

Martin D Berger

List of Publications by Year in Descending Order

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Version: 2024-04-18

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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.

135
papers

2,530
citations

19
h-index

49
g-index

146
ext. papers

3,646
ext. citations

4.4
avg, IF

4.78
L-index

#	Paper	IF	Citations
135	Impact on survival through consolidation radiotherapy for diffuse large B-cell lymphoma: a comprehensive meta-analysis. <i>Haematologica</i> , 2021 , 106, 1923-1931	6.6	1
134	Role of enterocyte-specific gene polymorphisms in response to adjuvant treatment for stage III colorectal cancer. <i>Pharmacogenetics and Genomics</i> , 2021 , 31, 10-16	1.9	1
133	Molecular differences between lymph nodes and distant metastases compared with primaries in colorectal cancer patients. <i>Npj Precision Oncology</i> , 2021 , 5, 95	9.8	1
132	Refining the ITBCC tumor budding scoring system with a "zero-budding" category in colorectal cancer. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 1	5.1	2
131	CCR5 is a potential therapeutic target for cancer. <i>Expert Opinion on Therapeutic Targets</i> , 2021 , 25, 311-324	3.4	3
130	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	0
129	LAG-3 Expression Predicts Outcome in Stage II Colon Cancer. <i>Journal of Personalized Medicine</i> , 2021 , 11,	3.6	1
128	RHAMM in liver metastases of stage IV colorectal cancer with mismatch-repair proficient status correlates with tumor budding, cytotoxic T-cells and PD-1/PD-L1. <i>Pathology Research and Practice</i> , 2021 , 223, 153486	3.4	2
127	Tumour budding in solid cancers. <i>Nature Reviews Clinical Oncology</i> , 2021 , 18, 101-115	19.4	43
126	Clinical significance of enterocyte-specific gene polymorphisms as candidate markers of oxaliplatin-based treatment for metastatic colorectal cancer. <i>Pharmacogenomics Journal</i> , 2021 , 21, 285-295	3.5	0
125	Potential Molecular Cross Talk Among CCR5 Pathway Predicts Regorafenib Responsiveness in Metastatic Colorectal Cancer Patients. <i>Cancer Genomics and Proteomics</i> , 2021 , 18, 317-324	3.3	0
124	Tumour budding and its clinical implications in gastrointestinal cancers. <i>British Journal of Cancer</i> , 2020 , 123, 700-708	8.7	6
123	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
122	Epidermal growth factor receptor mRNA expression: A potential molecular escape mechanism from regorafenib. <i>Cancer Science</i> , 2020 , 111, 441-450	6.9	5
121	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
120	Are tumour grade and tumour budding equivalent in colorectal cancer? A retrospective analysis of 771 patients. <i>European Journal of Cancer</i> , 2020 , 130, 139-145	7.5	4
119	Comprehensive molecular profiling of IDH1/2 mutant biliary cancers (BC).. <i>Journal of Clinical Oncology</i> , 2020 , 38, 479-479	2.2	6

