Fotios Loupakis

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.

329	12,169	54	105
papers	citations	h-index	g-index
361 ext. papers	14,713 ext. citations	5.7 avg, IF	5.93 L-index

#	Paper	IF	Citations
329	Complete pathological response of colorectal peritoneal metastases in Lynch syndrome after immunotherapy case report: is a paradigm shift in cytoreductive surgery needed?. <i>BMC Gastroenterology</i> , 2022 , 22, 17	3	O
328	Systematic review of randomised clinical trials and observational studies for patients with RAS wild-type or BRAF-mutant metastatic and/or unresectable colorectal cancer <i>Critical Reviews in Oncology/Hematology</i> , 2022 , 173, 103646	7	О
327	Prognostic impact of FGFR2/3 alterations in patients with biliary tract cancers receiving systemic chemotherapy: the BITCOIN study <i>European Journal of Cancer</i> , 2022 , 166, 165-175	7.5	1
326	Association of CLDN18 Protein Expression with Clinicopathological Features and Prognosis in Advanced Gastric and Gastroesophageal Junction Adenocarcinomas. <i>Journal of Personalized Medicine</i> , 2021 , 11,	3.6	4
325	RNA-Binding Protein Polymorphisms as Novel Biomarkers to Predict Outcomes of Metastatic Colorectal Cancer: A Meta-analysis from TRIBE, FIRE-3, and MAVERICC. <i>Molecular Cancer Therapeutics</i> , 2021 , 20, 1153-1160	6.1	O
324	NUTM1-rearranged colorectal sarcoma: a clinicopathologically and genetically distinctive malignant neoplasm with a poor prognosis. <i>Modern Pathology</i> , 2021 , 34, 1547-1557	9.8	5
323	RAS as a positive predictive biomarker: focus on lung and colorectal cancer patients. <i>European Journal of Cancer</i> , 2021 , 146, 74-83	7.5	8
322	MicroRNAs as Predictive Biomarkers of Resistance to Targeted Therapies in Gastrointestinal Tumors. <i>Biomedicines</i> , 2021 , 9,	4.8	2
321	Molecular profiling of appendiceal serrated lesions, polyps and mucinous neoplasms: a single-centre experience. <i>Journal of Cancer Research and Clinical Oncology</i> , 2021 , 147, 1897-1904	4.9	1
320	The Role of p53 Expression in Patients with RAS/BRAF Wild-Type Metastatic Colorectal Cancer Receiving Irinotecan and Cetuximab as Later Line Treatment. <i>Targeted Oncology</i> , 2021 , 16, 517-527	5	1
319	Trastuzumab deruxtecan (DS-8201) in patients with HER2-expressing metastatic colorectal cancer (DESTINY-CRC01): a multicentre, open-label, phase 2 trial. <i>Lancet Oncology, The</i> , 2021 , 22, 779-789	21.7	53
318	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
317	Germ line polymorphisms of genes involved in pluripotency transcription factors predict efficacy of cetuximab in metastatic colorectal cancer. <i>European Journal of Cancer</i> , 2021 , 150, 133-142	7.5	1
316	Detection of Molecular Residual Disease Using Personalized Circulating Tumor DNA Assay in Patients With Colorectal Cancer Undergoing Resection of Metastases. <i>JCO Precision Oncology</i> , 2021 , 5,	3.6	9
315	Outcome of patients with colorectal cancer undergoing lung metastases resection: a single-institution retrospective analysis. <i>Tumori</i> , 2021 , 107, 46-54	1.7	2
314	Hurrying up but not rushing, acting and not reacting, good sense and not common sense: Open thoughts and reasonable doubts on COVID-19 vaccination strategies in cancer patients. <i>Critical Reviews in Oncology/Hematology</i> , 2021 , 167, 103271	7	1
313	Encorafenib Plus Cetuximab as a New Standard of Care for Previously Treated V600E-Mutant Metastatic Colorectal Cancer: Updated Survival Results and Subgroup Analyses from the BEACON Study. <i>Journal of Clinical Oncology</i> , 2021 , 39, 273-284	2.2	60

