Shenying Fang

List of Publications by Year
in descending order

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68 papers	3,865 citations	172457 29 h-index	128289 60 g-index
69 all docs	69 docs citations	69 times ranked	8189 citing authors

#	Article	IF	Citations
1	Association of body-mass index and outcomes in patients with metastatic melanoma treated with targeted therapy, immunotherapy, or chemotherapy: a retrospective, multicohort analysis. Lancet Oncology, The, 2018, 19, 310-322.	10.7	486
2	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. JAMA Oncology, 2017, 3, 636.	7.1	376
3	Genome-wide association study identifies three new melanoma susceptibility loci. Nature Genetics, 2011, 43, 1108-1113.	21.4	230
4	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. Nature Genetics, 2015, 47, 987-995.	21.4	218
5	Racial disparity and socioeconomic status in association with survival in older men with local/regional stage prostate carcinoma. Cancer, 2006, 106, 1276-1285.	4.1	212
6	Genome-wide association study identifies novel loci predisposing to cutaneous melanomaâ€. Human Molecular Genetics, 2011, 20, 5012-5023.	2.9	187
7	Impact of Diabetes Mellitus on Complications and Outcomes of Adjuvant Chemotherapy in Older Patients With Breast Cancer. Journal of Clinical Oncology, 2009, 27, 2170-2176.	1.6	181
8	Racial disparities and socioeconomic status in association with survival in a large population-based cohort of elderly patients with colon cancer. Cancer, 2007, 110, 660-669.	4.1	157
9	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. Nature Genetics, 2020, 52, 494-504.	21.4	138
10	Acute Myeloid Leukemia After Adjuvant Breast Cancer Therapy in Older Women: Understanding Risk. Journal of Clinical Oncology, 2007, 25, 3871-3876.	1.6	134
11	A variant in FTO shows association with melanoma risk not due to BMI. Nature Genetics, 2013, 45, 428-432.	21.4	111
12	Mammography Before Diagnosis Among Women Age 80 Years and Older With Breast Cancer. Journal of Clinical Oncology, 2008, 26, 2482-2488.	1.6	93
13	Impact of Treatment and Socioeconomic Status on Racial Disparities in Survival Among Older Women With Breast Cancer. American Journal of Clinical Oncology: Cancer Clinical Trials, 2008, 31, 125-132.	1.3	93
14	Ethnic variations in diagnosis, treatment, socioeconomic status, and survival in a large populationâ€based cohort of elderly patients with nonâ€Hodgkin lymphoma. Cancer, 2008, 113, 3231-3241.	4.1	83
15	C-Reactive Protein As a Marker of Melanoma Progression. Journal of Clinical Oncology, 2015, 33, 1389-1396.	1.6	71
16	Association of Vitamin D Levels With Outcome in Patients With Melanoma After Adjustment For C-Reactive Protein. Journal of Clinical Oncology, 2016, 34, 1741-1747.	1.6	64
17	Socioeconomic status and cervical cancer survival among older women: Findings from the SEERâ€“Medicare linked data cohorts. Gynecologic Oncology, 2006, 102, 278-284.	1.4	60
18	Completion of adjuvant radiation therapy among women with breast cancer. Cancer, 2008, 113, 22-29.	4.1	60