118	Gene polymorphisms of CCL3, CCL4, CCL5, and CCR5 network in metastatic colorectal cancer patients treated with regorafenib.. <i>Journal of Clinical Oncology</i> , 2020 , 38, 199-199	2.2	
117	Variation in genetic polymorphisms and gene expression of HLA-E to predict outcomes in metastatic colorectal cancer (mCRC) patients (pts) treated with first-line FOLFIRI/cetuximab: Data from the phase III FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2020 , 38, 245-245	2.2	0
116	Polymorphisms of genes encoding for regulatory proteins in the coagulation cascade to predict outcome for stage II and III colon cancer.. <i>Journal of Clinical Oncology</i> , 2020 , 38, 227-227	2.2	
115	Comprehensive gene expression analysis of IDH1/2 mutant biliary cancers (BC).. <i>Journal of Clinical Oncology</i> , 2020 , 38, 4598-4598	2.2	
114	Simple acute phase protein score to predict long-term survival in patients with acute myeloid leukemia. <i>Hematological Oncology</i> , 2020 , 38, 74-81	1.3	4
113	Preservation of Epstein-Barr virus status and mismatch repair protein status along the metastatic course of gastric cancer. <i>Histopathology</i> , 2020 , 76, 740-747	7.3	3
112	The impact of ARID1A mutation on molecular characteristics in colorectal cancer. <i>European Journal of Cancer</i> , 2020 , 140, 119-129	7.5	13
111	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
110	Comprehensive Genomic Profiling of Gastroenteropancreatic Neuroendocrine Neoplasms (GEP-NENs). <i>Clinical Cancer Research</i> , 2020 , 26, 5943-5951	12.9	17
109	Multicenter International Society for Immunotherapy of Cancer Study of the Consensus Immunoscore for the Prediction of Survival and Response to Chemotherapy in Stage III Colon Cancer. <i>Journal of Clinical Oncology</i> , 2020 , 38, 3638-3651	2.2	47
108	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
107	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
106	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
105	A polymorphism in the cachexia-associated gene INHBA predicts efficacy of regorafenib in patients with refractory metastatic colorectal cancer 2020 , 15, e0239439		
104	Molecular Profiling of Appendiceal Adenocarcinoma and Comparison with Right-sided and Left-sided Colorectal Cancer. <i>Clinical Cancer Research</i> , 2019 , 25, 3096-3103	12.9	30
103	Impact of polymorphisms within genes involved in regulating DNA methylation in patients with metastatic colorectal cancer enrolled in three independent, randomised, open-label clinical trials: a meta-analysis from TRIBE, MAVERICC and FIRE-3. <i>European Journal of Cancer</i> , 2019 , 111, 138-147	7.5	3
102	AMPK variant, a candidate of novel predictor for chemotherapy in metastatic colorectal cancer: A meta-analysis using TRIBE, MAVERICC and FIRE3. <i>International Journal of Cancer</i> , 2019 , 145, 2082-2090	7.5	0
101	Comprehensive molecular characterization of brain metastases (BM) from colorectal cancer (CRC). <i>Annals of Oncology</i> , 2019 , 30, v764	10.3	2

100	BRCA1 genetic variant to predict survival in metastatic colorectal cancer (mCRC) patients (pts) treated with FOLFIRI/bevacizumab (bev): Results from phase III TRIBE and FIRE-3 trials.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3145-3145	2.2	2
99	Gene mutations of SWI/SNF complex and molecular profile in colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3600-3600	2.2	1
98	Characteristics of colorectal cancer (CRC) patients with BRCA1 and BRCA2 mutations.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 606-606	2.2	3
97	Th17 cell pathway-related genetic variants in metastatic colorectal cancer: A meta-analysis using TRIBE, MAVERICC, and FIRE-3.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 594-594	2.2	
96	Genetic variants in the lipopolysaccharide (LPS) receptor complex and TLR4 expression levels to predict efficacy of cetuximab (cet) in patients (pts) with metastatic colorectal cancer (mCRC): Data from the FIRE-3 phase III trial.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 564-564	2.2	
95	Polymorphisms in the telomerase complex to predict outcome in patients (pts) with metastatic colorectal cancer (mCRC): Data from TRIBE and FIRE-3 phase III trials.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 566-566	2.2	
94	Genetic variations within the CD40L immune stimulating gene predict outcome for mCRC patients treated with first-line FOLFIRI/bevacizumab: Data from FIRE-3 and TRIBE.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 558-558	2.2	
93	Comprehensive molecular profiling of signet-ring-cell carcinoma (SRCC) from the stomach and colon.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 63-63	2.2	
92	Role of enterocyte-specific gene polymorphisms in adjuvant treatment for stage III colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 550-550	2.2	
91	Gene expression and genetic variants in Parkinson's disease (PD) genes to predict outcome in metastatic colorectal cancer (mCRC): Data from FIRE-3 phase III trial.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3595-3595	2.2	0
90	Molecular differences between lymph nodes (LNs) and distant metastases (mets) in colorectal cancer (CRC).. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3130-3130	2.2	
89	Polymorphisms in the dopamine (DA) signaling to predict outcome in patients (pts) with metastatic colorectal cancer (mCRC): Data from TRIBE, MAVERICC, and FIRE-3 phase III trials.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3048-3048	2.2	0
88	Association of genetic variations within the T-cell costimulatory LIGHT gene with outcome in stage II and III colon cancer.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 2633-2633	2.2	
87	Prognostic Effect of Adenosine-related Genetic Variants in Metastatic Colorectal Cancer Treated With Bevacizumab-based Chemotherapy. <i>Clinical Colorectal Cancer</i> , 2019 , 18, e8-e19	3.8	9
86	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
85	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
84	Validation of the International Tumor Budding Consensus Conference 2016 recommendations on tumor budding in stage I-IV colorectal cancer. <i>Human Pathology</i> , 2019 , 85, 145-151	3.7	34
83	B cell and B cell-related pathways for novel cancer treatments. <i>Cancer Treatment Reviews</i> , 2019 , 73, 10-19	4.4	59

