Dongyun Yang

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.

198
papers
6,277
citations
42
p-index
73
g-index

7,140
ext. papers
ext. citations
4.7
avg, IF
L-index

#	Paper	IF	Citations
198	High prevalence and inferior long-term outcomes for TP53 mutations in therapy-related acute lymphoblastic leukemia <i>American Journal of Hematology</i> , 2022 ,	7.1	O
197	A Randomized Open Label Pilot Study of Clostridium Butyricum Miyairi 588 (CBM588) in Recipients of Allogeneic Hematopoietic Cell Transplantation. <i>Blood</i> , 2021 , 138, 334-334	2.2	
196	Cytokine Release Syndrome Following Peripheral Blood Stem Cell Haploidentical HCT with Post-Transplant Cyclophosphamide: CRS post haploidentical HCT with PTCy. <i>Transplantation and Cellular Therapy</i> , 2021 ,		2
195	Donor-specific HLA antibodies associate with chronic graft-versus-host disease in haploidentical hematopoietic stem cell transplantation with post-transplant cyclophosphamide. <i>Bone Marrow Transplantation</i> , 2021 ,	4.4	
194	Allogeneic Hematopoietic Cell Transplantation for Relapsed and Refractory Philadelphia Negative B Cell ALL in the Era of Novel Salvage Therapies. <i>Transplantation and Cellular Therapy</i> , 2021 , 27, 255.e ⁻⁷	1-255.e	9 ¹
193	Cytokine gene polymorphisms are associated with response to blinatumomab in B-cell acute lymphoblastic leukemia. <i>European Journal of Haematology</i> , 2021 , 106, 851-858	3.8	1
192	Random survival forests identify pathways with polymorphisms predictive of survival in KRAS mutant and KRAS wild-type metastatic colorectal cancer patients. <i>Scientific Reports</i> , 2021 , 11, 12191	4.9	O
191	Long-Term Follow-up of High-Dose Chemotherapy with Autologous Stem Cell Transplantation in Children and Young Adults with Metastatic or Relapsed Ewing Sarcoma: A Single-Institution Experience. <i>Transplantation and Cellular Therapy</i> , 2021 , 27, 72.e1-72.e7		2
190	Late and very late relapsed acute lymphoblastic leukemia: clinical and molecular features, and treatment outcomes. <i>Blood Cancer Journal</i> , 2021 , 11, 125	7	O
189	Long-term follow-up of patients with poor-risk acute leukemia treated on a phase 2 trial undergoing intensified conditioning regimen prior to allogeneic hematopoietic cell transplantation. <i>Leukemia and Lymphoma</i> , 2021 , 1-7	1.9	
188	Comparing transplant outcomes in ALL patients after haploidentical with PTCy or matched unrelated donor transplantation. <i>Blood Advances</i> , 2020 , 4, 2073-2083	7.8	21
187	Iron Overload Is Associated with Delayed Engraftment and Increased Nonrelapse Mortality in Recipients of Umbilical Cord Blood Hematopoietic Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2020 , 26, 1697-1703	4.7	3
186	Immune phenotype and histopathological growth pattern in patients with colorectal liver metastases. <i>British Journal of Cancer</i> , 2020 , 122, 1518-1524	8.7	17
185	Single Nucleotide Polymorphisms in MiRNA Binding Sites of Nucleotide Excision Repair-Related Genes Predict Clinical Benefit of Oxaliplatin in FOLFOXIRI Plus Bevacizumab: Analysis of the TRIBE Trial. <i>Cancers</i> , 2020 , 12,	6.6	2
184	Outcomes of Allogeneic Hematopoietic Cell Transplantation after Salvage Therapy with Blinatumomab in Patients with Relapsed/Refractory Acute Lymphoblastic Leukemia. <i>Biology of Blood and Marrow Transplantation</i> , 2020 , 26, 1084-1090	4.7	11
183	Epidermal growth factor receptor mRNA expression: A potential molecular escape mechanism from regorafenib. <i>Cancer Science</i> , 2020 , 111, 441-450	6.9	5
