

Consulting physician		Patient		Sample	
Provider	General Hospital	Age	64	Accession Number	TSO500_CRC_MSI_TMB
Physician	Dr. E Schmidt	Gender	Male	Collection site	Colon
Pathologist	Dr. R Braun	Diagnosis	Colorectal Carcinoma	Type	Biopsy
Report Date	08.04.2021	Stage	IV	Collection date	08.04.2021

Panel Analysis: TSO500

The TruSight Oncology 500 panel enables in-house, pan-cancer comprehensive genomic profiling (CGP) for solid tumors from either blood or tissue biopsy samples; the panel is designed to identify relevant biomarkers in guidelines and trials, including the immuno-oncology markers TMB and MSI.

Analysis results: Positive

2 Biomarkers	Approved treatments	Other findings
Tumor Mutation Burden: TMB-high (37.2 Mutations/Megabase)	Pembrolizumab	Other Indications: ipilimumab /nivolumab, nivolumab Trials: 1 Phase 2
Microsatellite Status: MSI-high	Durvalumab Ipilimumab/nivolumab Nivolumab Pembrolizumab	Trials: 3 Phase 2 2 Phase 1
3 Variants of potential clinical significance, Tier 2	Approved treatments	Other findings
ARID1A: p.H688fs*129, Likely Pathogenic	-	-
MSH2: p.N538fs*5, Likely Pathogenic	-	-
MSH2: p.N835fs*4, Pathogenic	-	-
6 Variants of uncertain significance, Tier 3		

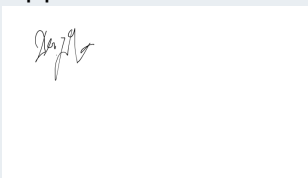
Interactions

None

Guidelines

Potentially relevant guidelines are reported in the "guidelines" section starting on page 2.

Approval



Electronically signed on: 08.04.2021 by Dr. Schmidt

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GUIDELINES

The ESMO Clinical Practice Guidelines suggest testing for microsatellite instability (MSI) in all newly diagnosed patients with colon cancer and note that the presence of MSI may predict sensitivity to immune check point inhibitors; the Guidelines additionally note that stage 2 MSI-high colorectal carcinoma patients may have a good prognosis and may not benefit from adjuvant 5FU therapy [PMID:32702383, PMID:27380959].

TREATMENT OPTIONS

Therapies with potential clinical benefit (6)

IPILIMUMAB/NIVOLUMAB

Nivolumab, a PD-1 blocking antibody, in combination with ipilimumab, a CTLA-4 blocking antibody, is FDA- and EMA-approved for treating patients with unresectable or metastatic melanoma; for treating adult patients with metastatic or recurrent non-small cell lung cancer with no EGFR or ALK genomic tumor aberrations as first-line treatment, in combination with 2 cycles of platinum-doublet chemotherapy; and for treating patients with intermediate or poor risk advanced renal cell carcinoma, as a first-line treatment; nivolumab, in combination with ipilimumab, is FDA-approved for treating adult patients with metastatic non-small cell lung cancer expressing PD-L1 ($\geq 1\%$) as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, as first-line treatment; for treating adult patients with unresectable malignant pleural mesothelioma, as first-line treatment; for treating adult and pediatric (12 years and older) patients with microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) metastatic colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan; and for treating patients with hepatocellular carcinoma who have been previously treated with sorafenib.

Sensitive

Biomarkers: Tumor Mutation Burden: TMB-high, Tier 1A
Microsatellite Status: MSI-high, Tier 1A

IPILIMUMAB/NIVOLUMAB

Nivolumab, a PD-1 blocking antibody, in combination with ipilimumab, a CTLA-4 blocking antibody, is FDA- and EMA-approved for treating patients with unresectable or metastatic melanoma; for treating adult patients with metastatic or recurrent non-small cell lung cancer with no EGFR or ALK genomic tumor aberrations as first-line treatment, in combination with 2 cycles of platinum-doublet chemotherapy; and for treating patients with intermediate or poor risk advanced renal cell carcinoma, as a first-line treatment; nivolumab, in combination with ipilimumab, is FDA-approved for treating adult patients with metastatic non-small cell lung cancer expressing PD-L1 ($\geq 1\%$) as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, as first-line treatment; for treating adult patients with unresectable malignant pleural mesothelioma, as first-line treatment; for treating adult and pediatric (12 years and older) patients with microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) metastatic colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan; and for treating patients with hepatocellular carcinoma who have been previously treated with sorafenib.

Sensitive

Biomarkers: Tumor Mutation Burden: TMB-high, Tier 1A
Microsatellite Status: MSI-high, Tier 1A

NIVOLUMAB

Nivolumab, a PD-1 blocking antibody, is FDA- and EMA-approved for treating patients with unresectable or metastatic melanoma, as a single agent or in combination with ipilimumab; melanoma with lymph node involvement or metastatic disease who have undergone complete resection, in the adjuvant setting; for treating adult patients with metastatic or recurrent non-small cell lung cancer with no EGFR or ALK genomic tumor aberrations as first-line treatment, in combination with ipilimumab and 2 cycles of platinum-doublet chemotherapy; for treating patients with metastatic non-small cell lung cancer with progression on or after platinum-based chemotherapy (patients with EGFR or ALK genomic tumor aberrations should have disease progression on FDA-approved therapy for these aberrations prior to receiving nivolumab); intermediate or poor risk advanced renal cell carcinoma, as a first-line treatment in combination with ipilimumab; advanced renal cell carcinoma who have received prior antiangiogenic therapy; for treating adult patients with classical Hodgkin lymphoma that has relapsed or progressed after autologous hematopoietic stem cell transplantation (HSCT) and brentuximab vedotin, or has relapsed or progressed after 3 or more lines of systemic therapy that includes autologous HSCT; for treating patients with recurrent or metastatic squamous cell carcinoma of the head and neck with disease progression on or after a platinum-based therapy; locally advanced or metastatic urothelial carcinoma who have disease progression during or following platinum-containing chemotherapy or have disease progression within 12 months of neoadjuvant or adjuvant treatment with platinum-containing chemotherapy; and unresectable advanced, recurrent or metastatic esophageal squamous cell carcinoma after prior fluoropyrimidine- and platinum-based chemotherapy; nivolumab is also FDA-approved for treating adult patients with metastatic non-small cell lung cancer expressing PD-L1 ($\geq 1\%$) as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, as first-line treatment in combination with ipilimumab; unresectable malignant pleural mesothelioma, as first-line treatment in combination with ipilimumab; for treating patients with advanced renal cell carcinoma, as a first-line treatment in combination with cabozantinib; for treating adult and pediatric (12 years and older) patients with microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) metastatic colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan, as a single agent or in combination with ipilimumab; and for treating patients with hepatocellular carcinoma who have been previously treated with sorafenib, as a single agent or in combination with ipilimumab.

Sensitive

Biomarkers: Tumor Mutation Burden: TMB-high, Tier 1A
Microsatellite Status: MSI-high, Tier 1A

NIVOLUMAB

Nivolumab, a PD-1 blocking antibody, is FDA- and EMA-approved for treating patients with unresectable or metastatic melanoma, as a single agent or in combination with ipilimumab; melanoma with lymph node involvement or metastatic disease who have undergone complete resection, in the adjuvant setting; for treating adult patients with metastatic or recurrent non-small cell lung cancer with no EGFR or ALK genomic tumor aberrations as first-line treatment, in combination with ipilimumab and 2 cycles of platinum-doublet chemotherapy; for treating patients with metastatic non-small cell lung cancer with progression on or after platinum-based chemotherapy (patients with EGFR or ALK genomic tumor aberrations should have disease progression on FDA-approved therapy for these aberrations prior to receiving nivolumab); intermediate or poor risk advanced renal cell carcinoma, as

Therapies with potential clinical benefit (6)

a first-line treatment in combination with ipilimumab; advanced renal cell carcinoma who have received prior antiangiogenic therapy; for treating adult patients with classical Hodgkin lymphoma that has relapsed or progressed after autologous hematopoietic stem cell transplantation (HSCT) and brentuximab vedotin, or has relapsed or progressed after 3 or more lines of systemic therapy that includes autologous HSCT; for treating patients with recurrent or metastatic squamous cell carcinoma of the head and neck with disease progression on or after a platinum-based therapy; locally advanced or metastatic urothelial carcinoma who have disease progression during or following platinum-containing chemotherapy or have disease progression within 12 months of neoadjuvant or adjuvant treatment with platinum-containing chemotherapy; and unresectable advanced, recurrent or metastatic esophageal squamous cell carcinoma after prior fluoropyrimidine- and platinum-based chemotherapy; nivolumab is also FDA-approved for treating adult patients with metastatic non-small cell lung cancer expressing PD-L1 ($\geq 1\%$) as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, as first-line treatment in combination with ipilimumab; unresectable malignant pleural mesothelioma, as first-line treatment in combination with ipilimumab; for treating patients with advanced renal cell carcinoma, as a first-line treatment in combination with cabozantinib; for treating adult and pediatric (12 years and older) patients with microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) metastatic colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan, as a single agent or in combination with ipilimumab; and for treating patients with hepatocellular carcinoma who have been previously treated with sorafenib, as a single agent or in combination with ipilimumab.

Sensitive

Biomarkers: Tumor Mutation Burden: TMB-high, Tier 1A
 Microsatellite Status: MSI-high, Tier 1A

PEMBROLIZUMAB

Pembrolizumab, a programmed death receptor-1 (PD-1)-blocking antibody, is FDA- and EMA-approved for treating patients with unresectable or metastatic melanoma; for the adjuvant treatment of patients with melanoma with involvement of lymph node(s) following complete resection; in combination with pemetrexed and platinum chemotherapy for the first-line treatment of patients with metastatic nonsquamous non-small cell lung cancer (NSCLC), with no EGFR or ALK genomic tumor aberrations; in combination with carboplatin and either paclitaxel or paclitaxel protein-bound, for the first-line treatment of patients with metastatic squamous NSCLC; as a single agent for treating patients with metastatic NSCLC whose tumors express PD-L1 (TPS $\geq 1\%$) as determined by an FDA-approved test, with disease progression on or after platinum-containing chemotherapy (patients with EGFR or ALK genomic tumor aberrations should have disease progression on FDA-approved therapy for these aberrations prior to receiving pembrolizumab); for treating patients with locally advanced or metastatic urothelial carcinoma who are not eligible for cisplatin-containing chemotherapy and whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 10] as determined by an FDA-approved test; and, in combination with axitinib, for the first-line treatment of patients with advanced renal cell carcinoma; pembrolizumab is also FDA-approved as a single agent for the first-line treatment of patients with NSCLC expressing PD-L1 [Tumor Proportion Score (TPS) $\geq 1\%$] as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, and is stage III where patients are not candidates for surgical resection or definitive chemoradiation, or metastatic; for treating patients with metastatic small cell lung cancer (SCLC) with disease progression on or after platinum-based chemotherapy and at least one other prior line of therapy; in combination with platinum and FU for the first-line treatment of patients with metastatic or with unresectable, recurrent head and neck squamous cell cancer (HNSCC); as a single agent for the first-line treatment of patients with metastatic or with unresectable, recurrent HNSCC whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 1] as determined by an FDA-approved test; as a single agent for treating patients with recurrent or metastatic HNSCC with disease progression on or after platinum-containing chemotherapy; for treating adult patients with relapsed or refractory classical Hodgkin lymphoma (cHL); for treating pediatric patients with refractory cHL, or cHL that has relapsed after 2 or more lines of therapy; for treating adult and pediatric patients with refractory primary mediastinal large B-cell lymphoma (PMBCL), or who have relapsed after 2 or more prior lines of therapy; for treating patients with locally advanced or metastatic urothelial carcinoma who are not eligible for any platinum-containing chemotherapy regardless of PD-L1 status; for treating patients with locally advanced or metastatic urothelial carcinoma who have disease progression during or following platinum-containing chemotherapy or within 12 months of neoadjuvant or adjuvant treatment with platinum-containing chemotherapy; for treating patients with Bacillus Calmette-Guerin (BCG)-unresponsive, high-risk, non-muscle invasive bladder cancer (NMIBC) with carcinoma in situ (CIS) with or without papillary tumors who are ineligible for or have elected not to undergo cystectomy; for treating adult and pediatric patients with unresectable or metastatic, microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) solid tumors that have progressed following prior treatment and who have no satisfactory alternative treatment options, or colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan; for the first-line treatment of patients with unresectable or metastatic microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) colorectal cancer (CRC); for treating patients with recurrent locally advanced or metastatic gastric or gastroesophageal junction adenocarcinoma whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 1] as determined by an FDA-approved test, with disease progression on or after two or more prior lines of therapy including fluoropyrimidine-and platinum-containing chemotherapy and if appropriate, HER2/neu-targeted therapy; for treating patients with recurrent locally advanced or metastatic squamous cell carcinoma of the esophagus whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 10] as determined by an FDA-approved test, with disease progression after one or more prior lines of systemic therapy; for treating patients with recurrent or metastatic cervical cancer with disease progression on or after chemotherapy whose tumors express PD-L1 (CPS ≥ 1) as determined by an FDA-approved test; for treating patients with hepatocellular carcinoma who have been previously treated with sorafenib; for treating adult and pediatric patients with recurrent locally advanced or metastatic Merkel cell carcinoma; in combination with lenvatinib, for treating patients with advanced endometrial carcinoma that is not microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR), who have disease progression following prior systemic therapy and are not candidates for curative surgery or radiation; as a single agent for treating adult and pediatric patients with unresectable or metastatic tumor mutational burden-high (TMB-H) [≥ 10 mutations/megabase (mut/Mb)] solid tumors, as determined by an FDA-approved test, that have progressed following prior treatment and who have no satisfactory alternative treatment options; for treating patients with recurrent or metastatic cutaneous squamous cell carcinoma that is not curable by surgery or radiation; and in combination with chemotherapy, for the treatment of patients with locally recurrent unresectable or metastatic Triple-Negative Breast Cancer (TNBC) whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 10] as determined by an FDA-approved test; pembrolizumab is also EMA-approved for treating adult patients with relapsed or refractory classical Hodgkin lymphoma (cHL) who have failed autologous stem cell transplant (ASCT) and brentuximab vedotin (BV), or who are transplant-ineligible and have failed BV; locally advanced or metastatic urothelial carcinoma who have received prior platinum-containing chemotherapy; as monotherapy or in combination with platinum and 5-fluorouracil (5-FU) chemotherapy for the first-line treatment of adult patients with metastatic or unresectable recurrent head and neck squamous cell carcinoma (HNSCC) whose tumours express PD-L1 with a CPS ≥ 1 ; for treating adult patients with recurrent or metastatic HNSCC whose tumours express PD-L1 with a $\geq 50\%$ TPS and progressing on or after platinum-containing chemotherapy; and as a single agent for the first-line treatment of adult patients with metastatic non-small cell lung carcinoma (NSCLC) whose tumours express PD-L1 with a $\geq 50\%$ tumour proportion score (TPS) with no EGFR or ALK positive tumour mutations.