82	Stromal PD-1/PD-L1 Expression Predicts Outcome in Colon Cancer Patients. <i>Clinical Colorectal Cancer</i> , 2019 , 18, e20-e38	3.8	37
81	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
80	Outlooks on Epstein-Barr virus associated gastric cancer. <i>Cancer Treatment Reviews</i> , 2018 , 66, 15-22	14.4	74
79	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
78	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
77	International validation of the consensus Immunoscore for the classification of colon cancer: a prognostic and accuracy study. <i>Lancet, The</i> , 2018 , 391, 2128-2139	40	910
76	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
75	Circadian clock gene PER1 mutations in colorectal cancer (CRC).. <i>Journal of Clinical Oncology</i> , 2018 , 36, 12106-12106	2.2	2
74	Polymorphism in the circadian clock pathway to predict outcome in patients (pts) with metastatic colorectal cancer (mCRC): Data from TRIBE and FIRE-3 phase III trials.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 3576-3576	2.2	1
73	Polymorphism in cancer-associated fibroblasts (CAFs) related genes and clinical outcome in metastatic colorectal cancer (mCRC) patients (pts) enrolled in two independent randomized phase III trials: TRIBE and FIRE-3.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 645-645	2.2	1
72	Matrix metalloproteinase-related gene polymorphisms to predict efficacy of regorafenib in patients with metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 692-692	2.2	1
71	Pharmacogenomics in colorectal cancer: current role in clinical practice and future perspectives. <i>Journal of Cancer Metastasis and Treatment</i> , 2018 , 4,	3.8	2
70	Association of genetic variations within the PD-L2 immune checkpoint gene with outcome in stage II and III colon cancer.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 626-626	2.2	
69	The impact of Tfh cell/ B cell pathway-related genetic variants in metastatic colorectal cancer patients with bevacizumab-based chemotherapy.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 651-651	2.2	
68	Single nucleotide polymorphisms in miRNA binding sites of nucleotide excision repair-related genes to predict clinical benefit of oxaliplatin in FOLFIRI plus bevacizumab in TRIBE trial.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 663-663	2.2	
67	Polymorphisms in beta-defensin pathways and clinical outcomes in metastatic colorectal cancer patients treated with FOLFIRI-bevacizumab in two randomized phase III trials.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 662-662	2.2	
66	Genetic variants in methylation and demethylation pathways to predict clinical outcome in metastatic colorectal cancer (mCRC) patients (pts) treated with first-line FOLFIRI/Bev: Data from TRIBE and FIRE-3 trials.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 646-646	2.2	
65	Clinical significance of enterocyte-specific gene polymorphisms as candidate marker of oxaliplatin-based treatment for metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 12066-12066	2.2	

64	Genetic variants within the glucocorticoids related genes to predict outcome in patients with metastatic colorectal cancer (mCRC).. <i>Journal of Clinical Oncology</i> , 2018 , 36, 12098-12098	2.2	
63	Molecular characterization of appendiceal cancer and comparison with right-sided (R-CRC) and left-sided colorectal cancer (L-CRC).. <i>Journal of Clinical Oncology</i> , 2018 , 36, 3611-3611	2.2	
62	Comprehensive genomic profiling of 724 gastroenteropancreatic neuroendocrine tumors (GEP-NETs).. <i>Journal of Clinical Oncology</i> , 2018 , 36, 4098-4098	2.2	
61	Genetic variations in the $\beta 2M$ /HLA-E immunomodulatory complex to predict outcomes in metastatic colorectal cancer (mCRC) patients (pts) treated with first line FOLFIRI/Cetuximab: Data from the phase III FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2018 , 36, 12107-12107	2.2	
60	The impact of Th17 cell pathway-related genetic variants in metastatic colorectal cancer patients treated with bevacizumab-based chemotherapy.. <i>Journal of Clinical Oncology</i> , 2018 , 36, e15578-e15578	2.2	
59	CXCL9, CXCL10, CXCL11/CXCR3 axis for immune activation - A target for novel cancer therapy. <i>Cancer Treatment Reviews</i> , 2018 , 63, 40-47	14.4	433
58	Neoadjuvant radiotherapy combined with capecitabine and sorafenib in patients with advanced KRAS-mutated rectal cancer: A phase I/II trial (SAKK 41/08). <i>European Journal of Cancer</i> , 2018 , 89, 82-89	7.5	10
57	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
56	CDX2 in colorectal cancer is an independent prognostic factor and regulated by promoter methylation and histone deacetylation in tumors of the serrated pathway. <i>Clinical Epigenetics</i> , 2018 , 10, 120	7.7	24
55	Histology of Nivolumab-Induced Thyroiditis. <i>Thyroid</i> , 2018 , 28, 1727-1728	6.2	17
54	Increased fibrinogen levels at diagnosis are associated with adverse outcome in patients with acute myeloid leukemia. <i>Hematological Oncology</i> , 2017 , 35, 789-796	1.3	14
53	Comprehensive assessment of tumour budding by cytokeratin staining in colorectal cancer. <i>Histopathology</i> , 2017 , 70, 1044-1051	7.3	20
52	What We Know About Stage II and III Colon Cancer: It's Still Not Enough. <i>Targeted Oncology</i> , 2017 , 12, 265-275	5	21
51	Digital analysis and epigenetic regulation of the signature of rejection in colorectal cancer. <i>Oncolimmunology</i> , 2017 , 6, e1288330	7.2	10
50	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
49	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
48	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
47	Consolidation with autologous stem cell transplantation in first remission is safe and effective in AML patients above 65 years. <i>Leukemia Research</i> , 2017 , 53, 28-34	2.7	14