182	A polymorphism within the R-spondin 2 gene predicts outcome in metastatic colorectal cancer patients treated with FOLFIRI/bevacizumab: data from FIRE-3 and TRIBE trials. <i>European Journal of Cancer</i> , 2020 , 131, 89-97	7.5	3

181	Hemorrhagic Cystitis in Patients Undergoing Allogeneic Hematopoietic Cell Transplant with Post Transplant Cyclophosphamide As GvHD Prophylaxis. <i>Blood</i> , 2020 , 136, 21-22	2.2	
180	Clinical Impact of Cytokine Release Syndrome on Outcomes of Peripheral Blood Stem Cell Haploidentical Hematopoietic Cell Transplantation with Post-Transplant Cyclophosphamide. <i>Blood</i> , 2020 , 136, 10-11	2.2	
179	Total Marrow and Lymphoid Irradiation (TMLI) at a Dose of 2000cGy in Combination with Post-Transplant Cyclophosphamide (PTCy)-Based Graft Versus Host Disease (GvHD) Prophylaxis Is Safe and Associated with Favorable GvHD-Free/Relapse-Free Survival at 1 Year in Patients with	2.2	1
178	Healthcare Resource Utilization in Transplant Patients Who Are at a Higher-Risk to Develop Cytomegalovirus Infection during Their Primary Transplant-Related Hospitalization. <i>Blood</i> , 2020 , 136, 16-17	2.2	
177	Pulmonary hypertension is associated with increased nonrelapse mortality after allogeneic hematopoietic cell transplantation for myelofibrosis. <i>Bone Marrow Transplantation</i> , 2020 , 55, 877-883	4.4	6
176	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer. <i>PLoS ONE</i> , 2020 , 15, e0239439	3.7	3
175	Red blood cell and platelet transfusion support in the first 30 and 100 days after allogeneic hematopoietic cell transplant. <i>Transfusion</i> , 2020 , 60, 2225-2242	2.9	2
174	Outcome of Allogeneic Hematopoietic Cell Transplantation after Venetoclax and Hypomethylating Agent Therapy for Acute Myelogenous Leukemia. <i>Biology of Blood and Marrow Transplantation</i> , 2020 , 26, e322-e327	4.7	19
173	Favorable outcomes for allogeneic hematopoietic cell transplantation in elderly patients with NPM1-mutated and FLT3-ITD-negative acute myeloid leukemia. <i>Bone Marrow Transplantation</i> , 2020 , 55, 473-475	4.4	9
172	Patient-reported outcomes of the relative severity of side effects from cancer radiotherapy. Supportive Care in Cancer, 2020 , 28, 309-316	3.9	5
171	Long-Term Outcomes of Patients with Acute Myelogenous Leukemia Treated with Myeloablative Fractionated Total Body Irradiation TBI-Based Conditioning with a Tacrolimus- and Sirolimus-Based Graft-versus-Host Disease Prophylaxis Regimen: 6-Year Follow-Up from a Single Center. <i>Biology of</i>	4.7	5
170	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
169	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
168	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
167	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
166	Radiation-Related Toxicities Using Organ Sparing Total Marrow Irradiation Transplant Conditioning Regimens. <i>International Journal of Radiation Oncology Biology Physics</i> , 2019 , 105, 1025-1033	4	19
165	Allogeneic Hematopoietic Cell Transplantation Outcomes in Patients Carrying Isocitrate Dehydrogenase Mutations. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019 , 19, e400-e405	2	3
164	MIPSS70+ v2.0 predicts long-term survival in myelofibrosis after allogeneic HCT with the Flu/Mel conditioning regimen. <i>Blood Advances</i> , 2019 , 3, 83-95	7.8	27

163	Influence of the facility caseload on the subsequent survival of men with localized prostate cancer undergoing radical prostatectomy. <i>Cancer</i> , 2019 , 125, 3853-3863	6.4	3