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19	Variation in modes of chemotherapy administration for breast carcinoma and association with hospitalization for chemotherapy-related toxicity. Cancer, 2005, 104, 913-924.	4.1	48
20	Genetic variants in Hippo pathway genesYAP1,TEAD1andTEAD4are associated with melanoma-specific survival. International Journal of Cancer, 2015, 137, 638-645.	5.1	48
21	Telomere structure and maintenance gene variants and risk of five cancer types. International Journal of Cancer, 2016, 139, 2655-2670.	5.1	43
22	Identification of a melanoma susceptibility locus and somatic mutation inTET2. Carcinogenesis, 2014, 35, 2097-2101.	2.8	41
23	Joint Effect of Multiple Common SNPs Predicts Melanoma Susceptibility. PLoS ONE, 2013, 8, e85642.	2.5	40
24	Association between Body Mass Index, C-Reactive Protein Levels, and Melanoma Patient Outcomes. Journal of Investigative Dermatology, 2017, 137, 1792-1795.	0.7	40
25	Effects of MDM2, MDM4 and TP53 Codon 72 Polymorphisms on Cancer Risk in a Cohort Study of Carriers of TP53 Germline Mutations. PLoS ONE, 2010, 5, e10813.	2.5	37
26	Elderly patients with non-Hodgkin lymphoma who receive chemotherapy are at higher risk for osteoporosis and fractures. Leukemia and Lymphoma, 2007, 48, 1514-1521.	1.3	35
27	Longâ€term survival after radical prostatectomy compared to other treatments in older men with local/regional prostate cancer. Journal of Surgical Oncology, 2008, 97, 583-591.	1.7	32
28	Temporal and Geographic Variation in the Use of Hematopoietic Growth Factors in Older Women Receiving Breast Cancer Chemotherapy: Findings From a Large Population-Based Cohort. Journal of Clinical Oncology, 2005, 23, 8620-8628.	1.6	31
29	Radiation Use and Long-Term Survival in Breast Cancer Patients With T1, T2 Primary Tumors and One to Three Positive Axillary Lymph Nodes. International Journal of Radiation Oncology Biology Physics, 2008, 71, 1022-1027.	0.8	29
30	Mitochondrial DNA Copy Number in Peripheral Blood and Melanoma Risk. PLoS ONE, 2015, 10, e0131649.	2.5	29
31	Genetic Variants in Fanconi Anemia Pathway Genes BRCA2 and FANCA Predict Melanoma Survival. Journal of Investigative Dermatology, 2015, 135, 542-550.	0.7	28
32	Psoriasis prediction from genome-wide SNP profiles. BMC Dermatology, 2011, 11, 1.	2.1	27
33	Risk factors shared by COPD and lung cancer and mediation effect of COPD: two center caseâ€“control studies. Cancer Causes and Control, 2015, 26, 11-24.	1.8	26
34	Use of Intravenous Bisphosphonates in Older Women with Breast Cancer. Oncologist, 2008, 13, 494-502.	3.7	25
35	Role of Immune Response, Inflammation, and Tumor Immune Responseâ€“Related Cytokines/Chemokines in Melanoma Progression. Journal of Investigative Dermatology, 2019, 139, 2352-2358.e3.	0.7	23
36	Genetic variants in the PIWIâ€piRNA pathway gene DCP1A predict melanoma diseaseâ€specific survival. International Journal of Cancer, 2016, 139, 2730-2737.	5.1	21