46	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
45	Colorectal cancer: epigenetic alterations and their clinical implications. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2017 , 1868, 439-448	11.2	35
44	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, 4506-4514 ⁴	6.4	14
43	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
42	Role of genetic polymorphisms in CCL5/CCR5 axis to predict efficacy of regorafenib in patients with refractory metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 596-596	2.2	2
41	Genetic variants of genes in CCL5/CCR5 pathway to predict regorafenib-induced hand-foot skin reaction in patients with refractory metastatic colorectal cancer: A report of ethnic difference.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 615-615	2.2	1
40	Polymorphisms in adipokine-related genes to predict treatment outcomes in patients (pts) with metastatic colorectal cancer (mCRC) treated with bevacizumab-based chemotherapy.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 600-600	2.2	
39	Effect of JAK2 SNP rs2274472 on outcome for mCRC patients treated with first-line FOLFIRI and bevacizumab: Data from FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 595-595	2.2	
38	Effect of polymorphisms of genes encoding regulatory proteins in the coagulation cascade on outcome for mCRC patients treated with FOLFIRI and bevacizumab: Data from FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 601-601	2.2	
37	Genetic variants in CCL5 and CCR5 genes and serum VEGF-A levels to predict efficacy of bevacizumab in metastatic colorectal cancer patients receiving first-line chemotherapy.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 11564-11564	2.2	
36	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
35	Genetic polymorphisms of CCL5 and CCR5 to predict efficacy of cetuximab-based treatment in metastatic colorectal cancer patients depending on primary tumor location.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 3594-3594	2.2	1
34	Genetic variations in semaphorin/neuropilin signaling to predict clinical outcome in patients (pts) with metastatic colorectal cancer (mCRC) receiving bevacizumab-based chemotherapy.. <i>Journal of Clinical Oncology</i> , 2017 , 35, 11608-11608	2.2	
33	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
32	Zevalin and BEAM (Z-BEAM) versus rituximab and BEAM (R-BEAM) conditioning chemotherapy prior to autologous stem cell transplantation in patients with mantle cell lymphoma. <i>Hematological Oncology</i> , 2016 , 34, 133-9	1.3	12
31	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
30	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
29	Females versus males: Clinical features and outcome differences in large molecularly selected cohort of mCRC patients.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3540-3540	2.2	1