162	The Cancer Bell: Too Much of a Good Thing?. <i>International Journal of Radiation Oncology Biology Physics</i> , 2019 , 105, 247-253	4	1
161	Association of leukemia genetics with response to venetoclax and hypomethylating agents in relapsed/refractory acute myeloid leukemia. <i>American Journal of Hematology</i> , 2019 , 94, E253-E255	7.1	26
160	RBC and platelet transfusion support in the first 30 and 100 days after haploidentical hematopoietic stem cell transplantation. <i>Transfusion</i> , 2019 , 59, 3371-3385	2.9	5
159	Updated Analysis: Novel Machine-Learning-Based Sepsis Prediction Model for Patients Undergoing Hematopoietic Stem Cell Transplantation (Early Sepsis Prediction/Identification for Transplant Recipients: ESPRIT). <i>Blood</i> , 2019 , 134, 4477-4477	2.2	О
158	Ruxolitinib as Salvage Therapy for Chronic Graft-versus-Host Disease. <i>Biology of Blood and Marrow Transplantation</i> , 2019 , 25, 265-269	4.7	44
157	Role of CCL5 and CCR5 gene polymorphisms in epidermal growth factor receptor signalling blockade in metastatic colorectal cancer: analysis of the FIRE-3 trial. <i>European Journal of Cancer</i> , 2019 , 107, 100-114	7.5	5
156	Genetic variants in CCL5 and CCR5 genes and serum VEGF-A levels predict efficacy of bevacizumab in metastatic colorectal cancer patients. <i>International Journal of Cancer</i> , 2019 , 144, 2567-2577	7.5	2
155	Prognostic Value of ACVRL1 Expression in Metastatic Colorectal Cancer Patients Receiving First-line Chemotherapy With Bevacizumab: Results From the Triplet Plus Bevacizumab (TRIBE) Study. <i>Clinical Colorectal Cancer</i> , 2018 , 17, e471-e488	3.8	4
154	Second primary acute lymphoblastic leukemia in adults: a SEER analysis of incidence and outcomes. <i>Cancer Medicine</i> , 2018 , 7, 499-507	4.8	20
153	Efficacy of the combination of venetoclax and hypomethylating agents in relapsed/refractory acute myeloid leukemia. <i>Haematologica</i> , 2018 , 103, e404-e407	6.6	131
152	Gene Polymorphisms in the CCL5/CCR5 Pathway as a Genetic Biomarker for Outcome and Hand-Foot Skin Reaction in Metastatic Colorectal Cancer Patients Treated With Regorafenib. <i>Clinical Colorectal Cancer</i> , 2018 , 17, e395-e414	3.8	16
151	Clinical relevance of EMT and stem-like gene expression in circulating tumor cells of metastatic colorectal cancer patients. <i>Pharmacogenomics Journal</i> , 2018 , 18, 29-34	3.5	25
150	A genetic variant in Rassf1a predicts outcome in mCRC patients treated with cetuximab plus chemotherapy: results from FIRE-3 and JACCRO 05 and 06 trials. <i>Pharmacogenomics Journal</i> , 2018 , 18, 43-48	3.5	1
149	Prognostic impact of FOXF1 polymorphisms in gastric cancer patients. <i>Pharmacogenomics Journal</i> , 2018 , 18, 262-269	3.5	2
148	Potential role of PIN1 genotypes in predicting benefit from oxaliplatin-based and irinotecan-based treatment in patients with metastatic colorectal cancer. <i>Pharmacogenomics Journal</i> , 2018 , 18, 623-632	3.5	4
147	Melphalan-Based Reduced-Intensity Conditioning is Associated with Favorable Disease Control and Acceptable Toxicities in Patients Older Than 70 with Hematologic Malignancies Undergoing Allogeneic Hematopoietic Stem Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> ,	4.7	9
146	2018, 24, 1828-1835 Immune-related Genes to Dominate Neutrophil-lymphocyte Ratio (NLR) Associated With Survival of Cetuximab Treatment in Metastatic Colorectal Cancer. Clinical Colorectal Cancer, 2018, 17, e741-e749	3.8	16