Therapies with potential clinical benefit (6)

Sensitive

Biomarkers: Tumor Mutation Burden: TMB-high, Tier 1A
 Microsatellite Status: MSI-high, Tier 1A

DURVALUMAB

Durvalumab, a programmed death-ligand 1 (PD-L1) blocking antibody, is FDA-approved for treating adult patients with locally advanced or metastatic urothelial carcinoma who have disease progression during or following platinum-containing chemotherapy, or who have disease progression within 12 months of neoadjuvant or adjuvant treatment with platinum-containing chemotherapy; for treating patients with unresectable, stage III non-small cell lung cancer (NSCLC) whose disease has not progressed following concurrent platinum-based chemotherapy and radiation therapy; and, in combination with etoposide and either carboplatin or cisplatin, as first-line treatment for adult patients with extensive-stage small cell lung cancer (ES-SCLC); durvalumab is EMA-approved for treating adult patients with locally advanced, unresectable non-small cell lung cancer (NSCLC) whose tumours express PD-L1 on $\geq 1\%$ of tumour cells and whose disease has not progressed following platinum-based chemoradiation therapy.

Sensitive

Biomarker: Microsatellite Status: MSI-high, Tier 1A

AVAILABLE CLINICAL TRIALS

Phase 2 clinical trials (3)

IPILIMUMAB, ATEZOLIZUMAB, NIVOLUMAB

The Rome Trial From Histology to Target: the Road to Personalize Target Therapy and Immunotherapy

[NCT04591431](#)

Qualifying variants

Biomarker	Classification	Score
TMB-high	Tier 1A Pathogenic	37.2 Mutations/Megabase
MSI-high	Tier 1A Pathogenic	-

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ATEZOLIZUMAB

A Phase II Study to Assess the Efficacy of the Anti-PD-L1 Antibody Atezolizumab (MPDL3280A) Administered With Stereotactic Ablative Radiotherapy (SABR) in Patients With Metastatic Tumours

[NCT02992912](#)

Qualifying variant

Biomarker	Classification	Score
MSI-high	Tier 1A Pathogenic	-

Contact

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CETUXIMAB, BEVACIZUMAB, AFLIBERCEPT, AVELUMAB, 5-FLUOROURACIL/IRINOTECAN/LEUCOVORIN, 5-FLUOROURACIL/LEUCOVORIN/OXALIPLATIN, PANITUMUMAB

Multicenter Randomized Phase II Study Comparing the Effectiveness and Tolerance of Avelumab Versus Standard 2nd Line Treatment Chemotherapy in Patients With Colorectal Metastatic Cancer With Microsatellite Instability (MSI)

[NCT03186326](#)

Qualifying variant

Biomarker	Classification	Score
MSI-high	Tier 1A Pathogenic	-

Contact

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Phase 1 clinical trials (2)

IPILIMUMAB, PEXASTIMOGENE DEVACIREPVEC

A Phase I Dose Escalation Trial Evaluating the Impact of an in Situ Immunization Strategy With Intra-Tumoral Injections of Pexa-Vec in Combination With Ipilimumab in Metastatic / Advanced Solid Tumors With Injectable Lesions.

[NCT02977156](#)

Qualifying variant

Biomarker	Classification	Score
MSI-high	Tier 1A Pathogenic	-

Contact

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MP0310

A First-In-Human, Single-Arm, Multi-Center, Open-Label, Repeated-Dose, Dose-Escalation Study of MP0310 in Patients With Advanced Solid Tumors

[NCT04049903](#)

Qualifying variant

Biomarker	Classification	Score
MSI-high	Tier 1A Pathogenic	-

Contact

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VARIANT DETAILS

Biomarkers (2)

Tumor Mutation Burden: TMB-high (37.2 Mutations/Megabase)

Biomarker: TMB-high

Classification: Tier 1A

Assessment: Pathogenic

Treatment options

3 Sensitive

1 Trial

Biomarker summary: Tumor Mutational Burden-high is an activating mutation.

Clinical relevance: Deregulation of multiple cellular processes is capable of introducing DNA alterations during tumorigenesis. Genetic mutations in tumor cells have been reported to result in the production of neoantigens, which are immunogenic peptides recognized by tumor-infiltrating lymphocytes (TILs) [106, 6, 153, 58]. Studies have shown high tumor mutational burden or high levels of neoantigens to be associated with high expression of cytotoxic T-cell markers; thus, immunotherapies may be relevant in tumors with high tumor mutational burden [72, 58, 17, 143]. Indeed, high tumor mutational burden has been associated with increased clinical benefit of several immune checkpoint inhibitors, including pembrolizumab, nivolumab, nivolumab plus ipilimumab, and atezolizumab in studies of NSCLC, urothelial carcinoma, and other solid tumors [141, 20, 54, 140, 52, 71, 197, 144, 80, 60, 148].

Disease summary: High mutational burden has been associated with microsatellite instability (MSI) and mismatch-repair deficiency (MMR-D) in studies of colorectal carcinoma [165, 105, 97, 21, 49, 98].

Molecular function: A test result demonstrating high tumor mutational burden has been reported in this sample.

Incidence: Hypermutation has been reported in 2-18% of colorectal carcinoma (CRC) samples analyzed in literature studies [21, 43]. In a study of 246 left-sided and 56 right-sided colorectal tumors in young adults and adolescents, TMB-high status was found more often in right-sided tumors, regardless of MSI status [146].

Role in disease: Multiple mechanisms, including oncogene-induced replication stress and deregulation of DNA replication, have been reported to introduce DNA alterations during tumorigenesis, resulting in variable frequencies of somatic mutations in different cancer subtypes [106, 6]. Genetic mutations in tumor cells can result in the production of neoantigens, which are presented in context of MHC molecules on cancer cells to tumor-infiltrating lymphocytes (TILs) [58, 153, 143]. High mutational burden has been associated with microsatellite instability (MSI) and mismatch-repair deficiency (MMR-D) in studies of colorectal carcinoma [165, 105, 97, 21, 49, 98].

Diagnostic significance: Unknown.

Prognostic significance: Unknown.

Drug sensitivity: Studies have shown high tumor mutational burden or high levels of neoantigens to be associated with high expression of cytotoxic T-cell markers; thus, immunotherapies may be relevant in tumors with high tumor mutational burden [72, 58, 17, 143]. A large study of advanced cancer patients reported that higher tumor mutation burden (TMB, defined as the highest 20% in each histology) was associated with significantly increased survival in patients treated with a variety of immune checkpoint inhibitors, although the numeric cutoff for TMB in each histology was highly variable. Tumor types with the most significant improvement were bladder, colorectal, head and neck, melanoma, and NSCLC. Breast cancer and glioma with high TMB were not associated with increased survival [148].

Drug resistance: None.

Approved Drugs: None.

Phase 3: The second interim analysis of the Phase 3 KEYNOTE-177 study comparing 1:1 pembrolizumab with investigator choice of chemotherapy in 307 patients with MSI-high or deficient MMR metastatic colorectal carcinoma has reported a progression-free survival of 16.5 and 8.2 months, a confirmed overall response rate of 43.8% and 33.1%, an estimated restricted mean survival of 13.7 and 10.8 months, and grade 3-5 serious adverse events in 22% and 66% of patients treated with pembrolizumab or chemotherapy, respectively. Ongoing responses at 24 months were observed in 83% and 35% of patients treated with pembrolizumab or chemotherapy, respectively [8]. A Phase 3 trial (IMblaze370) of atezolizumab with cobimetinib (AC), atezolizumab monotherapy (AM), or regorafenib in patients with metastatic colorectal carcinoma reported median overall survival of 8.87, 7.1, and 8.51 months in the AC, AM, and regorafenib arms, respectively. Treatment-related grade 3-4 adverse events were reported in 61% (109/179), 31% (28/90), and 58% (46/80), respectively [45].

Phase 2: The Phase 2 KEYNOTE-158 study of pembrolizumab in advanced solid tumor patients has reported an overall objective response in 29% (30/102) of patients with high tumor mutational burden (TMB-H), defined as ten or more mutations per megabase [115]. A Phase 2 trial of pembrolizumab with cyclophosphamide and the colon vaccine GVAX in 17 patients with advanced MMR-proficient colorectal carcinoma reported disease control rates of 18% and 29% by RECIST and irRC, respectively. No objective responses were reported; median progression-free and overall survival were 2.7 and 7.1 months. Grade 3-4 adverse events attributed to study therapy were reported in 11.8% (2/17) of patients [198]. A Phase 2 study of nivolumab in 74 dMMR or MSI-H CRC patients has reported investigator and independent radiology review committee objective response rates of 27% and 31% and disease control rates of 62% and 69% with a time to response of approximately 2.7

Biomarkers (2)

months; progression-free survival rates at 12 months were 46% and 48% with 83.4% overall survival at six months and 73.8% at 12 months [129]. A Phase 2 study of nivolumab in combination with trifluridine/tipiracil in 18 patients with MSS colorectal carcinoma has reported 13 grade 3 or higher adverse events, no tumor responses, stable disease in eight and ten patients, and a median progression-free survival of 2.2 and 2.8 months per immune-related response criteria (irRC) and RECIST, respectively [131]. A Phase 2 study of nivolumab with ipilimumab in 40 patients with MSS colorectal adenocarcinoma reported a disease control rate of 25%, and an objective response rate of 10% by intent to treat analysis. A Phase 2 trial of durvalumab with tremelimumab (DT) plus best supportive care (BSC) or BSC alone in 179 patients with advanced/refractory colorectal carcinoma reported median progression-free survival of 1.8 and 1.9 months, and median overall survival of 6.6 and 4.1 months in the DT+BSC and BSC alone arms, respectively. Disease control rates were 22.7% and 6.6%, respectively [28]. A Phase 2 study of durvalumab and trametinib in 29 microsatellite stable metastatic colorectal cancer patients has reported objective response in 3.4% and stable disease in 24% of patients. No grade 4 treatment-related adverse events were observed. The Phase 2 AVETUX study of avelumab plus cetuximab in combination with FOLFOX in 39 previously untreated metastatic colorectal cancer patients has reported overall response rate of 79.5%, with six complete and 25 partial responses, and disease control rate of 92.3%. The Phase 2 CAVE study of avelumab and cetuximab in RAS wild-type metastatic colorectal carcinoma patients with response to first-line chemotherapy in combination with anti-Egfr therapy has reported complete response in 1.5%, partial response in 4.6%, and stable disease in 49.2% of 65 evaluable patients. Median progression-free and overall survival were 3.6 months and 13.1 months, respectively, and grade 3 adverse events were observed in 22% (16/77) of patients. A Phase 2 study of avelumab monotherapy in 33 metastatic or unresectable colorectal carcinoma patients with dMMR/MSI-H or POLE mutations has reported objective response rate of 24.2% and median progression-free survival and overall survival of 3.9 and 13.2 months, respectively [88]. A Phase 2 study of short-course radiation followed by mFOLFOX6 with avelumab in 13 patients with locally advanced rectal cancer has reported pathologic complete response (pCR) in 3/12 and near pCR in 3/12 evaluable patients. No grade 4 serious adverse events were observed [156].

Phase 1: A Phase 1b trial of atezolizumab in combination with bevacizumab in refractory metastatic colorectal cancer patients reported an unconfirmed overall response rate (ORR) of 8% (1/13) and grades 3/4 adverse events in 64% of cases. In oxaliplatin-naïve patients treated with atezolizumab in combination with bevacizumab and FOLFOX, the unconfirmed ORR in evaluable patients was 36% (9/25), with 73% of cases reporting grades 3/4 adverse events. A Phase 1b trial of atezolizumab in combination with bevacizumab in microsatellite instability (MSI)-high metastatic colorectal cancer patients has reported partial response in 30% (3/10) and stable disease in 60% (6/10) of patients. Grade 3/4 adverse events were observed in 40% of patients. An ongoing Phase 1b trial of atezolizumab in combination with cobimetinib in 84 chemotherapy-refractory or locally advanced metastatic colorectal cancer patients has reported an overall response rate of 8%, including patients with MSS and MSI-low status, and disease control rate of 31%, with median progression-free and overall survival of 1.9 and 10.0 months, respectively. A Phase 1 study of durvalumab plus monalizumab in 55 solid tumor patients has reported confirmed partial response in three and stable disease in 11 patients in the expansion cohort of 40 metastatic microsatellite-stable colorectal cancer patients, with disease control rate at 16 weeks of 24%. A Phase 1 study of oleclumab and durvalumab in 66 solid tumor patients, with an expansion cohort in 41 advanced microsatellite-stable colorectal cancer (MSS-CRC) and pancreatic cancer patients has reported partial response in 5% (1/21) and 10% (2/20), and stable disease in 10% (2/21) and 15% (3/20) of MSS-CRC and pancreatic cancer patients, respectively.

