Ser65 mutations enhance HIF2 $\beta$ signaling and promote epithelial-mesenchymal transition of renal cancer cells.. <i>Cell and Bioscience</i> , <b>2022</b> , 12, 52	9.8	0
43	Large Multicohort Study Reveals a Prostate Cancer Susceptibility Allele at 5p15 Regulating Androgen Signaling-Orchestrated Chromatin Binding of E2F1 and MYC. <i>Frontiers in Oncology</i> , <b>2021</b> , 11, 754206	5.3	
42	The Amino-Terminal Oligomerization Domain of Angiotensin-2 Affects Vascular Remodeling, Mammary Gland Tumor Growth, and Lung Metastasis in Mice. <i>Cancer Research</i> , <b>2021</b> , 81, 129-143	10.1	5
41	CRISPRi screens reveal a DNA methylation-mediated 3D genome dependent causal mechanism in prostate cancer. <i>Nature Communications</i> , <b>2021</b> , 12, 1781	17.4	6
40	Optimized CRISPR/Cas9-mediated single nucleotide mutation in adherent cancer cell lines. <i>STAR Protocols</i> , <b>2021</b> , 2, 100419	1.4	1
39	Clinical characteristics and risk factors of COVID-19 patients with chronic hepatitis B: a multi-center retrospective cohort study. <i>Frontiers of Medicine</i> , <b>2021</b> , 1	12	2
38	Mechanistic insights into genetic susceptibility to prostate cancer. <i>Cancer Letters</i> , <b>2021</b> , 522, 155-163	9.9	0
37	A long hypoxia-inducible factor 3 isoform 2 is a transcription activator that regulates erythropoietin. <i>Cellular and Molecular Life Sciences</i> , <b>2020</b> , 77, 3627-3642	10.3	20
36	Exome Sequencing Reveals a Phenotype Modifying Variant in ZNF528 in Primary Osteoporosis With a COL1A2 Deletion. <i>Journal of Bone and Mineral Research</i> , <b>2020</b> , 35, 2381-2392	6.3	2
35	Association between homocysteine, vitamin B , folic acid and erectile dysfunction: a cross-sectional study in China. <i>BMJ Open</i> , <b>2019</b> , 9, e023003	3	4
34	Illumination of cell cycle progression by multi-fluorescent sensing system. <i>Cell Cycle</i> , <b>2019</b> , 18, 1364-1378	4.7	1
33	The Role of HOX Transcription Factors in Cancer Predisposition and Progression. <i>Cancers</i> , <b>2019</b> , 11,	6.6	54
32	Enhancer Dysfunction in 3D Genome and Disease. <i>Cells</i> , <b>2019</b> , 8,	7.9	8
31	A Large-Scale, Exome-Wide Association Study of Han Chinese Women Identifies Three Novel Loci Predisposing to Breast Cancer. <i>Cancer Research</i> , <b>2018</b> , 78, 3087-3097	10.1	17
30	Whole-genome and Transcriptome Sequencing of Prostate Cancer Identify New Genetic Alterations Driving Disease Progression. <i>European Urology</i> , <b>2018</b> , 73, 322-339	10.2	71

29	Biology and Clinical Implications of the 19q13 Aggressive Prostate Cancer Susceptibility Locus. <i>Cell</i> , <b>2018</b> , 174, 576-589.e18	56.2	72
28	Oncogenic regulatory circuits driven by 19q13 rs11672691 underlies prostate cancer aggressiveness. <i>Molecular and Cellular Oncology</i> , <b>2018</b> , 5, e1516451	1.2	1
27	Synergistic Interaction of and Predisposes to Aggressive Prostate Cancer. <i>Clinical Cancer Research</i> , <b>2018</b> , 24, 6265-6276	12.9	10
26	ANO7 is associated with aggressive prostate cancer. <i>International Journal of Cancer</i> , <b>2018</b> , 143, 2479-2487	7.5	17
25	Whole exome sequencing in Finnish families identifies new candidate genes for osteoarthritis. <i>PLoS ONE</i> , <b>2018</b> , 13, e0203313	3.7	4
24	High-throughput screening of prostate cancer risk loci by single nucleotide polymorphisms sequencing. <i>Nature Communications</i> , <b>2018</b> , 9, 2022	17.4	36
23	Systematic identification of regulatory variants associated with cancer risk. <i>Genome Biology</i> , <b>2017</b> , 18, 194	18.3	44