145	NOS2 polymorphisms in prediction of benefit from first-line chemotherapy in metastatic colorectal cancer patients. <i>PLoS ONE</i> , 2018 , 13, e0193640	3.7	3
144	A Machine-Learning Sepsis Prediction Model for Patients Undergoing Hematopoietic Cell Transplantation. <i>Blood</i> , 2018 , 132, 711-711	2.2	1
143	Cytokine Gene Polymorphisms Are Associated with Disease Response to Blinatumomab in Patients with B-Cell Acute Lymphoblastic Leukemia. <i>Blood</i> , 2018 , 132, 1549-1549	2.2	
142	Effect of Vancomycin-Resistance Enterococci Colonization Status Prior to Allogeneic Hematopoietic Cell Transplantation on Transplant Outcomes: A Single Center Retrospective Experience. <i>Blood</i> , 2018 , 132, 3386-3386	2.2	
141	Clinical Outcomes of MDS Patients Who Were Allogeneic Hematopoietic Stem Cell Transplant Candidates but Did Not Proceed with Transplantation. <i>Blood</i> , 2018 , 132, 2181-2181	2.2	
140	Outcomes after Allogeneic Stem Cell Transplantation in Patients with Double-Hit and Double-Expressor Lymphoma. <i>Biology of Blood and Marrow Transplantation</i> , 2018 , 24, 514-520	4.7	27
139	A Polymorphism within the Vitamin D Transporter Gene Predicts Outcome in Metastatic Colorectal Cancer Patients Treated with FOLFIRI/Bevacizumab or FOLFIRI/Cetuximab. <i>Clinical Cancer Research</i> , 2018 , 24, 784-793	12.9	14
138	Prediction of cardiovascular disease among hematopoietic cell transplantation survivors. <i>Blood Advances</i> , 2018 , 2, 1756-1764	7.8	23
137	Genetic variants associated with colorectal brain metastases susceptibility and survival. <i>Pharmacogenomics Journal</i> , 2017 , 17, 29-35	3.5	6
136	Estrogen receptor-beta genetic variations and overall survival in patients with locally advanced gastric cancer. <i>Pharmacogenomics Journal</i> , 2017 , 17, 36-41	3.5	8
135	Genetic variations in immunomodulatory pathways to predict survival in patients with locoregional gastric cancer. <i>Pharmacogenomics Journal</i> , 2017 , 17, 528-534	3.5	7
134	Genetic variants of DNA repair-related genes predict efficacy of TAS-102 in patients with refractory metastatic colorectal cancer. <i>Annals of Oncology</i> , 2017 , 28, 1015-1022	10.3	20
133	Predictive value of TLR7 polymorphism for cetuximab-based chemotherapy in patients with metastatic colorectal cancer. <i>International Journal of Cancer</i> , 2017 , 141, 1222-1230	7.5	14
132	Single nucleotide polymorphisms in the IGF-IRS pathway are associated with outcome in mCRC patients enrolled in the FIRE-3 trial. <i>International Journal of Cancer</i> , 2017 , 141, 383-392	7.5	5
131	Autophagy-related polymorphisms predict hypertension in patients with metastatic colorectal cancer treated with FOLFIRI and bevacizumab: Results from TRIBE and FIRE-3 trials. <i>European Journal of Cancer</i> , 2017 , 77, 13-20	7.5	15
130	Potential role of polymorphisms in the transporter genes ENT1 and MATE1/OCT2 in predicting TAS-102 efficacy and toxicity in patients with refractory metastatic colorectal cancer. <i>European Journal of Cancer</i> , 2017 , 86, 197-206	7.5	16
129	The presence of Philadelphia chromosome does not confer poor prognosis in adult pre-B acute lymphoblastic leukaemia in the tyrosine kinase inhibitor era - a surveillance, epidemiology, and end results database analysis. <i>British Journal of Haematology</i> , 2017 , 179, 618-626	4.5	7
128	Tandem repeat variation near the HIC1 (hypermethylated in cancer 1) promoter predicts outcome of oxaliplatin-based chemotherapy in patients with metastatic colorectal cancer. <i>Cancer</i> , 2017 , 123, 49	506-451	4 ⁴