28	Genetic variants of ATM and XRCC3 to predict efficacy of TAS-102 in patients with refractory metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3579-3579	2.2	3
27	Genetic variants of hENT-1 to predict efficacy of TAS-102 in patients with refractory metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3580-3580	2.2	2
26	Polymorphisms in toll-like receptor (TLR) genes in the prediction of outcome for cetuximab-based treatment in patients with metastatic colorectal cancer (mCRC).. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3588-3588	2.2	2
25	Genetic variations associated with cancer cachexia pathways to predict survival in metastatic colorectal cancer (mCRC): Results from FIRE-3 and TRIBE.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3590-3590	2.2	1
24	Association of TLR9 polymorphism with overall survival in metastatic colorectal cancer patients treated with FOLFIRI plus bevacizumab enrolled in FIRE3.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 498-498	2.2	1
23	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
22	Serum amyloid A (SAA-1) SNP rs12218 to predict outcome for mCRC patients treated with FOLFIRI and bevacizumab: Data from FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 586-586	2.2	
21	Polymorphisms of genes encoding for vitamin D binding protein and Wnt5a to predict outcome for mCRC patients treated with first-line FOLFIRI and bevacizumab: Data from FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3581-3581	2.2	
20	Genetic variants of Pin1 to predict benefit from irinotecan and oxaliplatin based treatment in patients with metastatic colorectal cancer (mCRC).. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11589-11589	2.2	
19	NOS2 polymorphisms in the prediction of benefit from FOLFIRI plus bevacizumab in mCRC patients enrolled in TRIBE trial.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11597-11597	2.2	
18	Genetic variants of R-spondin genes to predict clinical outcome in mCRC patients (pts) treated with first line FOLFIRI and bevacizumab (FOLFIRI/BEV) in FIRE-3 cohort.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3586-3586	2.2	
17	MKNK1 SNP rs8602 to predict outcome for mCRC patients treated with first-line FOLFIRI and bevacizumab: Data from FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11588-11588	2.2	
16	Genetic variants in immune response genes to predict clinical outcome in mCRC patients treated with FOLFIRI/cetuximab (FIRE-3) or with first line cetuximab-based chemotherapy (JACCRO CC-05/06 AR).. <i>Journal of Clinical Oncology</i> , 2016 , 34, 3595-3595	2.2	
15	Epidermal growth factor receptor mRNA expression in circulating tumor cells as a potential mechanism of molecular escape from regorafenib therapy.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11517-11517	2.2	
14	IRS1 and IRS2 polymorphisms and outcome in mCRC patients enrolled in the FIRE-3 trial.. <i>Journal of Clinical Oncology</i> , 2016 , 34, 11600-11600	2.2	
13	The safety of monoclonal antibodies for treatment of colorectal cancer. <i>Expert Opinion on Drug Safety</i> , 2016 , 15, 799-808	4.1	21
12	Somatic POLE proofreading domain mutation, immune response, and prognosis in colorectal cancer: a retrospective, pooled biomarker study. <i>The Lancet Gastroenterology and Hepatology</i> , 2016 , 1, 207-216	18.8	160
11	Tumor budding in colorectal cancer revisited: results of a multicenter interobserver study. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2015 , 466, 485-93	5.1	73

10	Cytokeratin-20 and Survivin-Expressing Circulating Tumor Cells Predict Survival in Metastatic Colorectal Cancer Patients by a Combined Immunomagnetic qRT-PCR Approach. <i>Molecular Cancer Therapeutics</i> , 2015 , 14, 2401-8	6.1	21
9	CD34+ selected versus unselected autologous stem cell transplantation in patients with advanced-stage mantle cell and diffuse large B-cell lymphoma. <i>Leukemia Research</i> , 2015 , 39, 561-7	2.7	8
8	TWIST1 and TWIST2 promoter methylation and protein expression in tumor stroma influence the epithelial-mesenchymal transition-like tumor budding phenotype in colorectal cancer. <i>Oncotarget</i> , 2015 , 6, 874-85	3.3	48
7	VE1 immunohistochemistry predicts BRAF V600E mutation status and clinical outcome in colorectal cancer. <i>Oncotarget</i> , 2015 , 6, 41453-63	3.3	15
6	Unusual case of progressive multifocal leukoencephalopathy after allogeneic hematopoietic stem-cell transplantation. <i>Journal of Clinical Oncology</i> , 2014 , 32, e33-4	2.2	6
5	CD8/CD45RO T-cell infiltration in endoscopic biopsies of colorectal cancer predicts nodal metastasis and survival. <i>Journal of Translational Medicine</i> , 2014 , 12, 81	8.5	41
4	Neoadjuvant radiotherapy (RT) combined with capecitabine (Cape) and sorafenib (Sor) in patients (pts) with locally advanced, k-ras-mutated rectal cancer (LARC): A phase I/II trial SAKK 41/08.. <i>Journal of Clinical Oncology</i> , 2014 , 32, 3531-3531	2.2	4
3	Investigation of IL-23 (p19, p40) and IL-23R identifies nuclear expression of IL-23 p19 as a favorable prognostic factor in colorectal cancer: a retrospective multicenter study of 675 patients. <i>Oncotarget</i> , 2014 , 5, 4671-82	3.3	10
2	Pitfalls in the diagnosis of intravascular large B-cell lymphoma. <i>European Journal of Haematology</i> , 2013 , 91, 563-4	3.8	1
1	Mucin-producing adenocarcinoma arising in an atrial myxoma. <i>Annals of Diagnostic Pathology</i> , 2013 , 17, 104-7	2.2	6