Preclinical: N/A: Preclinical data are not presented when higher level data are available.

Microsatellite Status: MSI-high

Biomarker: MSI-high
Classification: Tier 1A
Assessment: Pathogenic

Treatment options
 4 Sensitive
 5 Trials

Biomarker summary: MSI-high instability exhibits altered function compared to wild-type.

Clinical relevance: Tumors exhibiting microsatellite instability (MSI) have a higher mutational burden than microsatellite stable (MSS) tumors and express higher levels of immune checkpoint receptors [86, 5, 72, 174, 109]. Thus, checkpoint inhibitors, several of which have received agency approval for certain indications, may be clinically relevant for tumors exhibiting MSI [122, 176, 130, 172, 97, 187]. In fact, pembrolizumab has been FDA-approved as a second or later line of therapy for the treatment of pediatric and adult solid tumors with high microsatellite instability (MSI-H) or that are deficient in mismatch repair (dMMR) and as a front-line therapy by the FDA and EMA for colorectal carcinoma patients with MSI-H or dMMR [102, 97].

Disease summary: MSI has been associated with proximal colon location, mucinous histology, lower tumor grade, age at diagnosis of less 50 years old, and the presence of BRAF mutation in colorectal carcinoma studies [164, 173, 27, 53, 9]. Adjuvant 5-FU may not benefit colorectal cancer patients with stage II/III MSI-H tumors when given as monotherapy [139, 150, 171].

Molecular function: The revised Bethesda Guidelines recommended criteria for defining a tumor with high microsatellite instability (MSI-H) include detecting alterations in two or more of the five microsatellite markers included in the National Cancer Institute (NCI) microsatellite panel [12, 179].

Incidence: MSI has been reported in 3-24% of colorectal carcinoma samples and has been observed in both inherited and sporadic forms of the disease [59, 169, 110, 30, 164, 126, 178, 137, 70, 170, 53].

Biomarkers (2)

Role in disease: MSI is associated with the loss or dysfunction of DNA mismatch repair (MMR) proteins that are required for correcting errors that occur during DNA replication or recombination; germline mutations in genes encoding MMR proteins are associated with Lynch syndrome, a hereditary cancer-predisposition syndrome also known as hereditary nonpolyposis colorectal cancer (HNPCC) [111, 34, 70]. Tumors exhibiting MSI have been reported to have increased numbers of tumor-infiltrating lymphocytes (TILs) and a significantly higher mutational burden than microsatellite stable (MSS) tumors [86, 5, 72, 174]. MSI has been associated with proximal colon location, mucinous histology, lower tumor grade, age at diagnosis of less 50 years old, and the presence of BRAF mutation in colorectal carcinoma studies [164, 9, 173, 27, 53]. In addition, increased frequency of PD-L1 expression has been reported in colorectal carcinoma cases with MSI-H as compared with MSS/MSI-L cases [92, 181, 147, 99, 123].

Diagnostic significance: Unknown.

Prognostic significance: Several studies have reported MSI-high status to be associated with better prognosis in colorectal carcinoma patients compared with patients lacking MSI-high [183, 66, 188, 136].

Drug sensitivity: MSI has been reported to correlate with high levels of immune checkpoint gene expression in some types of cancer, including colorectal and endometrial carcinoma, and with clinical response to checkpoint inhibition in colorectal carcinoma; thus, immunotherapies may be relevant in tumors exhibiting MSI [109, 97, 72]. Checkpoint inhibitors are currently in clinical development, several of which have received agency approval for certain indications [122, 187]. A Phase 1 follow-up study of nivolumab in 39 treatment-refractory solid tumor patients reported that one patient with MSI-H colorectal cancer showed an ongoing complete response of at least three years [107].

Drug resistance: Adjuvant 5-FU may not benefit colorectal cancer patients with stage II/III MSI-H tumors when given as monotherapy [139, 150, 171]. The therapeutic implications of MLH1 mutation or hypermethylation have been best studied in colon cancer. High MSI has been associated with lack of benefit from 5-FU based regimens [139, 150, 171, 68].

Approved Drugs: Pembrolizumab.

Phase 3: The second interim analysis of the Phase 3 KEYNOTE-177 study comparing 1:1 pembrolizumab with investigator choice of chemotherapy in 307 patients with MSI-high or deficient MMR metastatic colorectal carcinoma has reported a progression-free survival of 16.5 and 8.2 months, a confirmed overall response rate of 43.8% and 33.1%, an estimated restricted mean survival of 13.7 and 10.8 months, and grade 3-5 serious adverse events in 22% and 66% of patients treated with pembrolizumab or chemotherapy, respectively. Ongoing responses at 24 months were observed in 83% and 35% of patients treated with pembrolizumab or chemotherapy, respectively [8]. A Phase 3 trial (IMblaze370) of atezolizumab with cobimetinib (AC), atezolizumab monotherapy (AM), or regorafenib in patients with metastatic colorectal carcinoma reported median overall survival of 8.87, 7.1, and 8.51 months in the AC, AM, and regorafenib arms, respectively. Treatment-related grade 3-4 adverse events were reported in 61% (109/179), 31% (28/90), and 58% (46/80), respectively [45].

Phase 2: A safety and efficacy study of pembrolizumab in 149 patients with tumors with high microsatellite instability (MSI-H) or deficient in mismatch repair (dMMR) across five Phase 1 and 2 uncontrolled trials and including 15 different cancer types has reported a complete or partial response in 39.6% of patients; 78% of patients experienced response for six months or more [116, 102, 97]. A Phase 1/2 study of durvalumab monotherapy in 62 patients with MSI-high solid tumors, including 36 patients with colorectal carcinoma (CRC), 17 patients with endometrial carcinoma, and 9 patients with other tumor types, has reported an objective response rate of 23% and 22% for all patients and CRC patients, respectively; treatment-related adverse events (TRAEs) were reported in 60% (37/62) of patients, including grade 3/4 TRAEs in 3% (2/62) of cases. A Phase 2 trial of pembrolizumab with cyclophosphamide and the colon vaccine GVAX in 17 patients with advanced MMR-proficient colorectal carcinoma reported disease control rates of 18% and 29% by RECIST and irRC, respectively. No objective responses were reported; median progression-free and overall survival were 2.7 and 7.1 months. Grade 3-4 adverse events attributed to study therapy were reported in 11.8% (2/17) of patients [198]. Long-term followup of the Phase 2 CheckMate 142 study of nivolumab plus ipilimumab in dMMR or MSI-H CRC patients who had received at least one prior therapy reported an investigator-assessed overall response rate of 58% and a disease control rate of 81% at a median followup of 25.4 months. Complete and partial responses were reported in 6% (7/119) and 52% (62/119) of patients; median progression-free and overall survival rates at 24 months were 60% and 74%. Grade 3-4 adverse events were reported in 31% of patients. A Phase 2 study of nivolumab plus ipilimumab with radiotherapy in 40 metastatic microsatellite stable colorectal carcinoma (CRC) patients and 25 metastatic pancreatic ductal adenocarcinoma (PDAC) patients with progression on previous lines of therapy has reported disease control in 25% (10/40) and 20% (5/25), and overall response in 10% (4/40) and 13% (3/25) of patients in the CRC and PDAC cohorts, respectively. Grade 3 or higher treatment-related adverse events were observed in 40% (26/65) of patients, with grade 5 events in 3.1% (2/65) of patients. The Phase 2 GERCOR NIPICOL study of nivolumab plus ipilimumab in 57 pretreated patients with MSI-H/dMMR metastatic colorectal cancer has reported 12-week disease control rate of 86.0% and 87.7%, and 12-month progression-free survival rate of 72.9% and 76.5%, according to RECIST 1.1 and irRECIST criteria, respectively. Overall response rate was 59.7% [33]. The Phase 2 NICHE study of neoadjuvant ipilimumab and nivolumab in early-stage colon cancer patients has reported pathological response in 100% (20/20) of patients with MMR-deficient tumors and in 27% (4/15) of patients with MMR-proficient tumors [25]. A Phase 2 study of nivolumab in 74 dMMR or MSI-H CRC patients has reported

Biomarkers (2)

investigator and independent radiology review committee objective response rates of 27% and 31% and disease control rates of 62% and 69% with a time to response of approximately 2.7 months; progression-free survival rates at 12 months were 46% and 48% with 83.4% overall survival at six months and 73.8% at 12 months [129]. A Phase 2 study of nivolumab in combination with trifluridine/tipiracil in 18 patients with MSS colorectal carcinoma has reported 13 grade 3 or higher adverse events, no tumor responses, stable disease in eight and ten patients, and a median progression-free survival of 2.2 and 2.8 months per immune-related response criteria (irRC) and RECIST, respectively [131]. A Phase 2 study of nivolumab with ipilimumab in 40 patients with MSS colorectal adenocarcinoma reported a disease control rate of 25%, and an objective response rate of 10% by intent to treat analysis. A Phase 2 trial of durvalumab with tremelimumab (DT) plus best supportive care (BSC) or BSC alone in 179 patients with advanced/refractory colorectal carcinoma reported median progression-free survival of 1.8 and 1.9 months, and median overall survival of 6.6 and 4.1 months in the DT+BSC and BSC alone arms, respectively. Disease control rates were 22.7% and 6.6%, respectively [28]. A Phase 2 study of durvalumab and trametinib in 29 microsatellite stable metastatic colorectal cancer patients has reported objective response in 3.4% and stable disease in 24% of patients. No grade 4 treatment-related adverse events were observed. The Phase 2 AVETUX study of avelumab plus cetuximab in combination with FOLFOX in 39 previously untreated metastatic colorectal cancer patients has reported overall response rate of 79.5%, with six complete and 25 partial responses, and disease control rate of 92.3%. The Phase 2 CAVE study of avelumab and cetuximab in RAS wild-type metastatic colorectal carcinoma patients with response to first-line chemotherapy in combination with anti-Egfr therapy has reported complete response in 1.5%, partial response in 4.6%, and stable disease in 49.2% of 65 evaluable patients. Median progression-free and overall survival were 3.6 months and 13.1 months, respectively, and grade 3 adverse events were observed in 22% (16/77) of patients. A Phase 2 study of avelumab monotherapy in 33 metastatic or unresectable colorectal carcinoma patients with dMMR/MSI-H or POLE mutations has reported objective response rate of 24.2% and median progression-free survival and overall survival of 3.9 and 13.2 months, respectively [88]. A Phase 2 study of short-course radiation followed by mFOLFOX6 with avelumab in 13 patients with locally advanced rectal cancer has reported pathologic complete response (pCR) in 3/12 and near pCR in 3/12 evaluable patients. No grade 4 serious adverse events were observed [156].

Phase 1: A Phase 1b trial of atezolizumab in combination with bevacizumab in refractory metastatic colorectal cancer patients reported an unconfirmed overall response rate (ORR) of 8% (1/13) and grades 3/4 adverse events in 64% of cases. In oxaliplatin-naïve patients treated with atezolizumab in combination with bevacizumab and FOLFOX, the unconfirmed ORR in evaluable patients was 36% (9/25), with 73% of cases reporting grades 3/4 adverse events. A Phase 1b trial of atezolizumab in combination with bevacizumab in microsatellite instability (MSI)-high metastatic colorectal cancer patients has reported partial response in 30% (3/10) and stable disease in 60% (6/10) of patients. Grade 3/4 adverse events were observed in 40% of patients. An ongoing Phase 1b trial of atezolizumab in combination with cobimetinib in 84 chemotherapy-refractory or locally advanced metastatic colorectal cancer patients has reported an overall response rate of 8%, including patients with MSS and MSI-low status, and disease control rate of 31%, with median progression-free and overall survival of 1.9 and 10.0 months, respectively. A Phase 1 study of durvalumab plus monalizumab in 55 solid tumor patients has reported confirmed partial response in three and stable disease in 11 patients in the expansion cohort of 40 metastatic microsatellite-stable colorectal cancer patients, with disease control rate at 16 weeks of 24%. A Phase 1 study of oleclumab and durvalumab in 66 solid tumor patients, with an expansion cohort in 41 advanced microsatellite-stable colorectal cancer (MSS-CRC) and pancreatic cancer patients has reported partial response in 5% (1/21) and 10% (2/20), and stable disease in 10% (2/21) and 15% (3/20) of MSS-CRC and pancreatic cancer patients, respectively.

Preclinical: N/A: Preclinical data are not presented when higher level data are available.

Variants of potential clinical significance (3)

ARID1A H688fs*129

Gene: ARID1A

Exon: 5

Nucleotide:

NM_006015.6:

g.27087482_2708748

3insC

c.2060dupC

Amino Acid: p.H688fs*129

Allelic Fraction: 16.0% (of 1147 reads)

Classification: Tier 2C

Assessment: Likely Pathogenic

Biomarker summary: ARID1A-H688fs*129 is an inactivating mutation.

Clinical relevance: ARID1A encodes Arid1a, also known as Baf250a, a member of the SWI/SNF chromatin remodeling complex. Mutation, loss, or inactivation of ARID1A has been reported in many cancers, and functional studies have implicated it as a tumor suppressor [190, 82, 67, 81]. There are no approved targeted therapies that directly target ARID1A alterations at this time. However, inactivating ARID1A mutations and loss of Arid1a expression may predict sensitivity to Ezh2 inhibitors [11]. Small molecule inhibitors of Ezh2, such as tazemetostat, are currently under investigation in clinical studies [91, 36]. In addition, ARID1A-deficient preclinical cancer models exhibit sensitivity to Atr, PARP, and BET domain inhibitors and clinical trials are evaluating these agents in patients with loss of Arid1a expression or ARID1A mutations [192, 159, 10, 24].