127	Impact of genetic variations in the MAPK signaling pathway on outcome in metastatic colorectal cancer patients treated with first-line FOLFIRI and bevacizumab: data from FIRE-3 and TRIBE trials. <i>Annals of Oncology</i> , 2017 , 28, 2780-2785	10.3	19
126	CXCR4 polymorphism predicts progression-free survival in metastatic colorectal cancer patients treated with first-line bevacizumab-based chemotherapy. <i>Pharmacogenomics Journal</i> , 2017 , 17, 543-550) ^{3.5}	9
125	Genetic variations within the vitamin C transporter genes to predict outcome in metastatic colorectal cancer patients treated with first-line FOLFIRI and bevacizumab: Data from FIRE-3 trial <i>Journal of Clinical Oncology</i> , 2017 , 35, 11507-11507	2.2	1
124	Association of immune-related genes to neutrophil-lymphocyte ratio (NLR) with survival of cetuximab treatment for metastatic colorectal cancer (mCRC): JACCRO CC-05/06AR <i>Journal of Clinical Oncology</i> , 2017 , 35, 11613-11613	2.2	1
123	Trends in colorectal cancer mortality in hispanics: a SEER analysis. <i>Oncotarget</i> , 2017 , 8, 108771-108777	3.3	12
122	Association of genetic variations in genes implicated in the axis with outcome in patients (pts) with metastatic colorectal cancer (mCRC) treated with cetuximab plus chemotherapy <i>Journal of Clinical Oncology</i> , 2017 , 35, 3585-3585	2.2	1
121	Germline polymorphisms in genes involved in the Hippo pathway as recurrence biomarkers in stages II/III colon cancer. <i>Pharmacogenomics Journal</i> , 2016 , 16, 312-9	3.5	11
120	Expression of Genes Involved in Vascular Morphogenesis and Maturation Predicts Efficacy of Bevacizumab-Based Chemotherapy in Patients Undergoing Liver Resection. <i>Molecular Cancer Therapeutics</i> , 2016 , 15, 2814-2821	6.1	6
119	A Phase II Biomarker-Embedded Study of Lapatinib plus Capecitabine as First-line Therapy in Patients with Advanced or Metastatic Gastric Cancer. <i>Molecular Cancer Therapeutics</i> , 2016 , 15, 2251-8	6.1	5
118	Clinical Significance of TLR1 I602S Polymorphism for Patients with Metastatic Colorectal Cancer Treated with FOLFIRI plus Bevacizumab. <i>Molecular Cancer Therapeutics</i> , 2016 , 15, 1740-5	6.1	7
117	Metastatic Colorectal Cancer in Hispanics: Treatment Outcomes in a Treated Population. <i>Clinical Colorectal Cancer</i> , 2016 , 15, e221-e227	3.8	6
116	Impact of access to NCI- and NCCN-designated cancer centers on outcomes for multiple myeloma patients: A SEER registry analysis. <i>Cancer</i> , 2016 , 122, 618-25	6.4	16
115	TWIST1 Polymorphisms Predict Survival in Patients with Metastatic Colorectal Cancer Receiving First-Line Bevacizumab plus Oxaliplatin-Based Chemotherapy. <i>Molecular Cancer Therapeutics</i> , 2016 , 15, 1405-11	6.1	8
114	Prognostic Impact of IL6 Genetic Variants in Patients with Metastatic Colorectal Cancer Treated with Bevacizumab-Based Chemotherapy. <i>Clinical Cancer Research</i> , 2016 , 22, 3218-26	12.9	16
113	Persistent Racial/Ethnic Disparities in Outcomes for Multiple Myeloma: A SEER-Database Update. <i>Blood</i> , 2016 , 128, 1191-1191	2.2	2
112	Racial/Ethnic Disparities in NK/T-Cell Non-Hodgkin Lymphomas in the US: A SEER Analysis. <i>Blood</i> , 2016 , 128, 3612-3612	2.2	1
111	Immune-related genes to predict clinical outcome of cetuximab (cet) treatment for metastatic colorectal cancer (mCRC): Immuno-Oncology assay research <i>Journal of Clinical Oncology</i> , 2016 , 34, 115	9 ² 1 ⁻² 11!	591