312	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	О
311	Synaptophysin expression in mutated advanced colorectal cancers identifies a new subgroup of tumours with worse prognosis. <i>European Journal of Cancer</i> , 2021 , 146, 145-154	7.5	6
310	Epstein-Barr virus associated gastric dysplasia: a new rare entity?. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 1	5.1	O
309	Efficacy and Safety of Immune Checkpoint Inhibitors in Patients with Microsatellite Instability-High End-Stage Cancers and Poor Performance Status Related to High Disease Burden. <i>Oncologist</i> , 2020 , 25, 803-809	5.7	17
308	The Pan-Immune-Inflammation Value is a new prognostic biomarker in metastatic colorectal cancer: results from a pooled-analysis of the Valentino and TRIBE first-line trials. <i>British Journal of Cancer</i> , 2020 , 123, 403-409	8.7	22
307	Glycolytic competence in gastric adenocarcinomas negatively impacts survival outcomes of patients treated with salvage paclitaxel-ramucirumab. <i>Gastric Cancer</i> , 2020 , 23, 1064-1074	7.6	4
306	Prognostic impact of immune-microenvironment in colorectal liver metastases resected after triplets plus a biologic agent: A pooled analysis of five prospective trials. <i>European Journal of Cancer</i> , 2020 , 135, 78-88	7.5	5
305	Combination of variations in inflammation- and endoplasmic reticulum-associated genes as putative biomarker for bevacizumab response in KRAS wild-type colorectal cancer. <i>Scientific Reports</i> , 2020 , 10, 9778	4.9	1
304	KRAS G12C Metastatic Colorectal Cancer: Specific Features of a New Emerging Target Population. <i>Clinical Colorectal Cancer</i> , 2020 , 19, 219-225	3.8	16
303	Upfront FOLFOXIRI plus bevacizumab and reintroduction after progression versus mFOLFOX6 plus bevacizumab followed by FOLFIRI plus bevacizumab in the treatment of patients with metastatic colorectal cancer (TRIBE2): a multicentre, open-label, phase 3, randomised, controlled trial. <i>Lancet</i>	21.7	98
302	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
301	Prediction of Benefit from Checkpoint Inhibitors in Mismatch Repair Deficient Metastatic Colorectal Cancer: Role of Tumor Infiltrating Lymphocytes. <i>Oncologist</i> , 2020 , 25, 481-487	5.7	43
300	The heterogeneous clinical and pathological landscapes of metastatic -mutated colorectal cancer. <i>Cancer Cell International</i> , 2020 , 20, 30	6.4	22
299	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
298	Encorafenib plus cetuximab with or without binimetinib for BRAF V600E metastatic colorectal cancer: Updated survival results from a randomized, three-arm, phase III study versus choice of either irinotecan or FOLFIRI plus cetuximab (BEACON CRC) <i>Journal of Clinical Oncology</i> , 2020 , 38, 4001	2.2 -4001	22
297	Thyroid hormones ratio is a major prognostic marker in advanced metastatic colorectal cancer: Results from the phase III randomised CORRECT trial. <i>European Journal of Cancer</i> , 2020 , 133, 66-73	7.5	4
296	PD-L1 expression in gastroesophageal dysplastic lesions. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2020 , 477, 151-156	5.1	10
295	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

294	Practical considerations in the use of regorafenib in metastatic colorectal cancer. <i>Therapeutic Advances in Medical Oncology</i> , 2020 , 12, 1758835920956862	5.4	6
293	Immunogenic cell death pathway polymorphisms for predicting oxaliplatin efficacy in metastatic colorectal cancer 2020 , 8,		4
292	Mutation Analysis in Gastric Cancer and Clinical Outcomes of Patients with Metastatic Disease Treated with Ramucirumab/Paclitaxel or Standard Chemotherapy. <i>Cancers</i> , 2020 , 12,	6.6	3
291	Impact of Pre-Analytical Factors on MSI Test Accuracy in Mucinous Colorectal Adenocarcinoma: A Multi-Assay Concordance Study. <i>Cells</i> , 2020 , 9,	7.9	7
290	Investigating the concordance in molecular subtypes of primary colorectal tumors and their matched synchronous liver metastasis. <i>International Journal of Cancer</i> , 2020 , 147, 2303-2315	7.5	7
289	Retreatment With Anti-EGFR Antibodies in Metastatic Colorectal Cancer Patients: A Multi-institutional Analysis. <i>Clinical Colorectal Cancer</i> , 2020 , 19, 191-199.e6	3.8	10
288	CK7 and consensus molecular subtypes as major prognosticators in BRAF mutated metastatic colorectal cancer. <i>British Journal of Cancer</i> , 2019 , 121, 593-599	8.7	14
287	Negative Hyperselection of Patients With and Wild-Type Metastatic Colorectal Cancer Who Received Panitumumab-Based Maintenance Therapy. <i>Journal of Clinical Oncology</i> , 2019 , 37, 3099-3110	2.2	35
286	Relationship Between Tumor Response and Tumor-Related Symptoms in RAS Wild-Type Metastatic Colorectal Cancer: Retrospective Analyses From 3 Panitumumab Trials. <i>Clinical Colorectal Cancer</i> , 2019 , 18, 245-256.e5	3.8	0
285	Encorafenib, Binimetinib, and Cetuximab in V600E-Mutated Colorectal Cancer. <i>New England Journal of Medicine</i> , 2019 , 381, 1632-1643	59.2	481
284	Early modifications of circulating microRNAs levels in metastatic colorectal cancer patients treated with regorafenib. <i>Pharmacogenomics Journal</i> , 2019 , 19, 455-464	3.5	3
283	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
282	An overview on clinical, pathological and molecular features of lung metastases from colorectal cancer. <i>Expert Review of Respiratory Medicine</i> , 2019 , 13, 635-644	3.8	2
281	Quantitative evidence for early metastatic seeding in colorectal cancer. <i>Nature Genetics</i> , 2019 , 51, 1113	-3632	164
280	Claudin-18 expression in oesophagogastric adenocarcinomas: a tissue microarray study of 523 molecularly profiled cases. <i>British Journal of Cancer</i> , 2019 , 121, 257-263	8.7	20
279	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
278	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
277	Class 1, 2, and 3 -Mutated Metastatic Colorectal Cancer: A Detailed Clinical, Pathologic, and Molecular Characterization. <i>Clinical Cancer Research</i> , 2019 , 25, 3954-3961	12.9	36