Disease summary: Loss of Arid1a expression in colorectal carcinoma has been correlated with mismatch repair (MMR) deficiency and poor tumor differentiation [200, 31, 189, 100, 195]. ARID1A bi-allelic deletion has been reported to result in the development of colon adenocarcinoma in mice, however, in a separate study, ARID1A depletion has been observed to reduce cell viability in KRAS-mutant colorectal carcinoma cell lines [119, 154].

Molecular function: This frameshift alteration is expected to effectively truncate the Arid1a protein prior to the ARID domain, resulting in the loss of the entire ARID domain and three of four LXXLL motifs (UniProt), which

Variants of potential clinical significance (3)

are protein-protein interaction motifs and may mediate the binding of Arid1a with nuclear receptors [35]. The ARID domain is required for Arid1a-DNA interactions and promoter occupancy by SWI/SNF chromatin-remodeling complex [26]. ARID1A mutations, which are mostly truncating mutations, have been shown to be correlated with loss of Arid1a protein and predicted to be inactivating [81, 190, 82]. Therefore, this alteration is predicted to lead to a loss of Arid1a function.

Incidence: ARID1A mutations have been reported in 10% (371/3643) of Colorectal carcinoma (CRC) samples analyzed in COSMIC (May 2020). ARID1A mutations have been reported in 9.4-11% of Colorectal carcinoma (CRC) samples (cBioPortal for Cancer Genomics, May 2020). Other studies have reported ARID1A mutations in 6-10% of CRC cases; however, one study of microsatellite-unstable CRC reports ARID1A mutation in 39% of cases [21, 81, 19, 101].

Role in disease: Loss of Arid1a has been associated with mismatch repair deficiency in cancer, including endometrial and colorectal carcinoma [158, 31, 7]. Loss of Arid1a expression in colorectal carcinoma has been correlated with mismatch repair (MMR) deficiency and poor tumor differentiation [200, 31, 189, 100, 195]. ARID1A bi-allelic deletion has been reported to result in the development of colon adenocarcinoma in mice, however, in a separate study, ARID1A depletion has been observed to reduce cell viability in KRAS-mutant colorectal carcinoma cell lines [119, 154].

Diagnostic significance: Unknown.

Prognostic significance: Unknown.

Drug sensitivity: There are no approved targeted therapies to address ARID1A alterations at this time; however, ARID1A alterations may predict sensitivity to Ezh2 inhibitors [11]. Ezh2 inhibitors, such as tazemetostat, are currently being evaluated in clinical trials in patients with solid tumors or B-cell lymphomas [94, 91, 36]. In addition, ARID1A-deficient preclinical cancer models exhibit sensitivity to Atr, PARP, and BET domain inhibitors and clinical trials are evaluating these agents in patients with loss of Arid1a expression or ARID1A mutations [192, 159, 10, 24].

Drug resistance: None.

Approved Drugs: None.

Phase 3: None.

Phase 2: A Phase 2 trial of olaparib in 20 patients with microsatellite stable (MSS) colorectal cancer (CRC) and 13 with CRC with high-level microsatellite instability (MSI) has reported no complete or partial responses, with a median progression-free survival for all patients of 1.84 months; similar median progression-free and overall survival times were noted regardless of MSS or MSI status [103]. A Phase 2 study of veliparib with temozolomide in 75 metastatic colorectal carcinoma patients has reported a disease control rate of 24% with two confirmed partial responses; median progression-free survival was 1.8 months, and median overall survival was 6.6 months, with dose reductions required in some patients due to myelosuppression [133]. A Phase 2 study of FOLFIRI with veliparib or placebo in 130 patients with metastatic colorectal carcinoma reported objective response rates of 56.9% and 61.5%, median progression-free survival of 12 and 11 months, and median overall survival of 25 and 27 months, in the veliparib and placebo arms, respectively [61, 62].

Phase 1: A Phase 1 trial of niraparib in patients with advanced solid tumors has reported partial responses in 2/4 breast cancer patients with germline BRCA1/2 mutations and in 40% (8/20) of ovarian cancer patients with BRCA1/2 mutations, as well as anti-tumor activity in sporadic high-grade serous ovarian cancer, non-small cell lung cancer, and prostate cancer [149]. A Phase 1 trial of talazoparib in 110 patients with advanced solid tumors, including 71 patients in the expansion cohort, has reported an overall response rate of 22% (16/72), including two and 14 patients with complete and partial responses, respectively, and 16 stable diseases per RECIST in the expansion cohort [204]. A Phase 1 trial of talazoparib combined with irinotecan with or without temozolomide in 41 pediatric patients with recurrent or refractory solid malignancies has reported an overall response rate of 10.3% in the talazoparib combined with irinotecan arm compared with 25% in the talazoparib combined with irinotecan and temozolomide arm. In addition, a complete response was reported in an Ewing sarcoma patient and partial responses were reported in one synovial sarcoma and four Ewing sarcoma patients [50]. A Phase 1/2 study of rucaparib in 56 patients with advanced solid tumors and 42 patients with germline BRCA1/2-mutant high-grade ovarian carcinoma has reported two complete responses, six partial responses, and 22 stable diseases in the solid tumor cohort and four complete responses, 21 partial responses, and 12 stable diseases in the BRCA1/2 cohort per RECIST [93]. A Phase 1b study of veliparib in combination with capecitabine plus radiotherapy in patients with locally advanced rectal cancer reported tumor downstaging after surgery in 71% of 31 evaluable patients, with a pathologic complete response in 29% of patients. An acceptable safety profile was reported, with a recommended phase 2 dose of 400 mg BID; the maximum tolerated dose was not reached [38]. A Phase 1 study of tazemetostat in 43 patients with solid tumors and 21 patients with B-cell non-Hodgkin lymphoma (NHL) has reported response rates of 5% (2/43) and 38% (8/21) in solid tumor and NHL patients, respectively. Grade 4 thrombocytopenia was the only dose-limiting toxicity observed [76].

Preclinical: A preclinical study has reported that niraparib treatment of microsatellite stable and unstable colorectal cancer cell lines inhibits proliferation and enhances the anti-proliferative effects of SN-38 in vitro, as

Variants of potential clinical significance (3)

well as further delays tumor regrowth when combined with irinotecan in vivo, compared with irinotecan treatment alone [56]. A preclinical study analyzing 93 colorectal carcinoma cell lines reported that eight exhibited sensitivity to PARPi treatment, and that among these eight cell lines, talazoparib demonstrated greater efficacy than olaparib, niraparib, veliparib, or rucaparib [162]. A preclinical study has reported that tazemetostat treatment decreased tumor growth in a colorectal carcinoma xenograft mouse model [29].

MSH2 N538fs*5

Gene: MSH2

Exon: 10

Nucleotide:

NM_000251.3:

g.47693895delA

c.1613delA

Amino Acid: p.N538fs*5

Allelic Fraction: 18.0% (of 335 reads)

Classification: Tier 2C

Assessment: Likely Pathogenic

Biomarker summary: MSH2-N538fs*5 is an inactivating mutation.

Clinical relevance: MSH2 encodes MutS protein homolog 2 (Msh2), a member of the mismatch repair (MMR) gene family; defective MMR as a result of inactivating MSH2 mutation can result in microsatellite instability (MSI) [125, 51]. Germline mutations in MSH2 or genes encoding other mismatch repair proteins such as MLH1, MSH6, and PMS2 are associated with Lynch syndrome, which is a hereditary cancer-predisposition syndrome also known as hereditary nonpolyposis colorectal cancer (HNPCC) [111]. While there are currently no approved therapies directly addressing loss or mutation in MSH2, PD-1/PD-L1 inhibitors have been reported to be effective in tumors harboring mismatch repair defects; thus, a tumor with Msh2 inactivation may be sensitive to these therapies [97, 141, 23, 57].

Disease summary: Alterations in MSH2 or other mismatch repair genes (such as MLH1, MSH6, and PMS2) have been reported to underlie hereditary nonpolyposis colorectal cancer (HNPCC). Additionally, research suggests that carriers of MSH2 mutations have an increased risk of developing colorectal cancer [79, 44, 1].

Molecular function: The MSH2 frameshift alteration reported here is expected to effectively truncate the Msh2 protein, resulting in the loss of a portion of the C-terminal domain, which is involved in MutS dimer formation (InterPro). Truncating mutations in the C-terminal domain have been reported as germline alterations in Lynch syndrome families, and truncation of the C-terminal 60 amino acids of Msh2 has been reported to result in reduced mismatch repair capacity [191]. Therefore, this mutation is predicted to lead to a loss of Msh2 function.

Incidence: MSH2 mutations have been reported in 5.0% (208/4169) of Colorectal carcinoma (CRC) samples analyzed in COSMIC (May 2020). MSH2 mutations have been reported in 1.9-2.2% of Colorectal carcinoma (CRC) samples (cBioPortal for Cancer Genomics, May 2020). Scientific studies have reported MSH2 mutation in approximately 2-12% of colorectal carcinoma specimens analyzed [202, 2, 84, 132].

Role in disease: Defective mismatch repair (MMR), occurring as a result of mutation(s) in the MMR family (MLH1, MSH2, MSH6, or PMS2) can result in microsatellite instability (MSI), common in colon, endometrium and stomach cancers [117]. MSH2 alterations have been reported to underlie hereditary nonpolyposis colorectal cancer (HNPCC) and research suggests that carriers of MSH2 mutations have an increased risk of developing colorectal cancer [79, 44, 1].

Diagnostic significance: Unknown.

Prognostic significance: Unknown.

Drug sensitivity: While there are currently no approved therapies directly addressing loss or mutation in MSH2, PD-1/PD-L1 inhibitors have been reported to be effective in tumors harboring mismatch repair defects; thus, a tumor with Msh2 inactivation may be sensitive to PD-1/PD-L1 inhibitors [97, 141, 23, 57].

Drug resistance: None.

Approved Drugs: None.

Phase 3: The second interim analysis of the Phase 3 KEYNOTE-177 study comparing 1:1 pembrolizumab with investigator choice of chemotherapy in 307 patients with MSI-high or deficient MMR metastatic colorectal carcinoma has reported a progression-free survival of 16.5 and 8.2 months, a confirmed overall response rate of 43.8% and 33.1%, an estimated restricted mean survival of 13.7 and 10.8 months, and grade 3-5 serious adverse events in 22% and 66% of patients treated with pembrolizumab or chemotherapy, respectively. Ongoing responses at 24 months were observed in 83% and 35% of patients treated with pembrolizumab or chemotherapy, respectively [8]. A Phase 3 trial (IMblaze370) of atezolizumab with cobimetinib (AC), atezolizumab monotherapy (AM), or regorafenib in patients with metastatic colorectal carcinoma reported median overall survival of 8.87, 7.1, and 8.51 months in the AC, AM, and regorafenib arms, respectively. Treatment-related grade 3-4 adverse events were reported in 61% (109/179), 31% (28/90), and 58% (46/80), respectively [45].

Phase 2: A safety and efficacy study of pembrolizumab in 149 patients with tumors with high microsatellite instability (MSI-H) or deficient in mismatch repair (dMMR) across five Phase 1 and 2 uncontrolled trials and including 15 different cancer types has reported a complete or partial response in 39.6% of patients; 78% of patients experienced response for six months or more [116, 102, 97]. A Phase 2 study of pembrolizumab in patients with advanced dMMR cancers, including 12 different tumor types, has reported complete and partial responses in 23.1% (18/78) and 35.9% (28/78) of evaluable patients, respectively [96]. Long-term followup of the Phase 2 CheckMate 142 study of nivolumab plus ipilimumab in dMMR or MSI-H CRC patients who had received at least one prior therapy reported an investigator-assessed overall response rate of 58% and a

Variants of potential clinical significance (3)