110	Impact of sex, age, and ethnicity/race on the survival of patients with rectal cancer in the United States from 1988 to 2012. <i>Oncotarget</i> , 2016 , 7, 53668-53678	3.3	16

Gastric Cancer Among Asian Americans 2016, 249-269 109 1 AXIN2 expression predicts prostate cancer recurrence and regulates invasion and tumor growth. 108 4.2 13 Prostate, 2016, 76, 597-608 Prognostic factors for melanoma brain metastases treated with stereotactic radiosurgery. Journal 8 107 3.2 of Neurosurgery, 2016, 125, 31-39 Combined assessment of EGFR-related molecules to predict outcome of 1st-line cetuximab-containing chemotherapy for metastatic colorectal cancer. Cancer Biology and Therapy, 106 4.6 13 **2016**, 17, 751-9 Association of variants in genes encoding for macrophage-related functions with clinical outcome 105 10.3 20 in patients with locoregional gastric cancer. Annals of Oncology, 2015, 26, 332-9 Primary tumor location as a prognostic factor in metastatic colorectal cancer. Journal of the 298 104 9.7 National Cancer Institute, 2015, 107, Cytokeratin-20 and Survivin-Expressing Circulating Tumor Cells Predict Survival in Metastatic Colorectal Cancer Patients by a Combined Immunomagnetic qRT-PCR Approach. Molecular Cancer 6.1 103 21 Therapeutics, 2015, 14, 2401-8 Variations in genes involved in dormancy associated with outcome in patients with resected 10.3 6 102 colorectal liver metastases. Annals of Oncology, 2015, 26, 1728-33 BRAF and RAS mutations as prognostic factors in metastatic colorectal cancer patients undergoing 8.7 101 111 liver resection. British Journal of Cancer, 2015, 112, 1921-8 Genetic variants within obesity-related genes are associated with tumor recurrence in patients with 1.9 stages II/III colon cancer. Pharmacogenetics and Genomics, 2015, 25, 30-7 FCGR polymorphisms and cetuximab efficacy in chemorefractory metastatic colorectal cancer: an 99 19.2 17 international consortium study. Gut, 2015, 64, 921-8 Variations in genes regulating tumor-associated macrophages (TAMs) to predict outcomes of 98 bevacizumab-based treatment in patients with metastatic colorectal cancer: results from TRIBE and 10.3 20 FIRE3 trials. Annals of Oncology, 2015, 26, 2450-6 Polymorphisms in Genes Involved in EGFR Turnover Are Predictive for Cetuximab Efficacy in 6.1 97 3 Colorectal Cancer. Molecular Cancer Therapeutics, 2015, 14, 2374-81 Persistent Disparities Among Patients With T-Cell Non-Hodgkin Lymphomas and B-Cell Diffuse Large Cell Lymphomas Over 40 Years: A SEER Database Review. Clinical Lymphoma, Myeloma and 96 2 20 Leukemia, **2015**, 15, 578-85 Genetic variants of kinase suppressors of Ras (KSR1) to predict survival in patients with 95 3.5 1 EREpositive advanced breast cancer. Pharmacogenomics Journal, 2015, 15, 235-40 Genes involved in pericyte-driven tumor maturation predict treatment benefit of first-line FOLFIRI plus bevacizumab in patients with metastatic colorectal cancer. Pharmacogenomics Journal, 2015, 94 3.5 24 15, 69-76 Variations in genes involved in immune response checkpoints and association with outcomes in 8 93 3.5 patients with resected colorectal liver metastases. Pharmacogenomics Journal, 2015, 15, 521-9 Genetic variations in angiopoietin and pericyte pathways and clinical outcome in patients with 92 6.4 9 resected colorectal liver metastases. Cancer, 2015, 121, 1898-905

91	Distinct gene expression profiles of proximal and distal colorectal cancer: implications for cytotoxic and targeted therapy. <i>Pharmacogenomics Journal</i> , 2015 , 15, 354-62	3.5	36