22	Multi-factors including Inflammatory/Immune, Hormones, Tumor-related Proteins and Nutrition associated with Chronic Prostatitis NIH IIIa+b and IV based on FAMHES project. <i>Scientific Reports</i> , <b>2017</b> , 7, 9143	4.9	3
21	Ataxin-10 is involved in Golgi membrane dynamics. <i>Journal of Genetics and Genomics</i> , <b>2017</b> , 44, 549-552	4	1
20	Genetic association analysis of the RTK/ERK pathway with aggressive prostate cancer highlights the potential role of CCND2 in disease progression. <i>Scientific Reports</i> , <b>2017</b> , 7, 4538	4.9	13
19	Genomic Insight into the Role of lncRNA in Cancer Susceptibility. <i>International Journal of Molecular Sciences</i> , <b>2017</b> , 18,	6.3	52
18	SIRT1 deacetylates the cardiac transcription factor Nkx2.5 and inhibits its transcriptional activity. <i>Scientific Reports</i> , <b>2016</b> , 6, 36576	4.9	23
17	Structural basis for DNA recognition by STAT6. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2016</b> , 113, 13015-13020	11.5	26
16	Gene regulatory mechanisms underpinning prostate cancer susceptibility. <i>Nature Genetics</i> , <b>2016</b> , 48, 387-97	36.3	72
15	Meta-analysis of gene expression and integrin-associated signaling pathways in papillary renal cell carcinoma subtypes. <i>Oncotarget</i> , <b>2016</b> , 7, 84178-84189	3.3	3
14	Chromatin interactions and candidate genes at ten prostate cancer risk loci. <i>Scientific Reports</i> , <b>2016</b> , 6, 23202	4.9	30
13	Identification of several potential chromatin binding sites of HOXB7 and its downstream target genes in breast cancer. <i>International Journal of Cancer</i> , <b>2015</b> , 137, 2374-83	7.5	23
12	Systematic enrichment analysis of potentially functional regions for 103 prostate cancer risk-associated loci. <i>Prostate</i> , <b>2015</b> , 75, 1264-76	4.2	30

11	A prostate cancer susceptibility allele at 6q22 increases RFX6 expression by modulating HOXB13 chromatin binding. <i>Nature Genetics</i> , <b>2014</b> , 46, 126-35	36.3	142
10	TP53 supports basal-like differentiation of mammary epithelial cells by preventing translocation of deltaNp63 into nucleoli. <i>Scientific Reports</i> , <b>2014</b> , 4, 4663	4.9	7
9	DNA-binding specificities of human transcription factors. <i>Cell</i> , <b>2013</b> , 152, 327-39	56.2	763
8	Genome-wide analysis of ETS-family DNA-binding in vitro and in vivo. <i>EMBO Journal</i> , <b>2010</b> , 29, 2147-60	13	403
7	Multiplexed massively parallel SELEX for characterization of human transcription factor binding specificities. <i>Genome Research</i> , <b>2010</b> , 20, 861-73	9.7	292
6	The common colorectal cancer predisposition SNP rs6983267 at chromosome 8q24 confers potential to enhanced Wnt signaling. <i>Nature Genetics</i> , <b>2009</b> , 41, 885-90	36.3	422
5	SATB1 regulates beta-like globin genes through matrix related nuclear relocation of the cluster. <i>Biochemical and Biophysical Research Communications</i> , <b>2009</b> , 383, 11-5	3.4	9
4	Exploring cellular memory molecules marking competent and active transcriptions. <i>BMC Molecular Biology</i> , <b>2007</b> , 8, 31	4.5	23
3	Unravelling the world of cis-regulatory elements. <i>Medical and Biological Engineering and Computing</i> , <b>2007</b> , 45, 709-18	3.1	10
2	Mechanisms of human gamma-globin transcriptional induction by apicidin involves p38 signaling to chromatin. <i>Biochemical and Biophysical Research Communications</i> , <b>2007</b> , 363, 889-94	3.4	17
1	Charting gene regulatory networks: strategies, challenges and perspectives. <i>Biochemical Journal</i> , <b>2004</b> , 381, 1-12	3.8	61