disease control rate of 81% at a median followup of 25.4 months. Complete and partial responses were reported in 6% (7/119) and 52% (62/119) of patients; median progression-free and overall survival rates at 24 months were 60% and 74%. Grade 3-4 adverse events were reported in 31% of patients. A Phase 2 study of nivolumab plus ipilimumab with radiotherapy in 40 metastatic microsatellite stable colorectal carcinoma (CRC) patients and 25 metastatic pancreatic ductal adenocarcinoma (PDAC) patients with progression on previous lines of therapy has reported disease control in 25% (10/40) and 20% (5/25), and overall response in 10% (4/40) and 13% (3/25) of patients in the CRC and PDAC cohorts, respectively. Grade 3 or higher treatment-related adverse events were observed in 40% (26/65) of patients, with grade 5 events in 3.1% (2/65) of patients. The Phase 2 GERCOR NIPICOL study of nivolumab plus ipilimumab in 57 pretreated patients with MSI-H/dMMR metastatic colorectal cancer has reported 12-week disease control rate of 86.0% and 87.7%, and 12-month progression-free survival rate of 72.9% and 76.5%, according to RECIST 1.1 and iRECIST criteria, respectively. Overall response rate was 59.7% [33]. The Phase 2 NICHE study of neoadjuvant ipilimumab and nivolumab in early-stage colon cancer patients has reported pathological response in 100% (20/20) of patients with MMR-deficient tumors and in 27% (4/15) of patients with MMR-proficient tumors [25]. A Phase 2 study of nivolumab in 74 dMMR or MSI-H CRC patients has reported investigator and independent radiology review committee objective response rates of 27% and 31% and disease control rates of 62% and 69% with a time to response of approximately 2.7 months; progression-free survival rates at 12 months were 46% and 48% with 83.4% overall survival at six months and 73.8% at 12 months [129]. A Phase 2 study of nivolumab in combination with trifluridine/tipiracil in 18 patients with MSS colorectal carcinoma has reported 13 grade 3 or higher adverse events, no tumor responses, stable disease in eight and ten patients, and a median progression-free survival of 2.2 and 2.8 months per immune-related response criteria (irRC) and RECIST, respectively [131]. A Phase 2 study of nivolumab with ipilimumab in 40 patients with MSS colorectal adenocarcinoma reported a disease control rate of 25%, and an objective response rate of 10% by intent to treat analysis. A Phase 2 trial of pembrolizumab with cyclophosphamide and the colon vaccine GVAX in 17 patients with advanced MMR-proficient colorectal carcinoma reported disease control rates of 18% and 29% by RECIST and irRC, respectively. No objective responses were reported; median progression-free and overall survival were 2.7 and 7.1 months. Grade 3-4 adverse events attributed to study therapy were reported in 11.8% (2/17) of patients [198]. A Phase 2 trial of durvalumab with tremelimumab (DT) plus best supportive care (BSC) or BSC alone in 179 patients with advanced/refractory colorectal carcinoma reported median progression-free survival of 1.8 and 1.9 months, and median overall survival of 6.6 and 4.1 months in the DT+BSC and BSC alone arms, respectively. Disease control rates were 22.7% and 6.6%, respectively [28]. A Phase 2 study of durvalumab and trametinib in 29 microsatellite stable metastatic colorectal cancer patients has reported objective response in 3.4% and stable disease in 24% of patients. No grade 4 treatment-related adverse events were observed. The Phase 2 AVETUX study of avelumab plus cetuximab in combination with FOLFOX in 39 previously untreated metastatic colorectal cancer patients has reported overall response rate of 79.5%, with six complete and 25 partial responses, and disease control rate of 92.3%. The Phase 2 CAVE study of avelumab and cetuximab in RAS wild-type metastatic colorectal carcinoma patients with response to first-line chemotherapy in combination with anti-Egfr therapy has reported complete response in 1.5%, partial response in 4.6%, and stable disease in 49.2% of 65 evaluable patients. Median progression-free and overall survival were 3.6 months and 13.1 months, respectively, and grade 3 adverse events were observed in 22% (16/77) of patients. A Phase 2 study of avelumab monotherapy in 33 metastatic or unresectable colorectal carcinoma patients with dMMR/MSI-H or POLE mutations has reported objective response rate of 24.2% and median progression-free survival and overall survival of 3.9 and 13.2 months, respectively [88]. A Phase 2 study of short-course radiation followed by mFOLFOX6 with avelumab in 13 patients with locally advanced rectal cancer has reported pathologic complete response (pCR) in 3/12 and near pCR in 3/12 evaluable patients. No grade 4 serious adverse events were observed [156].

Phase 1: A Phase 1 study of durvalumab plus monalizumab in 55 solid tumor patients has reported confirmed partial response in three and stable disease in 11 patients in the expansion cohort of 40 metastatic microsatellite-stable colorectal cancer patients, with disease control rate at 16 weeks of 24%. A Phase 1 study of oleclumab and durvalumab in 66 solid tumor patients, with an expansion cohort in 41 advanced microsatellite-stable colorectal cancer (MSS-CRC) and pancreatic cancer patients has reported partial response in 5% (1/21) and 10% (2/20), and stable disease in 10% (2/21) and 15% (3/20) of MSS-CRC and pancreatic cancer patients, respectively. A Phase 1b trial of atezolizumab in combination with bevacizumab in refractory metastatic colorectal cancer patients reported an unconfirmed overall response rate (ORR) of 8% (1/13) and grades 3/4 adverse events in 64% of cases. In oxaliplatin-naïve patients treated with atezolizumab in combination with bevacizumab and FOLFOX, the unconfirmed ORR in evaluable patients was 36% (9/25), with 73% of cases reporting grades 3/4 adverse events. A Phase 1b trial of atezolizumab in combination with bevacizumab in microsatellite instability (MSI)-high metastatic colorectal cancer patients has reported partial response in 30% (3/10) and stable disease in 60% (6/10) of patients. Grade 3/4 adverse events were observed in 40% of patients. An ongoing Phase 1b trial of atezolizumab in combination with cobimetinib in 84 chemotherapy-refractory or locally advanced metastatic colorectal cancer patients has reported an overall response rate of 8%, including patients with MSS and MSI-low status, and disease control rate of 31%, with median progression-free and overall survival of 1.9 and 10.0 months, respectively.

Preclinical: N/A: Preclinical data are not presented when higher level data are available.

MSH2 N835fs*4

Gene: MSH2
Exon: 15
Nucleotide:

Biomarker summary: MSH2-N835fs*4 is an inactivating mutation.

Variants of potential clinical significance (3)

NM_000251.3:
 g.47707877_4770788
 3delCTAATTT
 c.2502_2508delTAAT
 TTC
Amino Acid: p.N835fs*4
Allelic Fraction: 35.0% (of 421
 reads)
Classification: Tier 2C
Assessment: Pathogenic

Clinical relevance: MSH2 encodes MutS protein homolog 2 (Msh2), a member of the mismatch repair (MMR) gene family; defective MMR as a result of inactivating MSH2 mutation can result in microsatellite instability (MSI) [125, 51]. Germline mutations in MSH2 or genes encoding other mismatch repair proteins such as MLH1, MSH6, and PMS2 are associated with Lynch syndrome, which is a hereditary cancer-predisposition syndrome also known as hereditary nonpolyposis colorectal cancer (HNPCC) [111]. While there are currently no approved therapies directly addressing loss or mutation in MSH2, PD-1/PD-L1 inhibitors have been reported to be effective in tumors harboring mismatch repair defects; thus, a tumor with Msh2 inactivation may be sensitive to these therapies [97, 141, 23, 57].

Disease summary: Alterations in MSH2 or other mismatch repair genes (such as MLH1, MSH6, and PMS2) have been reported to underlie hereditary nonpolyposis colorectal cancer (HNPCC). Additionally, research suggests that carriers of MSH2 mutations have an increased risk of developing colorectal cancer [79, 44, 1].

Molecular function: The MSH2 frameshift alteration reported here is expected to effectively truncate the Msh2 protein, resulting in the loss of a portion of the C-terminal domain, which is involved in MutS dimer formation (InterPro). Truncating mutations in the C-terminal domain have been reported as germline alterations in Lynch syndrome families, and truncation of the C-terminal 60 amino acids of Msh2 has been reported to result in reduced mismatch repair capacity [191]. Therefore, this mutation is predicted to lead to a loss of Msh2 function.

Incidence: MSH2 mutations have been reported in 5.0% (208/4169) of Colorectal carcinoma (CRC) samples analyzed in COSMIC (May 2020). MSH2 mutations have been reported in 1.9-2.2% of Colorectal carcinoma (CRC) samples (cBioPortal for Cancer Genomics, May 2020). Scientific studies have reported MSH2 mutation in approximately 2-12% of colorectal carcinoma specimens analyzed [202, 2, 84, 132].

Role in disease: Defective mismatch repair (MMR), occurring as a result of mutation(s) in the MMR family (MLH1, MSH2, MSH6, or PMS2) can result in microsatellite instability (MSI), common in colon, endometrium and stomach cancers [117]. MSH2 alterations have been reported to underlie hereditary nonpolyposis colorectal cancer (HNPCC) and research suggests that carriers of MSH2 mutations have an increased risk of developing colorectal cancer [79, 44, 1].

Diagnostic significance: Unknown.

Prognostic significance: Unknown.

Drug sensitivity: While there are currently no approved therapies directly addressing loss or mutation in MSH2, PD-1/PD-L1 inhibitors have been reported to be effective in tumors harboring mismatch repair defects; thus, a tumor with Msh2 inactivation may be sensitive to PD-1/PD-L1 inhibitors [97, 141, 23, 57].

Drug resistance: None.

Approved Drugs: None.

Phase 3: The second interim analysis of the Phase 3 KEYNOTE-177 study comparing 1:1 pembrolizumab with investigator choice of chemotherapy in 307 patients with MSI-high or deficient MMR metastatic colorectal carcinoma has reported a progression-free survival of 16.5 and 8.2 months, a confirmed overall response rate of 43.8% and 33.1%, an estimated restricted mean survival of 13.7 and 10.8 months, and grade 3-5 serious adverse events in 22% and 66% of patients treated with pembrolizumab or chemotherapy, respectively. Ongoing responses at 24 months were observed in 83% and 35% of patients treated with pembrolizumab or chemotherapy, respectively [8]. A Phase 3 trial (IMblaze370) of atezolizumab with cobimetinib (AC), atezolizumab monotherapy (AM), or regorafenib in patients with metastatic colorectal carcinoma reported median overall survival of 8.87, 7.1, and 8.51 months in the AC, AM, and regorafenib arms, respectively. Treatment-related grade 3-4 adverse events were reported in 61% (109/179), 31% (28/90), and 58% (46/80), respectively [45].

Phase 2: A safety and efficacy study of pembrolizumab in 149 patients with tumors with high microsatellite instability (MSI-H) or deficient in mismatch repair (dMMR) across five Phase 1 and 2 uncontrolled trials and including 15 different cancer types has reported a complete or partial response in 39.6% of patients; 78% of patients experienced response for six months or more [116, 102, 97]. A Phase 2 study of pembrolizumab in patients with advanced dMMR cancers, including 12 different tumor types, has reported complete and partial responses in 23.1% (18/78) and 35.9% (28/78) of evaluable patients, respectively [96]. Long-term followup of the Phase 2 CheckMate 142 study of nivolumab plus ipilimumab in dMMR or MSI-H CRC patients who had received at least one prior therapy reported an investigator-assessed overall response rate of 58% and a disease control rate of 81% at a median followup of 25.4 months. Complete and partial responses were reported in 6% (7/119) and 52% (62/119) of patients; median progression-free and overall survival rates at 24 months were 60% and 74%. Grade 3-4 adverse events were reported in 31% of patients. A Phase 2 study of nivolumab plus ipilimumab with radiotherapy in 40 metastatic microsatellite stable colorectal carcinoma (CRC) patients and 25 metastatic pancreatic ductal adenocarcinoma (PDAC) patients with progression on previous lines of therapy has reported disease control in 25% (10/40) and 20% (5/25), and overall response in 10% (4/40) and 13% (3/25) of patients in the CRC and PDAC cohorts, respectively. Grade 3 or higher treatment-related adverse events were observed in 40% (26/65) of patients, with grade 5 events in 3.1% (2/65) of patients. The Phase 2 GERCOR NIPICOL study of nivolumab plus ipilimumab in 57 pretreated patients with MSI-H/dMMR metastatic colorectal cancer has reported 12-week disease control rate of 86.0% and 87.7%,

Variants of potential clinical significance (3)

and 12-month progression-free survival rate of 72.9% and 76.5%, according to RECIST 1.1 and iRECIST criteria, respectively. Overall response rate was 59.7% [33]. The Phase 2 NICHE study of neoadjuvant ipilimumab and nivolumab in early-stage colon cancer patients has reported pathologic response in 100% (20/20) of patients with MMR-deficient tumors and in 27% (4/15) of patients with MMR-proficient tumors [25]. A Phase 2 study of nivolumab in 74 dMMR or MSI-H CRC patients has reported investigator and independent radiology review committee objective response rates of 27% and 31% and disease control rates of 62% and 69% with a time to response of approximately 2.7 months; progression-free survival rates at 12 months were 46% and 48% with 83.4% overall survival at six months and 73.8% at 12 months [129]. A Phase 2 study of nivolumab in combination with trifluridine/tipiracil in 18 patients with MSS colorectal carcinoma has reported 13 grade 3 or higher adverse events, no tumor responses, stable disease in eight and ten patients, and a median progression-free survival of 2.2 and 2.8 months per immune-related response criteria (irRC) and RECIST, respectively [131]. A Phase 2 study of nivolumab with ipilimumab in 40 patients with MSS colorectal adenocarcinoma reported a disease control rate of 25%, and an objective response rate of 10% by intent to treat analysis. A Phase 2 trial of pembrolizumab with cyclophosphamide and the colon vaccine GVAX in 17 patients with advanced MMR-proficient colorectal carcinoma reported disease control rates of 18% and 29% by RECIST and irRC, respectively. No objective responses were reported; median progression-free and overall survival were 2.7 and 7.1 months. Grade 3-4 adverse events attributed to study therapy were reported in 11.8% (2/17) of patients [198]. A Phase 2 trial of durvalumab with tremelimumab (DT) plus best supportive care (BSC) or BSC alone in 179 patients with advanced/refractory colorectal carcinoma reported median progression-free survival of 1.8 and 1.9 months, and median overall survival of 6.6 and 4.1 months in the DT+BSC and BSC alone arms, respectively. Disease control rates were 22.7% and 6.6%, respectively [28]. A Phase 2 study of durvalumab and trametinib in 29 microsatellite stable metastatic colorectal cancer patients has reported objective response in 3.4% and stable disease in 24% of patients. No grade 4 treatment-related adverse events were observed. The Phase 2 AVETUX study of avelumab plus cetuximab in combination with FOLFOX in 39 previously untreated metastatic colorectal cancer patients has reported overall response rate of 79.5%, with six complete and 25 partial responses, and disease control rate of 92.3%. The Phase 2 CAVE study of avelumab and cetuximab in RAS wild-type metastatic colorectal carcinoma patients with response to first-line chemotherapy in combination with anti-Egfr therapy has reported complete response in 1.5%, partial response in 4.6%, and stable disease in 49.2% of 65 evaluable patients. Median progression-free and overall survival were 3.6 months and 13.1 months, respectively, and grade 3 adverse events were observed in 22% (16/77) of patients. A Phase 2 study of avelumab monotherapy in 33 metastatic or unresectable colorectal carcinoma patients with dMMR/MSI-H or POLE mutations has reported objective response rate of 24.2% and median progression-free survival and overall survival of 3.9 and 13.2 months, respectively [88]. A Phase 2 study of short-course radiation followed by mFOLFOX6 with avelumab in 13 patients with locally advanced rectal cancer has reported pathologic complete response (pCR) in 3/12 and near pCR in 3/12 evaluable patients. No grade 4 serious adverse events were observed [156].