90	Integrin genetic variants and stage-specific tumor recurrence in patients with stage II and III colon cancer. <i>Pharmacogenomics Journal</i> , 2015 , 15, 226-34	3.5	14
89	A let-7 microRNA-binding site polymorphism in KRAS predicts improved outcome in patients with metastatic colorectal cancer treated with salvage cetuximab/panitumumab monotherapy. <i>Clinical Cancer Research</i> , 2014 , 20, 4499-4510	12.9	44
88	Association of common gene variants in the WNT/Latenin pathway with colon cancer recurrence. <i>Pharmacogenomics Journal</i> , 2014 , 14, 142-50	3.5	26
87	CXCR2 inhibition enhances sulindac-mediated suppression of colon cancer development. <i>International Journal of Cancer</i> , 2014 , 135, 232-7	7.5	11
86	The cyclin D1 (CCND1) rs9344 G>A polymorphism predicts clinical outcome in colon cancer patients treated with adjuvant 5-FU-based chemotherapy. <i>Pharmacogenomics Journal</i> , 2014 , 14, 130-4	3.5	15
85	Insulin-like growth factor receptor polymorphism defines clinical outcome in estrogen receptor-positive breast cancer patients treated with tamoxifen. <i>Pharmacogenomics Journal</i> , 2014 , 14, 28-34	3.5	25
84	Outcome disparities among ethnic subgroups of Waldenstrfhß macroglobulinemia: a population-based study. <i>Oncology</i> , 2014 , 86, 253-62	3.6	20
83	Variable risk of second primary malignancy in multiple myeloma patients of different ethnic subgroups. <i>Blood Cancer Journal</i> , 2014 , 4, e243	7	13
82	Impact of sex on the survival of patients with hepatocellular carcinoma: a Surveillance, Epidemiology, and End Results analysis. <i>Cancer</i> , 2014 , 120, 3707-16	6.4	80
81	Gender-specific profiling in SCN1A polymorphisms and time-to-recurrence in patients with stage II/III colorectal cancer treated with adjuvant 5-fluoruracil chemotherapy. <i>Pharmacogenomics Journal</i> , 2014 , 14, 135-41	3.5	10
80	Prospective study of EGFR intron 1 (CA)n repeats variants as predictors of benefit from cetuximab and irinotecan in chemo-refractory metastatic colorectal cancer (mCRC) patients. <i>Pharmacogenomics Journal</i> , 2014 , 14, 322-7	3.5	10
79	Plastin polymorphisms predict gender- and stage-specific colon cancer recurrence after adjuvant chemotherapy. <i>Molecular Cancer Therapeutics</i> , 2014 , 13, 528-39	6.1	27
78	Single nucleotide polymorphisms in AREG and EREG are prognostic biomarkers in locally advanced gastric cancer patients after surgery with curative intent. <i>Pharmacogenetics and Genomics</i> , 2014 , 24, 539	9 ¹ 49	2
77	Prognostic impact of the c-MET polymorphism on the clinical outcome in locoregional gastric cancer patients. <i>Pharmacogenetics and Genomics</i> , 2014 , 24, 588-96	1.9	8
76	Secondary Acute Lymphoblastic Leukemia after Primary Solid Organ Malignancy: A SEER Analysis of Incidence and Outcomes. <i>Blood</i> , 2014 , 124, 935-935	2.2	2
75	mRNA expression levels of candidate genes and clinical outcome in mCRC patients treated with FOLFOXIRI plus bevacizumab (bev) or FOLFIRI plus bev in the TRIBE study <i>Journal of Clinical Oncology</i> , 2014 , 32, 3640-3640	2.2	1
74	Pharmacogenetic profiling of CD133 is associated with response rate (RR) and progression-free survival (PFS) in patients with metastatic colorectal cancer (mCRC), treated with bevacizumab-based chemotherapy. <i>Pharmacogenomics Journal</i> , 2013 , 13, 173-80	3.5	32

(2012-2013)