276	Benefit from anti-EGFRs in and wild-type metastatic transverse colon cancer: a clinical and molecular proof of concept study. <i>ESMO Open</i> , 2019 , 4, e000489	6	8
275	Ramucirumab: the long and winding road toward being an option for mCRC treatment. <i>Expert Opinion on Biological Therapy</i> , 2019 , 19, 399-409	5.4	3
274	Chemotherapeutic and antiangiogenic drugs beyond tumor progression in colon cancer: Evaluation of the effects of switched schedules and related pharmacodynamics. <i>Biochemical Pharmacology</i> , 2019 , 164, 94-105	6	9
273	DPYD*6 plays an important role in fluoropyrimidine toxicity in addition to DPYD*2A and c.2846A>T: a comprehensive analysis in 1254 patients. <i>Pharmacogenomics Journal</i> , 2019 , 19, 556-563	3.5	20
272	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 146-157	9.7	67
271	Tumor mutation burden: from comprehensive mutational screening to the clinic. <i>Cancer Cell International</i> , 2019 , 19, 209	6.4	74
270	Pathological Tumor Regression Grade Classifications in Gastrointestinal Cancers: Role on Patients' Prognosis. <i>International Journal of Surgical Pathology</i> , 2019 , 27, 816-835	1.2	4
269	A validated prognostic classifier for BRAF-mutated metastatic colorectal cancer: the 'BRAF BeCool' study. <i>European Journal of Cancer</i> , 2019 , 118, 121-130	7.5	29
268	Aryl hydrocarbon receptor nuclear translocator-like (ARNTL/BMAL1) is associated with bevacizumab resistance in colorectal cancer via regulation of vascular endothelial growth factor A. <i>EBioMedicine</i> , 2019 , 45, 139-154	8.8	19
267	High Circulating Methylated DNA Is a Negative Predictive and Prognostic Marker in Metastatic Colorectal Cancer Patients Treated With Regorafenib. <i>Frontiers in Oncology</i> , 2019 , 9, 622	5.3	17
266	Treatment with checkpoint inhibitors in a metastatic colorectal cancer patient with molecular and immunohistochemical heterogeneity in MSI/dMMR status 2019 , 7, 297		10
265	PTEN in Colorectal Cancer: Shedding Light on Its Role as Predictor and Target. <i>Cancers</i> , 2019 , 11,	6.6	30
264	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
263	Targeted therapies in metastatic gastric cancer: Current knowledge and future perspectives. <i>World Journal of Gastroenterology</i> , 2019 , 25, 5773-5788	5.6	45
262	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	
261	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	
260	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	
259	Genetic variants in RNA binding protein (RBP) to predict outcome in metastatic colorectal cancer (mCRC): Data from FIRE-3, TRIBE, and MAVERICC trials <i>Journal of Clinical Oncology</i> , 2019 , 37, 3545-35	545 ^{2.2}	