Phase 1: A Phase 1 study of durvalumab plus monalizumab in 55 solid tumor patients has reported confirmed partial response in three and stable disease in 11 patients in the expansion cohort of 40 metastatic microsatellite-stable colorectal cancer patients, with disease control rate at 16 weeks of 24%. A Phase 1 study of oleclumab and durvalumab in 66 solid tumor patients, with an expansion cohort in 41 advanced microsatellite-stable colorectal cancer (MSS-CRC) and pancreatic cancer patients has reported partial response in 5% (1/21) and 10% (2/20), and stable disease in 10% (2/21) and 15% (3/20) of MSS-CRC and pancreatic cancer patients, respectively. A Phase 1b trial of atezolizumab in combination with bevacizumab in refractory metastatic colorectal cancer patients reported an unconfirmed overall response rate (ORR) of 8% (1/13) and grades 3/4 adverse events in 64% of cases. In oxaliplatin-naïve patients treated with atezolizumab in combination with bevacizumab and FOLFOX, the unconfirmed ORR in evaluable patients was 36% (9/25), with 73% of cases reporting grades 3/4 adverse events. A Phase 1b trial of atezolizumab in combination with bevacizumab in microsatellite instability (MSI)-high metastatic colorectal cancer patients has reported partial response in 30% (3/10) and stable disease in 60% (6/10) of patients. Grade 3/4 adverse events were observed in 40% of patients. An ongoing Phase 1b trial of atezolizumab in combination with cobimetinib in 84 chemotherapy-refractory or locally advanced metastatic colorectal cancer patients has reported an overall response rate of 8%, including patients with MSS and MSI-low status, and disease control rate of 31%, with median progression-free and overall survival of 1.9 and 10.0 months, respectively.

Preclinical: N/A: Preclinical data are not presented when higher level data are available.

Variants of uncertain significance (6)

Gene	Variant	Allelic fraction	Classification
DNMT3A	c.557C>T p.P186L	20.0% (of 606 reads)	Tier 3, Uncertain Significance
INPP4A	c.2792G>A p.C931Y	22.0% (of 459 reads)	Tier 3, Uncertain Significance
NTRK1	c.253C>T p.R85C	16.0% (of 680 reads)	Tier 3, Uncertain Significance
PAX8	c.172G>A p.V58I	17.0% (of 1014 reads)	Tier 3, Uncertain Significance
PIK3C2B	c.1771G>A p.A591T	18.0% (of 997 reads)	Tier 3, Uncertain Significance
RAD54L	c.1345T>C p.S449P	49.0% (of 790 reads)	Tier 3, Uncertain Significance

REPORT INFORMATION

Genes tested (523)

DIGER1, DHX15, DDX41, DDR2, DCUN1D1, DAXX, CYLD, CXCR4, CUX1, CUL3, CTNNB1, CTNNA1, CTLA4, CTCF, CSNK1A1, CSF3R, CSF1R, CRLF2, CRKL, CREBBP, CIC, CHEK2, CHEK1, CHD4, CHD2, CENPA, CEBPA, CDKN2C, CDKN2B, CDKN2A, CDKN1B, CDKN1A, CDK8, CDK6, CDK4, CDK12, CDH1, CDC73, CD79B, CD79A, CD74, CD276, CD274, CCNE1, CCND3, CCND2, CCND1, CBL, CBFB, CASP8, CARD11, CALR, BTK, BTG1, BRIP1, BRD4, BRCA2, BRCA1, BRAF, BMPR1A, BLM, BIRC3, BCR, BCORL1, BCOR, BCL6, BCL2L1, BCL2L11, BCL2L1, BCL2, BCL10, BBC3, BARD1, BAP1, B2M, AXL, AXIN2, AXIN1, AURKB, AURKA, ATRX, ATR, ATM, ASXL2, ASXL1, ARID5B, ARID2, ARID1B, ARID1A, ARFRP1, ARAF, AR, APC, ANKRD26, ANKRD11, ALOX12B, ALK, AKT3, AKT2, AKT1, ACVR1B, ACVR1, ABL2, ABL1, H3-5, H3-3B, H3-3A, GSK3B, GRM3, GRIN2A, GREM1, GPS2, ADGRA2, GNAS, GNAQ, GNA13, GNA11, GLI1, GID4, GEN1, GATA6, GATA4, GATA3, GATA2, GATA1, GABRA6, FYN, FUBP1, FRS2, FOXP1, FOXO1, FOXL2, FOXA1, FLT4, FLT3, FLT1, FLI1, FLCN, FH, FGFR4, FGFR3, FGFR2, FGFR1, FGF9, FGF8, FGF7, FGF6, FGF5, FGF4, FGF3, FGF23, FGF2, FGF19, FGF14, FGF10, FGF1, FBXW7, FAT1, FAS, FANCL, FANCI, FANCG, FANCF, FANCE, FANCD2, FANCC, FANCA, TENT5C, ABRAXAS1, AMER1, EZH2, EWSR1, ETV6, ETV5, ETV4, ETV1, ETS1, ESR1, ERFF1, ERG, ERCC5, ERCC4, ERCC3, ERCC2, ERCC1, ERBB4, ERBB3, ERBB2, EPHB1, EPHA7, EPHA5, EPHA3, EPCAM, EP300, EMSY, EML4, EIF4E, EIF4A2, EIF1AX, EGFR, EGFL7, EED, E2F3, DOT1L, DNMT3B, DNMT3A, DNMT1, DNAJB1, DIS3, MST1R, MST1, MSH6, MSH3, MSH2, MRE11, MPL, MLLT3, KMT2A, MLH1, MITF, MGA, MET, MEN1, MEF2B, MED12, MDM4, MDM2, MDC1, MCL1, MAX, MAPK3, MAPK1, MAP3K4, MAP3K14, MAP3K13, MAP3K1, MAP2K4, MAP2K2, MAP2K1, MALT1, MAGI2, LZTR1, LYN, LRP1B, LMO1, LATS2, LATS1, LAMP1, KRAS, KMT2D, KMT2C, KMT2B, KLHL6, KLF4, KIT, KIF5B, KEL, KEAP1, KDR, KDM6A, KDM5C, KDM5A, KAT6A, JUN, JAK3, JAK2, JAK1, IRS2, IRS1, IRF4, IRF2, INSR, INPP4B, INPP4A, INHBA, INHA, IL7R, IL10, IKZF1, IKBKE, IGF2, IGF1R, IGF1, IFNGR1, IDH2, IDH1, ID3, ICOSLG, HSP90AA1, HSD3B1, HRAS, HOXB13, HNRNPB, HNF1A, HLA-C, HLA-B, HLA-A, H3-4, H3C13, H3C14, H3C15, H3C12, H3C11, H3C10, H3C8, H3C7, H3C6, H3C4, H3C3, H3C2, H3C1, H2BC5, H1-2, HGF, RBM10, RB1, RASA1, RARA, RANBP2, RAF1, RAD54L, RAD52, RAD51D, RAD51C, RAD51B, RAD51, RAD50, RAD21, RAC1, RAB35, QKI, PTPRT, PTPRS, PTPRD, PTPN11, PTEN, PTCH1, PRSS8, PRKDC, PRKCI, PRKAR1A, PREX2, PRDM1, PPP6C, PPP2R2A, PPP2R1A, PPM1D, PPARG, POLE, POLD1, PNRC1, PMS2, PMS1, PMAIP1, PLK2, PLCG2, PIM1, PIK3R3, PIK3R2, PIK3R1, PIK3CG, PIK3CD, PIK3CB, PIK3CA, PIK3C3, PIK3C2G, PIK3C2B, PHOX2B, PHF6, PGR, PDPK1, PDK1, PDGFRB, PDGFRA, PDCD1LG2, PDCD1, PBRM1, PAX8, PAX7, PAX5, PAX3, PARP1, PRKN, PALB2, PAK5, PAK3, PAK1, NUTM1, NUP93, NTRK3, NTRK2, NTRK1, NSD1, NRG1, NRAS, NPM1, NOTCH4, NOTCH3, NOTCH2, NOTCH1, NKX3-1, NKX2-1, NFKBIA, NFE2L2, NF2, NF1, NEGR1, NCOR1, NCOA3, NBN, NAB2, MYO11, MYD88, MYCN, MYCL, MYC, MYB, MUTYH, MTOR, ZRSR2, ZNF703, ZNF217, ZFH3, ZBTB7A, ZBTB2, YES1, YAP1, XRCC2, XPO1, XIAP, WT1, CCN6, VTCN1, VHL, VEGFA, U2AF1, TSHR, TSC2, TSC1, TRAF7, TRAF2, TP63, TP53, TOP2A, TOP1, TNFRSF14, TNFAIP3, TMPRSS2, TMEM127, TGFB2, TGFB1, TFRC, TFE3, TET2, TET1, TERT, TERC, TCF7L2, TCF3, ELOC, TBX3, TAF1, SYK, SUZ12, SUFU, STK40, STK11, STAT5B, STAT5A, STAT4, STAT3, STAG2, STAG1, SRSF2, SRC, SPTA1, SPOP, SPEN, SOX9, SOX2, SOX17, SOX10, SOCS1, SNCAIP, SMO, SMC3, SMC1A, SMARCD1, SMARCB1, SMARCA4, SMAD4, SMAD3, SMAD2, SLX4, SLIT2, SHQ1, SH2D1A, SH2B3, SF3B1, SETD2, SETBP1, SDHB, SDHC, SDHB, SDHAF2, SDHA, RYBP, RUNX1T1, RUNX1, RPTOR, RPS6KB2, RPS6KB1, RPS6KA4, ROS1, RNF43, RIT1, RICTOR, RHOA, RHEB, COP1, RET, REL, RECQL4

Methods and limitations

QIAGEN Clinical Insight (QCI™) is a variant analysis, interpretation and decision support tool for research and clinical labs analyzing human genetics data and is not intended to be used for diagnostic purposes. QCI Interpret software includes the following underlying databases, data reference sets and tools; QIAGEN Clinical Insight-Interpret (7.1.20210316), Ingenuity Knowledge Base (B-release), CADD (v1.6), Allele Frequency Community (2019-09-25), EVS (ESP6500SI-V2), Refseq Gene Model (2020-04-06), JASPAR (2013-11), Ingenuity Knowledge Base Snapshot Timestamp (2021-03-18 14:15:50.073), Vista Enhancer hg18 (2012-07), Vista Enhancer hg19 (2012-07), Clinical Trials (B-release), MITOMAP: A Human Mitochondrial Genome Database. <http://www.mitomap.org>, 2019 (2020-06-19), PolyPhen-2 (v2.2.2), 1000 Genome Frequency (phase3v5b), ExAC (0.3.1), iva (Nov 20 02:39), TargetScan (7.2), phyloP hg18 (NCBI36 (hg18) 2009-11, GRCh37 (hg19) 2014-02, GRCh38 2015-05), phyloP hg19 (NCBI36 (hg18) 2009-11, GRCh37 (hg19) 2014-02, GRCh38 2015-05), GENCODE (Release 33), CentoMD (5.3), OMIM (July 06, 2020), gnomAD (2.1.1), BSIFT (2016-02-23), TCGA (2013-09-05), Clinvar (2020-09-15), DGV (2016-05-15), COSMIC (v92), HGMD (2020.4), OncoTree (oncotree_2019_03_01), dbSNP (NCBI36 (hg18) 151, GRCh37 (hg19) 153, GRCh38 153), SIFT4G (2016-02-23)

Clinical significance of variants based on AMP / ASCO / CAP guidelines*

Strong clinical significance	Tier 1A	Biomarker predicts response or resistance to an FDA or EMA approved therapy, according to drug label or professional guidelines for this diagnosis Biomarker included in professional guidelines is prognostic or diagnostic for this diagnosis
	Tier 1B	Biomarker predicts response or resistance to a therapy for this diagnosis based on well-powered studies Biomarker is prognostic or diagnostic for this diagnosis based on well-powered studies
Potential clinical significance	Tier 2C	Biomarker is associated with response or resistance to an FDA or EMA approved therapy, according to drug label or professional guidelines but only for different diagnosis Biomarker is an inclusion criterion for an active clinical trial Biomarker is prognostic or diagnostic based on multiple small studies
	Tier 2D	Biomarker shows plausible response or resistance based on case or preclinical studies Biomarker may assist in disease diagnosis or prognosis based on small studies
Uncertain clinical significance	Tier 3	Biomarker has uncertain clinical significance and not known to be likely benign or benign

*Adapted from PMID:27993330 [jmd.amjpathol.org/article/S1525-1578\(16\)30223-9/pdf](https://pubmed.ncbi.nlm.nih.gov/27993330/)

SELECTED REFERENCES

1. Aaltonen LA, Peltomäki P (1994) Genes involved in hereditary nonpolyposis colorectal carcinoma. *Anticancer Res* 1994 Jul-Aug;14(4B):1657-60 (PMID: 7979203)