73	The endoplasmic reticulum stress marker, glucose-regulated protein-78 (GRP78) in visceral adipocytes predicts endometrial cancer progression and patient survival. <i>Gynecologic Oncology</i> , 2013 , 128, 552-9	4.9	46
72	Timeliness of adjuvant chemotherapy for stage III adenocarcinoma of the colon: a measure of quality of care. <i>Clinical Colorectal Cancer</i> , 2013 , 12, 275-9	3.8	22
71	Intratumoral expression profiling of genes involved in angiogenesis in colorectal cancer patients treated with chemotherapy plus the VEGFR inhibitor PTK787/ZK 222584 (vatalanib). <i>Pharmacogenomics Journal</i> , 2013 , 13, 410-6	3.5	12
70	Reply to J. Zhang et al. <i>Journal of Clinical Oncology</i> , 2013 , 31, 816-7	2.2	
69	Prognostic role of lemur tyrosine kinase-3 germline polymorphisms in adjuvant gastric cancer in Japan and the United States. <i>Molecular Cancer Therapeutics</i> , 2013 , 12, 2261-72	6.1	17
68	An individual coding polymorphism and the haplotype of the SPARC gene predict gastric cancer recurrence. <i>Pharmacogenomics Journal</i> , 2013 , 13, 342-8	3.5	4
67	Prospective validation of candidate SNPs of VEGF/VEGFR pathway in metastatic colorectal cancer patients treated with first-line FOLFIRI plus bevacizumab. <i>PLoS ONE</i> , 2013 , 8, e66774	3.7	55
66	Gene expression profiles and tumor locations in colorectal cancer (left vs. right vs. rectum) <i>Journal of Clinical Oncology</i> , 2013 , 31, 3527-3527	2.2	7
65	Ethnic Disparities and Their Association With Outcomes In Chronic Myeloid Leukemia. <i>Blood</i> , 2013 , 122, 2917-2917	2.2	0
64	Outcome disparities in multiple myeloma: a SEER-based comparative analysis of ethnic subgroups. British Journal of Haematology, 2012 , 158, 91-8	4.5	69
63	9-cis retinoic acid promotes lymphangiogenesis and enhances lymphatic vessel regeneration: therapeutic implications of 9-cis retinoic acid for secondary lymphedema. <i>Circulation</i> , 2012 , 125, 872-82	16.7	65
62	The CXCR2 antagonist, SCH-527123, shows antitumor activity and sensitizes cells to oxaliplatin in preclinical colon cancer models. <i>Molecular Cancer Therapeutics</i> , 2012 , 11, 1353-64	6.1	79
61	A randomized phase II of gemcitabine and sorafenib versus sorafenib alone in patients with metastatic pancreatic cancer. <i>Investigational New Drugs</i> , 2012 , 30, 1175-83	4.3	33
60	The Cyclin D1 (CCND1) A870G polymorphism predicts clinical outcome to lapatinib and capecitabine in HER2-positive metastatic breast cancer. <i>Annals of Oncology</i> , 2012 , 23, 1455-64	10.3	8
59	Interleukin-8 and its receptor CXCR2 in the tumour microenvironment promote colon cancer growth, progression and metastasis. <i>British Journal of Cancer</i> , 2012 , 106, 1833-41	8.7	197
58	Influence of sex on the survival of patients with esophageal cancer. <i>Journal of Clinical Oncology</i> , 2012 , 30, 2265-72	2.2	85
57	Prospective evaluation of candidate SNPs of VEGF/VEGFR pathway in metastatic colorectal cancer (mCRC) patients (pts) treated with first-line FOLFIRI plus bevacizumab (BV) <i>Journal of Clinical Oncology</i> , 2012 , 30, 3518-3518	2.2	1
56	Ethnic gene profile of genes involved in angiogenesis to predict regional bevacizumab efficacy difference in gastric cancer <i>Journal of Clinical Oncology</i> , 2012 , 30, 4026-4026	2.2	4

55	LMTK3 polymorphism in patients with metastatic colon cancer <i>Journal of Clinical Oncology</i> , 2012 , 30, 471-471	2.2	4
54	Ethnic Disparities in Chronic Lymphocytic Leukemia Survival: A SEER Database Review. <i>Blood</i> , 2012 , 120, 757-757	2.2	О
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