258	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	О
257	Another Chapter of the : Is Primary Tumor Location a Prognostic Feature in Mutant Metastatic Colorectal Cancer?. <i>Oncologist</i> , 2019 , 24, e77-e79	5.7	3
256	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
255	miR-224 Is Significantly Upregulated and Targets Caspase-3 and Caspase-7 During Colorectal Carcinogenesis. <i>Translational Oncology</i> , 2019 , 12, 282-291	4.9	10
254	Safety and Tolerability of Anti-Angiogenic Protein Kinase Inhibitors and Vascular-Disrupting Agents in Cancer: Focus on Gastrointestinal Malignancies. <i>Drug Safety</i> , 2019 , 42, 159-179	5.1	13
253	Safety and Tolerability of c-MET Inhibitors in Cancer. <i>Drug Safety</i> , 2019 , 42, 211-233	5.1	40
252	Pharmacokinetic analysis of metronomic capecitabine in refractory metastatic colorectal cancer patients. <i>Investigational New Drugs</i> , 2018 , 36, 709-714	4.3	5
251	Primary tumor sidedness and benefit from FOLFOXIRI plus bevacizumab as initial therapy for metastatic colorectal cancer. Retrospective analysis of the TRIBE trial by GONO. <i>Annals of Oncology</i> , 2018 , 29, 1528-1534	10.3	58
250	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
249	The role of tumor angiogenesis as a therapeutic target in colorectal cancer. <i>Expert Review of Anticancer Therapy</i> , 2018 , 18, 251-266	3.5	29
248	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
247	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
246	The PANDA study: a randomized phase II study of first-line FOLFOX plus panitumumab versus 5FU plus panitumumab in RAS and BRAF wild-type elderly metastatic colorectal cancer patients. <i>BMC Cancer</i> , 2018 , 18, 98	4.8	6
245	Noninferiority of three months versus six months of oxaliplatin-based adjuvant chemotherapy for resected colon cancer. How should IDEA findings affect clinical practice?. <i>International Journal of Cancer</i> , 2018 , 143, 2342-2350	7.5	5
244	Prognostic Value of Thyroid Hormone Ratios in Patients With Advanced Metastatic Colorectal Cancer Treated With Regorafenib: The TOREADOR Study. <i>Clinical Colorectal Cancer</i> , 2018 , 17, e601-e61	5 ^{3.8}	8
243	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
242	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
241	Clinico-pathological and molecular characterisation of BRAF mutant metastatic colorectal cancer (mCRC): Are all mutations created equal?. <i>Journal of Clinical Oncology</i> , 2018 , 36, 3590-3590	2.2	4

2 40	Clinical prognostic score of BRAF V600E mutated (BM) metastatic colorectal cancer (mCRC): Results from the B RAF, BeCooliplatform <i>Journal of Clinical Oncology</i> , 2018 , 36, 639-639	2.2	2
239	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
238	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
237	Efficacy outcomes with bevacizumab added to chemotherapy (bev+CT) compared with chemotherapy alone (CT) in left- and right-sided tumors in metastatic colorectal cancer (mCRC) <i>Journal of Clinical Oncology</i> , 2018 , 36, 726-726	2.2	3
236	and genotyping to predict adverse events during first-line FOLFIRI or FOLFOXIRI plus bevacizumab in metastatic colorectal cancer. <i>Oncotarget</i> , 2018 , 9, 7859-7866	3.3	18
235	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	
234	Single nucleotide polymorphisms in miRNA binding sites of nucleotide excision repair-related genes to predict clinical benefit of oxaliplatin in FOLFOXIRI plus bevacizumab in TRIBE trial <i>Journal of Clinical Oncology</i> , 2018 , 36, 663-663	2.2	
233	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	
232	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	
231	Histopathologic response and growth patterns of colorectal cancer liver metastases (CRCLM) in patients treated with triplets plus bevacizumab (bev) or anti-EGFRs <i>Journal of Clinical Oncology</i> , 2018 , 36, 636-636	2.2	
230	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, 120	0 66 -12	066
229	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	
228	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	
227	Genetic variation in TET3 and survival in metastatic colorectal cancer (mCRC) from FIRE-3, TRIBE, and MAVERICC clinical trials <i>Journal of Clinical Oncology</i> , 2018 , 36, 3575-3575	2.2	
226	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
225	Loss of Chromosome 18q11.2-q12.1 Is Predictive for Survival in Patients With Metastatic Colorectal Cancer Treated With Bevacizumab. <i>Journal of Clinical Oncology</i> , 2018 , 36, 2052-2060	2.2	19
224	The DISTINCTIVE study: A biologically enriched phase II study of seconD-line folfiri/aflibercept in proSpecTively stratified, anti-EGFR resistaNt, metastatic coloreCTal cancer patients with RAS Validated wild typE status - Trial in progress. <i>Annals of Oncology</i> , 2018 , 29, v82	10.3	3
223	Impact of primary tumour location on efficacy of bevacizumab plus chemotherapy in metastatic colorectal cancer. <i>British Journal of Cancer</i> , 2018 , 119, 1451-1455	8.7	14