2. Abdul Murad NA, Othman Z, Khalid M, Abdul Razak Z, Hussain R, Nadesan S, Sagap I, Mohamed Rose I, Wan Ngah WZ, Jamal R (2012) Missense mutations in MLH1, MSH2, KRAS, and APC genes in colorectal cancer patients in Malaysia. *Dig Dis Sci.* 2012 Nov;57(11):2863-72. Epub 2012 Jun 6 ([PMID: 22669205](#))
3. Agarwal S, van Cappellen WA, Guénolé A, Eppink B, Linsen SE, Meijering E, Houtsmuller A, Kanaar R, Essers J (2011) ATP-dependent and independent functions of Rad54 in genome maintenance. *J Cell Biol.* 2011 Mar 07;192(5):735-50. Epub 2011 Feb 28 ([PMID: 21357745](#))
4. Alberti L, Carniti C, Miranda C, Roccato E, Pierotti MA (2003) RET and NTRK1 proto-oncogenes in human diseases. *J Cell Physiol.* 2003 May;195(2):168-86 ([PMID: 12652644](#))
5. Alexander J, Watanabe T, Wu TT, Rashid A, Li S, Hamilton SR (2001) Histopathological identification of colon cancer with microsatellite instability. *Am J Pathol* 2001 Feb;158(2):527-35 ([PMID: 11159189](#))
6. Alexandrov LB, Nik-Zainal S, Wedge DC, Aparicio SA, Behjati S, Biankin AV, Bignell GR, Bolli N, Borg A, Børresen-Dale AL, Boyault S, Burkhardt B, Butler AP, Caldas C, Davies HR, Desmedt C, Eils R, Eyfjörd JE, Foekens JA, Greaves M, Hosoda F, Hutter B, Ilicic T, Imbeaud S, Imielinski M, Jäger N, Jones DT, Jones D, Knappskog S, Kool M, Lakhani SR, López-Otín C, Martin S, Munshi NC, Nakamura H, Northcott PA, Pajic M, Papaemmanuil E, Paradiso A, Pearson JV, Puente XS, Raine K, Ramakrishna M, Richardson AL, Richter J, Rosenstiel P, Schlesner M, Schumacher TN, Span PN, Teague JW, Totoki Y, Tutt AN, Valdés-Mas R, van Buuren MM, van 't Veer L, Vincent-Salomon A, Waddell N, Yates LR, Australian Pancreatic Cancer Genome Initiative, ICGC Breast Cancer Consortium, ICGC MMML-Seq Consortium, ICGC PedBrain, Zucman-Rossi J, Futreal PA, McDermott U, Lichter P, Meyerson M, Grimmond SM, Siebert R, Campo E, Shibata T, Pfister SM, Campbell PJ, Stratton MR (2013) Signatures of mutational processes in human cancer. *Nature.* 2013 Aug 22;500(7463):415-21. Epub 2013 Aug 14 ([PMID: 23945592](#))
7. Allo G, Bernardini MQ, Wu RC, Shih IeM, Kalloger S, Pollett A, Gilks CB, Clarke BA (2013) ARID1A loss correlates with mismatch repair deficiency and intact p53 expression in high-grade endometrial carcinomas. *Mod Pathol.* 2014 Feb;27(2):255-61. Epub 2013 Jul 26 ([PMID: 23887303](#))
8. André T, Shiu KK, Kim TW, Jensen BV, Jensen LH, Punt C, Smith D, Garcia-Carbonero R, Benavides M, Gibbs P, de la Fouchardiere C, Rivera F, Elez E, Bendell J, Le DT, Yoshino T, Van Cutsem E, Yang P, Farooqui MZH, Marinello P, Diaz LA Jr (2020) Pembrolizumab in Microsatellite-Instability-High Advanced Colorectal Cancer. *N Engl J Med* 2020 Dec 3;383(23):2207-2218 ([PMID: 33264544](#))
9. Bai H, Wang R, Cheng W, Shen Y, Li H, Xia W, Ding Z, Zhang Y (2020) Evaluation of Concordance Between Deficient Mismatch Repair and Microsatellite Instability Testing and Their Association with Clinicopathological Features in Colorectal Cancer. *Cancer Manag Res* 2020;12:2863-2873 ([PMID: 32425600](#))
10. Berns K, Caumanns JJ, Hijmans EM, Gennissen AMC, Severson TM, Evers B, Wisman GBA, Jan Meersma G, Liefink C, Beijersbergen RL, Itamochi H, van der Zee AGJ, de Jong S, Bernards R (2018) ARID1A mutation sensitizes most ovarian clear cell carcinomas to BET inhibitors. *Oncogene* 2018 Aug;37(33):4611-4625 ([PMID: 29760405](#))
11. Bitler BG, Aird KM, Garipov A, Li H, Amatangelo M, Kossenkov AV, Schultz DC, Liu Q, Shih IeM, Conejo-Garcia JR, Speicher DW, Zhang R (2015) Synthetic lethality by targeting EZH2 methyltransferase activity in ARID1A-mutated cancers. *Nat Med.* 2015 Mar;21(3):231-8. Epub 2015 Feb 16 ([PMID: 25686104](#))
12. Boland CR, Thibodeau SN, Hamilton SR, Sidransky D, Eshleman JR, Burt RW, Meltzer SJ, Rodriguez-Bigas MA, Fodde R, Ranzani GN, Srivastava S (1998) A National Cancer Institute Workshop on Microsatellite Instability for cancer detection and familial predisposition: development of international criteria for the determination of microsatellite instability in colorectal cancer. *Cancer Res* 1998 Nov 15;58(22):5248-57 ([PMID: 9823339](#))
13. Boller D, Doepfner KT, De Laurentiis A, Guerreiro AS, Marinov M, Shalaby T, Depledge P, Robson A, Saghir N, Hayakawa M, Kaizawa H, Koizumi T, Ohishi T, Fattet S, Delattre O, Schweri-Olac A, Höland K, Grotzer MA, Frei K, Spertini O, Waterfield MD, Arcaro A (2012) Targeting PI3K2β impairs proliferation and survival in acute leukemia, brain tumours and neuroendocrine tumours. *Anticancer Res.* 2012 Aug;32(8):3015-27 ([PMID: 22843869](#))
14. Bong IPN, Ng CC, Baharuddin P, Zakaria Z (2017) MicroRNA expression patterns and target prediction in multiple myeloma development and malignancy. *Genes Genomics.* 2017;39(5):533-540. Epub 2017 Feb 9 ([PMID: 28458781](#))
15. Bouchard M, Souabni A, Mandler M, Neubüser A, Busslinger M (2002) Nephric lineage specification by Pax2 and Pax8. *Genes Dev.* 2002 Nov 15;16(22):2958-70 ([PMID: 12435636](#))
16. Brown RA, Ho LK, Weber-Hall SJ, Shipley JM, Fry MJ (1997) Identification and cDNA cloning of a novel mammalian C2 domain-containing phosphoinositide 3-kinase, HsC2-PI3K. *Biochem Biophys Res Commun.* 1997 Apr 17;233(2):537-44 ([PMID: 9144573](#))
17. Brown SD, Warren RL, Gibb EA, Martin SD, Spinelli JJ, Nelson BH, Holt RA (2014) Neo-antigens predicted by tumor genome meta-analysis correlate with increased patient survival. *Genome Res* 2014 May;24(5):743-50 ([PMID: 24782321](#))
18. Błajęcka K, Marinov M, Leitner L, Uth K, Posern G, Arcaro A (2012) Phosphoinositide 3-kinase C2β regulates RhoA and the actin cytoskeleton through an interaction with Dbl. *PLoS One.* 2012;7(9):e44945. Epub 2012 Sep 12 ([PMID: 22984590](#))
19. Cajuso T, Hänninen UA, Kondelin J, Gylfe AE, Tanskanen T, Katainen R, Pitkänen E, Ristolainen H, Kaasinen E, Taipale M, Taipale J, Böhm J, Renkonen-Sinisalo L, Mecklin JP, Järvinen H, Tuupanen S, Kilpivaara O, Vahteristo P (2014) Exome sequencing reveals frequent inactivating mutations in ARID1A, ARID1B, ARID2 and ARID4A in microsatellite unstable colorectal cancer. *Int J Cancer.* 2014 Aug 01;135(3):611-23. Epub 2014 Jan 13 ([PMID: 24382590](#))
20. Campesato LF, Barroso-Sousa R, Jimenez L, Correa BR, Sabbaga J, Hoff PM, Reis LF, Galante PA, Camargo AA (2015) Comprehensive cancer-gene panels can be used to estimate mutational load and predict clinical benefit to PD-1 blockade in clinical practice. *Oncotarget.* 2015 Oct 27;6(33):34221-7 ([PMID: 26439694](#))
21. Cancer Genome Atlas Network (2012) Comprehensive molecular characterization of human colon and rectal cancer. *Nature.* 2012 Jul 18;487(7407):330-7 ([PMID: 22810696](#))
22. Carbone DP, Reck M, Paz-Ares L, Creelan B, Horn L, Steins M, Felip E, van den Heuvel MM, Ciuleanu TE, Badin F, Ready N, Hiltermann T, Nair S, Juergens R, Peters S, Minenza E, Wrangle JM, Rodriguez-Abreu D, Borghaei H, Blumenschein GR, Villaruz LC, Havel L, Krejci J, Corral Jaime J, Chang H, Geese WJ, Bhagavatheswaran P, Chen AC, Socinski MA, CheckMate 026 Investigators (2017) First-Line Nivolumab in Stage IV or Recurrent Non-Small-Cell Lung Cancer. *N Engl J Med.* 2017 Jun 22;376(25):2415-2426 ([PMID: 28636851](#))
23. Castro MP, Goldstein N (2015) Mismatch repair deficiency associated with complete remission to combination programmed cell death ligand immune therapy in a patient with sporadic urothelial carcinoma: immunotherapeutic considerations. *J Immunother Cancer.* 2015;3:58. Epub 2015 Dec 15 ([PMID: 26674132](#))

