

# Yi-Ping Fu

## List of Publications by Year in descending order

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23  
papers

1,621  
citations

430874

18  
h-index

713466

21  
g-index

23  
all docs

23  
docs citations

23  
times ranked

3081  
citing authors

#	ARTICLE	IF	CITATIONS
1	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 978-984.	21.4	493
2	Breast cancer risk associated with genotypic polymorphism of the nonhomologous end-joining genes: a multigenic study on cancer susceptibility. <i>Cancer Research</i> , 2003, 63, 2440-6.	0.9	132
3	Breast cancer risk associated with genotype polymorphism of the catechol estrogen-metabolizing genes: A multigenic study on cancer susceptibility. <i>International Journal of Cancer</i> , 2005, 113, 345-353.	5.1	109
4	Breast Cancer Risk and the DNA Double-Strand Break End-Joining Capacity of Nonhomologous End-Joining Genes Are Affected by BRCA1. <i>Cancer Research</i> , 2004, 64, 5013-5019.	0.9	108
5	Common Genetic Polymorphisms Modify the Effect of Smoking on Absolute Risk of Bladder Cancer. <i>Cancer Research</i> , 2013, 73, 2211-2220.	0.9	107
6	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011, 20, 4282-4289.	2.9	100
7	Pesticide exposure on southwestern Taiwanese with MnSOD and NQO1 polymorphisms is associated with increased risk of Parkinson's disease. <i>Clinica Chimica Acta</i> , 2007, 378, 136-141.	1.1	79
8	Common genetic variants in the <i>PSCA</i> gene influence gene expression and bladder cancer risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 4974-4979.	7.1	79
9	Mapping of the UGT1A locus identifies an uncommon coding variant that affects mRNA expression and protects from bladder cancer. <i>Human Molecular Genetics</i> , 2012, 21, 1918-1930.	2.9	71
10	Trans-ethnic Meta-analysis and Functional Annotation Illuminates the Genetic Architecture of Fasting Glucose and Insulin. <i>American Journal of Human Genetics</i> , 2016, 99, 56-75.	6.2	55
11	NOTCH2 in breast cancer: association of SNP rs11249433 with gene expression in ER-positive breast tumors without TP53 mutations. <i>Molecular Cancer</i> , 2010, 9, 113.	19.2	52
12	The clinical implications of MMP-11 and CK-20 expression in human breast cancer. <i>Clinica Chimica Acta</i> , 2010, 411, 234-241.	1.1	45
13	Polymorphism of cytosolic serine hydroxymethyltransferase, estrogen and breast cancer risk among Chinese women in Taiwan. <i>Breast Cancer Research and Treatment</i> , 2008, 111, 145-155.	2.5	41
14	Large-Scale Pathway-Based Analysis of Bladder Cancer Genome-Wide Association Data from Five Studies of European Background. <i>PLoS ONE</i> , 2012, 7, e29396.	2.5	36
15	Association of MTHFR, MTR, and MTRR polymorphisms with Parkinson's disease among ethnic Chinese in Taiwan. <i>Clinica Chimica Acta</i> , 2011, 412, 332-338.	1.1	32
16	Breast cancer risk associated with genotypic polymorphism of the genes involved in the estrogen-receptor-signaling pathway: a multigenic study on cancer susceptibility. <i>Journal of Biomedical Science</i> , 2006, 13, 419-432.	7.0	25
17	Genetic Variant as a Selection Marker for Anti-Prostate Stem Cell Antigen Immunotherapy of Bladder Cancer. <i>Journal of the National Cancer Institute</i> , 2013, 105, 69-73.	6.3	25
18	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. <i>Cancer Research</i> , 2014, 74, 5808-5818.	0.9	24

#	ARTICLE	IF	CITATIONS
19	Detection of bladder, breast and prostate cancer using serum and tissue miRNA profiling. Genome Biology, 2011, 12, .	9.6	3
20	Prostate stem cell antigen (PSCA) and risk of bladder cancer: linking genotypes to functional mechanisms. Genome Biology, 2011, 12, .	8.8	3
21	Functional exploration of CCNE1 splicing forms as a possible link to bladder cancer susceptibility. Genome Biology, 2011, 12, .	8.8	2
22	An unusual suspect: an uncommon human-specific synonymous coding variant within the UGT1A6 gene explains a GWAS signal and protects against bladder cancer. Genome Biology, 2011, 12, .	8.8	0
23	A novel functional variant in 8q24 is associated with regulation of prostate stem cell antigen (PSCA) gene expression and bladder cancer risk. Genome Biology, 2011, 12, .	8.8	0