222	Copy number load predicts outcome of metastatic colorectal cancer patients receiving bevacizumab combination therapy. <i>Nature Communications</i> , 2018 , 9, 4112	17.4	36
221	LONG-NONCODING RNAs in gastroesophageal cancers. <i>Non-coding RNA Research</i> , 2018 , 3, 195-212	6	28
220	Assessment of intratumor immune-microenvironment in colorectal cancers with extranodal extension of nodal metastases. <i>Cancer Cell International</i> , 2018 , 18, 131	6.4	6
219	Precision medicine in cholangiocarcinoma. <i>Translational Gastroenterology and Hepatology</i> , 2018 , 3, 40	5.2	39
218	Glycolysis gene expression analysis and selective metabolic advantage in the clinical progression of colorectal cancer. <i>Pharmacogenomics Journal</i> , 2017 , 17, 258-264	3.5	55
217	Variant alleles in factor V, prothrombin, plasminogen activator inhibitor-1, methylenetetrahydrofolate reductase and risk of thromboembolism in metastatic colorectal cancer patients treated with first-line chemotherapy plus bevacizumab. <i>Pharmacogenomics Journal</i> , 2017 ,	3.5	6
216	Serum LDH predicts benefit from bevacizumab beyond progression in metastatic colorectal cancer. British Journal of Cancer, 2017 , 116, 318-323	8.7	20
215	The role of pharmacogenetics in the new ESMO colorectal cancer guidelines. <i>Pharmacogenomics</i> , 2017 , 18, 197-200	2.6	7
214	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
213	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
212	Immunotherapy for colorectal cancer: where are we heading?. <i>Expert Opinion on Biological Therapy</i> , 2017 , 17, 709-721	5.4	71
211	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
210	Homeobox B9 Mediates Resistance to Anti-VEGF Therapy in Colorectal Cancer Patients. <i>Clinical Cancer Research</i> , 2017 , 23, 4312-4322	12.9	27
209	Efficacy of FOLFOXIRI plus bevacizumab in liver-limited metastatic colorectal cancer: A pooled analysis of clinical studies by Gruppo Oncologico del Nord Ovest. <i>European Journal of Cancer</i> , 2017 , 73, 74-84	7.5	32
208	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
207	QoL is a cool tool. <i>Annals of Oncology</i> , 2017 , 28, 2032-2033	10.3	2
206	BRAF p.V600E-specific immunohistochemical assessment in colorectal cancer endoscopy biopsies is consistent with the mutational profiling. <i>Histopathology</i> , 2017 , 71, 1008-1011	7.3	8
205	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, 450	06 -4 51	44

(2016-2017)

204	Anti-EGFR monoclonal antibody panitumumab for the treatment of patients with metastatic colorectal cancer: an overview of current practice and future perspectives. <i>Expert Opinion on Biological Therapy</i> , 2017 , 17, 1297-1308	5.4	17	
203	New perspectives for TAS-102: TASK successful?. <i>Lancet Oncology, The</i> , 2017 , 18, 1139-1141	21.7	1	
202	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	
201	Vinorelbine in BRAF V600E mutated metastatic colorectal cancer: a prospective multicentre phase II clinical study. <i>ESMO Open</i> , 2017 , 2, e000241	6	6	
200	TRIBE-2: a phase III, randomized, open-label, strategy trial in unresectable metastatic colorectal cancer patients by the GONO group. <i>BMC Cancer</i> , 2017 , 17, 408	4.8	20	
199	Estimating 12-week death probability in patients with refractory metastatic colorectal cancer: the Colon Life nomogram. <i>Annals of Oncology</i> , 2017 , 28, 555-561	10.3	32	
198	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	o ^{3.5}	9	
197	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	
196	Stereotactic Body Radiotherapy in Patients with Lung Oligometastases from Colorectal Cancer. <i>Anticancer Research</i> , 2017 , 37, 315-319	2.3	16	
195	Surrogate Endpoints in Second-Line Trials of Targeted Agents in Metastatic Colorectal Cancer: A Literature-Based Systematic Review and Meta-Analysis. <i>Cancer Research and Treatment</i> , 2017 , 49, 834-8	345 ²	9	
194	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		
193	Partial splenic embolization in chemotherapy-induced thrombocytopenia: A retrospective analysis with long term follow-up <i>Journal of Clinical Oncology</i> , 2017 , 35, e21654-e21654	2.2		
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169	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	

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