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37	Functional Variants in Notch Pathway Genes NCOR2, NCSTN, and MAML2 Predict Survival of Patients with Cutaneous Melanoma. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1101-1110.	2.5	20
38	Genetic variants in the vitamin <scp>D</scp> pathway genes <scp>VDBP</scp>Âand <scp>RXRA</scp> modulate cutaneous melanoma diseaseâ€specific survival. Pigment Cell and Melanoma Research, 2016, 29, 176-185.	3.3	19
39	On the Interplay of Telomeres, Nevi and the Risk of Melanoma. PLoS ONE, 2012, 7, e52466.	2.5	18
40	Gene Variants in Angiogenesis and Lymphangiogenesis and Cutaneous Melanoma Progression. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 827-834.	2.5	17
41	Chemotherapy and Survival for Patients With Multiple Myeloma. American Journal of Clinical Oncology: Cancer Clinical Trials, 2007, 30, 540-548.	1.3	16
42	Complications associated with erythropoietin-stimulating agents in patients with metastatic breast cancer. Cancer, 2011, 117, 3641-3649.	4.1	16
43	Integrated pathway and epistasis analysis reveals interactive effect of genetic variants at <scp>TERF1</scp> and <scp>AFAP1L2</scp> loci on melanoma risk. International Journal of Cancer, 2015, 137, 1901-1909.	5.1	16
44	Functional annotation of melanoma risk loci identifies novel susceptibility genes. Carcinogenesis, 2020, 41, 452-457.	2.8	15
45	Variations in Chemotherapy and Radiation Therapy in a Large Nationwide and Community-Based Cohort of Elderly Patients With Non-Hodgkin Lymphoma. American Journal of Clinical Oncology: Cancer Clinical Trials, 2007, 30, 163-171.	1.3	13
46	Genetic variants in RORA and DNMT1 associated with cutaneous melanoma survival. International Journal of Cancer, 2018, 142, 2303-2312.	5.1	13
47	Global methylation of blood leukocyte DNA and risk of melanoma. International Journal of Cancer, 2017, 140, 1503-1509.	5.1	12
48	Genetic variants in ELOVL2 and HSD17B12 predict melanomaâ€specific survival. International Journal of Cancer, 2019, 145, 2619-2628.	5.1	11
49	Ordered Subset Analysis Identifies Loci Influencing Lung Cancer Risk on Chromosomes 6q and 12q. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 3157-3166.	2.5	10
50	Sex-specific effect of the TP53 PIN3 polymorphism on cancer risk in a cohort study of TP53 germline mutation carriers. Human Genetics, 2011, 130, 789-794.	3.8	10
51	Natural and orthogonal model for estimating geneâ€“gene interactions applied to cutaneous melanoma. Human Genetics, 2014, 133, 559-574.	3.8	10
52	Conditional Generative Adversarial Networks for Individualized Treatment Effect Estimation and Treatment Selection. Frontiers in Genetics, 2020, 11, 585804.	2.3	9
53	A comprehensive genomeâ€wide analysis of melanoma Breslow thickness identifies interaction between CDC42 and SCIN genetic variants. International Journal of Cancer, 2016, 139, 2012-2020.	5.1	8
54	Genetic variants in the genes encoding rho GTPases and related regulators predict cutaneous melanomaâ€specific survival. International Journal of Cancer, 2017, 141, 721-730.	5.1	8

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55	Evaluation of Plasma IL-6 in Patients with Melanoma as a Prognostic and Checkpoint Immunotherapy Predictive Biomarker. Journal of Investigative Dermatology, 2022, 142, 2046-2049.e3.	0.7	8
56	Association of Common Genetic Polymorphisms with Melanoma Patient IL-12p40 Blood Levels, Risk, and Outcomes. Journal of Investigative Dermatology, 2015, 135, 2266-2272.	0.7	7
57	Mitochondrial DNA 4977â€base pair common deletion in blood leukocytes and melanoma risk. Pigment Cell and Melanoma Research, 2016, 29, 372-378.	3.3	7
58	No prognostic value added by vitamin D pathway SNPs to current prognostic system for melanoma survival. PLoS ONE, 2017, 12, e0174234.	2.5	7
59	Genetic variants in the calcium signaling pathway genes are associated with cutaneous melanoma-specific survival. Carcinogenesis, 2019, 40, 279-288.	2.8	6
60	The relationship between blood <scp>IL</scp>â€12p40 level and melanoma progression. International Journal of Cancer, 2015, 136, 1874-1880.	5.1	5
61	Genetic Variants in WNT2B and BTRC Predict Melanoma Survival. Journal of Investigative Dermatology, 2017, 137, 1749-1756.	0.7	5
62	Genetic variants in the metzincin metallopeptidase family genes predict melanoma survival. Molecular Carcinogenesis, 2018, 57, 22-31.	2.7	5
63	Characterization of novel neutralizing mouse monoclonal antibody JM1-24-3 developed against MUC18 in metastatic melanoma. Journal of Experimental and Clinical Cancer Research, 2020, 39, 273.	8.6	5
64	Genetic variants in the integrin signaling pathway genes predict cutaneous melanoma survival. International Journal of Cancer, 2017, 140, 1270-1279.	5.1	4
65	Genetic variants of PDGF signaling pathway genes predict cutaneous melanoma survival. Oncotarget, 2017, 8, 74595-74606.	1.8	3
66	Building a Statistical Model for Predicting Cancer Genes. PLoS ONE, 2012, 7, e49175.	2.5	2
67	Melanoma Expression Genes Identified through Genome-Wide Association Study ofÂBreslow Tumor Thickness. Journal of Investigative Dermatology, 2017, 137, 253-257.	0.7	2
68	Reply to Z. Li et al. Journal of Clinical Oncology, 2015, 33, 3674-3675.	1.6	1