24. Caumanns JJ, Wisman GBA, Berns K, van der Zee AGJ, de Jong S (2018) ARID1A mutant ovarian clear cell carcinoma: A clear target for synthetic lethal strategies. *Biochim Biophys Acta Rev Cancer* 2018 Dec;1870(2):176-184 ([PMID: 30025943](#))
25. Chalabi M, Fanchi LF, Dijkstra KK, Van den Berg JG, Aalbers AG, Sikorska K, Lopez-Yurda M, Grootsholten C, Beets GL, Snaebjornsson P, Maas M, Mertz M, Veninga V, Bounova G, Broeks A, Beets-Tan RG, de Wijkerslooth TR, van Lent AU, Marsman HA, Nuijten E, Kok NF, Kuiper M, Verbeek WH, Kok M, Van Leerdam ME, Schumacher TN, Voest EE, Haanen JB (2020) Neoadjuvant immunotherapy leads to pathological responses in MMR-proficient and MMR-deficient early-stage colon cancers. *Nat Med* 2020 Apr;26(4):566-576 ([PMID: 32251400](#))
26. Chandler RL, Brennan J, Schisler JC, Serber D, Patterson C, Magnuson T (2012) ARID1a-DNA interactions are required for promoter occupancy by SWI/SNF. *Mol Cell Biol*. 2013 Jan;33(2):265-80. Epub 2012 Nov 5 ([PMID: 23129809](#))
27. Chang CC, Lin PC, Lin CC, Lan YT, Lin HH, Lin CH, Yang SH, Liang WY, Chen WS, Jiang JK, Lin JK, Chang SC (2017) Molecular and Clinicopathological Differences by Age at the Diagnosis of Colorectal Cancer. *Int J Mol Sci*. 2017 Jul 05;18(7). Epub 2017 Jul 5 ([PMID: 28678173](#))
28. Chen EX, Jonker DJ, Loree JM, Kennecke HF, Berry SR, Couture F, Ahmad CE, Goffin JR, Kavan P, Harb M, Colwell B, Samimi S, Samson B, Abbas T, Aucoin N, Aubin F, Koski SL, Wei AC, Magoski NM, Tu D, O'Callaghan CJ (2020) Effect of Combined Immune Checkpoint Inhibition vs Best Supportive Care Alone in Patients With Advanced Colorectal Cancer: The Canadian Cancer Trials Group CO.26 Study. *JAMA Oncol* 2020 Jun 1; 6(6):831-838 ([PMID: 32379280](#))
29. Chen JF, Luo X, Xiang LS, Li HT, Zha L, Li N, He JM, Xie GF, Xie X, Liang HJ (2016) EZH2 promotes colorectal cancer stem-like cell expansion by activating p21cip1-Wnt/ β -catenin signaling. *Oncotarget* 2016 Jul 5;7(27):41540-41558 ([PMID: 27172794](#))
30. Chen MH, Chang SC, Lin PC, Yang SH, Lin CC, Lan YT, Lin HH, Lin CH, Lai JI, Liang WY, Lu ML, Yang MH, Chao Y (2019) Combined Microsatellite Instability and Elevated Microsatellite Alterations at Selected Tetranucleotide Repeats (EMAST) Might Be a More Promising Immune Biomarker in Colorectal Cancer. *Oncologist* 2019 Dec;24(12):1534-1542 ([PMID: 31292272](#))
31. Chou A, Toon CW, Clarkson A, Sioson L, Houang M, Watson N, DeSilva K, Gill AJ (2014) Loss of ARID1A expression in colorectal carcinoma is strongly associated with mismatch repair deficiency. *Hum Pathol*. 2014 Aug;45(8):1697-703. Epub 2014 Apr 24 ([PMID: 24925223](#))
32. Chédin F (2011) The DNMT3 family of mammalian de novo DNA methyltransferases. *Prog Mol Biol Transl Sci*. 2011;101:255-85 ([PMID: 21507354](#))
33. Cohen R, Bennouna J, Meurisse A, Tournigand C, De La Fouchardière C, Tougeron D, Borg C, Mazard T, Chibaudel B, Garcia-Larnicol ML, Svrcek M, Vernerey D, Menu Y, André T (2020) RECIST and iRECIST criteria for the evaluation of nivolumab plus ipilimumab in patients with microsatellite instability-high/mismatch repair-deficient metastatic colorectal cancer: the GERCOR NIPICOL phase II study. *J Immunother Cancer*. 2020 Nov;8(2) ([PMID: 33148693](#))
34. Colas C, Coulet F, Svrcek M, Collura A, Fléjou JF, Duval A, Hamelin R (2012) Lynch or not Lynch? Is that always a question? *Adv Cancer Res*. 2012; 113:121-66 ([PMID: 22429854](#))
35. Collingwood TN, Urnov FD, Wolffe AP (1999) Nuclear receptors: coactivators, corepressors and chromatin remodeling in the control of transcription. *J Mol Endocrinol* 1999 Dec;23(3):255-75 ([PMID: 10601972](#))
36. Copeland RA (2013) Molecular pathways: protein methyltransferases in cancer. *Clin Cancer Res* 2013 Dec 1;19(23):6344-50 ([PMID: 23958745](#))
37. Cui W, Cai Y, Wang W, Liu Z, Wei P, Bi R, Chen W, Sun M, Zhou X (2014) Frequent copy number variations of PI3K/AKT pathway and aberrant protein expressions of PI3K subunits are associated with inferior survival in diffuse large B cell lymphoma. *J Transl Med*. 2014 Jan 13;12:10 ([PMID: 24418330](#))
38. Czito BG, Deming DA, Jameson GS, Mulcahy MF, Vaghefi H, Dudley MW, Holen KD, DeLuca A, Mittapalli RK, Munasinghe W, He L, Zalcborg JR, Ngan SY, Komarnitsky P, Michael M (2017) Safety and tolerability of veliparib combined with capecitabine plus radiotherapy in patients with locally advanced rectal cancer: a phase 1b study. *Lancet Gastroenterol Hepatol* 2017 Jun;2(6):418-426 ([PMID: 28497757](#))
39. Das M, Scappini E, Martin NP, Wong KA, Dunn S, Chen YJ, Miller SL, Domin J, O'Bryan JP (2007) Regulation of neuron survival through an intersectin-phosphoinositide 3'-kinase C2beta-AKT pathway. *Mol Cell Biol*. 2007 Nov;27(22):7906-17. Epub 2007 Sep 17 ([PMID: 17875942](#))
40. Daskalos A, Oleksiewicz U, Filia A, Nikolaidis G, Xinarianos G, Gosney JR, Malliri A, Field JK, Liloglou T (2010) UHRF1-mediated tumor suppressor gene inactivation in nonsmall cell lung cancer. *Cancer*. 2011 Mar 01;117(5):1027-37. Epub 2010 Nov 8 ([PMID: 21351083](#))
41. Di Palma T, Lucci V, de Cristofaro T, Filippone MG, Zannini M (2014) A role for PAX8 in the tumorigenic phenotype of ovarian cancer cells. *BMC Cancer*. 2014 Apr 26;14:292 ([PMID: 24766781](#))
42. Domin J, Harper L, Aubyn D, Wheeler M, Florey O, Haskard D, Yuan M, Zicha D (2005) The class II phosphoinositide 3-kinase PI3K-C2beta regulates cell migration by a PtdIns3P dependent mechanism. *J Cell Physiol*. 2005 Dec;205(3):452-62 ([PMID: 16113997](#))
43. Donehower LA, Creighton CJ, Schultz N, Shinbrot E, Chang K, Gunaratne PH, Muzny D, Sander C, Hamilton SR, Gibbs RA, Wheeler D (2013) MLH1-silenced and non-silenced subgroups of hypermutated colorectal carcinomas have distinct mutational landscapes. *J Pathol*. 2013 Jan;229(1): 99-110 ([PMID: 22899370](#))
44. Dowty JG, Win AK, Buchanan DD, Lindor NM, Macrae FA, Clendenning M, Antill YC, Thibodeau SN, Casey G, Gallinger S, Marchand LL, Newcomb PA, Haile RW, Young GP, James PA, Giles GG, Gunawardena SR, Leggett BA, Gattas M, Boussioutas A, Ahnen DJ, Baron JA, Parry S, Goldblatt J, Young JP, Hopper JL, Jenkins MA (2012) Cancer risks for MLH1 and MSH2 mutation carriers. *Hum Mutat* 2013 Mar;34(3):490-7 ([PMID: 23255516](#))
45. Eng C, Kim TW, Bendell J, Argilés G, Tebbutt NC, Di Bartolomeo M, Falcone A, Fakih M, Kozloff M, Segal NH, Sobrero A, Yan Y, Chang I, Uyei A, Roberts L, Ciardiello F (2019) Atezolizumab with or without cobimetinib versus regorafenib in previously treated metastatic colorectal cancer (IMblaze370): a multicentre, open-label, phase 3, randomised, controlled trial. *Lancet Oncol* 2019 Jun;20(6):849-861 ([PMID: 31003911](#))
46. Engelman JA (2009) Targeting PI3K signalling in cancer: opportunities, challenges and limitations. *Nat Rev Cancer*. 2009 Aug;9(8):550-62 ([PMID: 19629070](#))
47. Eppink B, Tafel AA, Hanada K, van Drunen E, Hickson ID, Essers J, Kanaar R (2011) The response of mammalian cells to UV-light reveals Rad54-dependent and independent pathways of homologous recombination. *DNA Repair (Amst)*. 2011 Nov 10;10(11):1095-105. Epub 2011 Aug 31 ([PMID: 21885354](#))
48. Fabbri M, Garzon R, Cimmino A, Liu Z, Zanesi N, Callegari E, Liu S, Alder H, Costinean S, Fernandez-Cymering C, Volinia S, Guler G, Morrison CD, Chan KK, Marcucci G, Calin GA, Huebner K, Croce CM (2007) MicroRNA-29 family reverts aberrant methylation in lung cancer by targeting DNA methyltransferases 3A and 3B. *Proc Natl Acad Sci U S A*. 2007 Oct 02;104(40):15805-10. Epub 2007 Sep 21 ([PMID: 17890317](#))

49. Fabrizio DA, George TJ Jr, Dunne RF, Frampton G, Sun J, Gowen K, Kennedy M, Greenbowe J, Schrock AB, Hezel AF, Ross JS, Stephens PJ, Ali SM, Miller VA, Fakhri M, Klemperer SJ (2018) Beyond microsatellite testing: assessment of tumor mutational burden identifies subsets of colorectal cancer who may respond to immune checkpoint inhibition. *J Gastrointest Oncol* 2018 Aug;9(4):610-617 ([PMID: 30151257](#))
50. Federico SM, Pappo AS, Sahr N, Sykes A, Campagne O, Stewart CF, Clay MR, Bahrami A, McCarville MB, Kaste SC, Santana VM, Helmig S, Gartrell J, Shelat A, Brennan RC, Hawkins D, Godwin K, Bishop MW, Furman WL, Stewart E (2020) A phase I trial of talazoparib and irinotecan with and without temozolomide in children and young adults with recurrent or refractory solid malignancies. *Eur J Cancer* 2020 Sep;137:204-213 ([PMID: 32795876](#))
51. Fishel R, Ewel A, Lescoe MK (1994) Purified human MSH2 protein binds to DNA containing mismatched nucleotides. *Cancer Res.* 1994 Nov 01;54(21):5539-42 ([PMID: 7923193](#))
52. Forde PM, Chafft JE, Smith KN, Anagnostou V, Cottrell TR, Hellmann MD, Zahurak M, Yang SC, Jones DR, Broderick S, Battafarano RJ, Velez MJ, Rekhtman N, Olah Z, Naidoo J, Marrone KA, Verde F, Guo H, Zhang J, Caushi JX, Chan HY, Sidhom JW, Scharpf RB, White J, Gabrielson E, Wang H, Rosner GL, Rusch V, Wolchok JD, Merghoub T, Taube JM, Velculescu VE, Topalian SL, Brahmer JR, Pardoll DM (2018) Neoadjuvant PD-1 Blockade in Resectable Lung Cancer. *N Engl J Med.* 2018 May 24;378(21):1976-1986. Epub 2018 Apr 16 ([PMID: 29658848](#))
53. Fujiyoshi K, Yamaguchi T, Kakuta M, Takahashi A, Arai Y, Yamada M, Yamamoto G, Ohde S, Takao M, Horiguchi SI, Natsume S, Kazama S, Nishizawa Y, Nishimura Y, Akagi Y, Sakamoto H, Akagi K (2017) Predictive model for high-frequency microsatellite instability in colorectal cancer patients over 50 years of age. *Cancer Med.* 2017 Jun;6(6):1255-1263. Epub 2017 May 23 ([PMID: 28544821](#))
54. Gandara DR, Paul SM, Kowanetz M, Schleifman E, Zou W, Li Y, Rittmeyer A, Fehrenbacher L, Otto G, Malboeuf C, Lieber DS, Lipson D, Siltrerra J, Amler L, Riehl T, Cummings CA, Hegde PS, Sandler A, Ballinger M, Fabrizio D, Mok T, Shames DS (2018) Blood-based tumor mutational burden as a predictor of clinical benefit in non-small-cell lung cancer patients treated with atezolizumab. *Nat Med.* 2018 Sep;24(9):1441-1448. Epub 2018 Aug 6 ([PMID: 30082870](#))
55. Gao Q, Steine EJ, Barrasa MI, Hockemeyer D, Pawlak M, Fu D, Reddy S, Bell GW, Jaenisch R (2011) Deletion of the de novo DNA methyltransferase Dnmt3a promotes lung tumor progression. *Proc Natl Acad Sci U S A.* 2011 Nov 01;108(44):18061-6. Epub 2011 Oct 19 ([PMID: 22011581](#))
56. Genther Williams SM, Kuznicki AM, Andrade P, Dolinski BM, Elbi C, O'Hagan RC, Toniatti C (2015) Treatment with the PARP inhibitor, niraparib, sensitizes colorectal cancer cell lines to irinotecan regardless of MSI/MSS status. *Cancer Cell Int* 2015;15(1):14 ([PMID: 25685067](#))
57. Ghatalia P, Nagarathinam R, Cooper H, Geynisman DM, El-Deiry WS (2017) Mismatch repair deficient metastatic colon cancer and urothelial cancer: A case report of sequential immune checkpoint therapy. *Cancer Biol Ther* 2017 Sep 2;18(9):651-654 ([PMID: 28726535](#))
58. Giannakis M, Mu XJ, Shukla SA, Qian ZR, Cohen O, Nishihara R, Bahl S, Cao Y, Amin-Mansour A, Yamauchi M, Sukawa Y, Stewart C, Rosenberg M, Mima K, Inamura K, Noshro K, Nowak JA, Lawrence MS, Giovannucci EL, Chan AT, Ng K, Meyerhardt JA, Van Allen EM, Getz G, Gabriel SB, Lander ES, Wu CJ, Fuchs CS, Ogino S, Garraway LA (2016) Genomic Correlates of Immune-Cell Infiltrates in Colorectal Carcinoma. *Cell Rep.* 2016 Apr 26;15(4):857-865. Epub 2016 Apr 14 ([PMID: 27149842](#))
59. Gekas I, Novotny J, Pecan L, Strigård K, Palmqvist R, Gunnarsson U (2017) Microsatellite Instability as a Prognostic Factor in Stage II Colon Cancer Patients, a Meta-Analysis of Published Literature. *Anticancer Res* 2017 Dec;37(12):6563-6574 ([PMID: 29187431](#))
60. Goodman AM, Kato S, Bazhenova L, Patel SP, Frampton GM, Miller V, Stephens PJ, Daniels GA, Kurzrock R (2017) Tumor Mutational Burden as an Independent Predictor of Response to Immunotherapy in Diverse Cancers. *Mol Cancer Ther.* 2017 Nov;16(11):2598-2608. Epub 2017 Aug 23 ([PMID: 28835386](#))
61. Gorbunova V, Beck JT, Hofheinz RD, Garcia-Alfonso P, Nechaeva M, Cubillo Gracian A, Mangel L, Elez Fernandez E, Deming DA, Ramanathan RK, Torres AH, Sullivan D, Luo Y, Berlin JD (2018) A phase 2 randomised study of veliparib plus FOLFIRI±bevacizumab versus placebo plus FOLFIRI±bevacizumab in metastatic colorectal cancer. *Br J Cancer* 2019 Jan;120(2):183-189 ([PMID: 30531832](#))
62. Gorbunova V, Beck JT, Hofheinz RD, Garcia-Alfonso P, Nechaeva M, Gracian AC, Mangel L, Fernandez EE, Deming DA, Ramanathan RK, Torres AH, Sullivan D, Luo Y, Berlin JD (2019) Correction: A phase 2 randomised study of veliparib plus FOLFIRI±bevacizumab versus placebo plus FOLFIRI±bevacizumab in metastatic colorectal cancer. *Br J Cancer* 2019 Aug;121(5):429-430 ([PMID: 31350526](#))
63. Greco A, Mariani C, Miranda C, Lupas A, Pagliardini S, Pomati M, Pierotti MA (1995) The DNA rearrangement that generates the TRK-T3 oncogene involves a novel gene on chromosome 3 whose product has a potential coiled-coil domain. *Mol Cell Biol.* 1995 Nov;15(11):6118-27 ([PMID: 7565764](#))
64. Greco A, Mariani C, Miranda C, Pagliardini S, Pierotti MA (1993) Characterization of the NTRK1 genomic region involved in chromosomal rearrangements generating TRK oncogenes. *Genomics.* 1993 Nov;18(2):397-400 ([PMID: 8288244](#))
65. Greco A, Pierotti MA, Bongarzone I, Pagliardini S, Lanzi C, Della Porta G (1992) TRK-T1 is a novel oncogene formed by the fusion of TPR and TRK genes in human papillary thyroid carcinomas. *Oncogene.* 1992 Feb;7(2):237-42 ([PMID: 1532241](#))
66. Gryfe R, Kim H, Hsieh ET, Aronson MD, Holowaty EJ, Bull SB, Redston M, Gallinger S (2000) Tumor microsatellite instability and clinical outcome in young patients with colorectal cancer. *N Engl J Med* 2000 Jan 13;342(2):69-77 ([PMID: 10631274](#))
67. Guan B, Wang TL, Shih IeM (2011) ARID1A, a factor that promotes formation of SWI/SNF-mediated chromatin remodeling, is a tumor suppressor in gynecologic cancers. *Cancer Res.* 2011 Nov 01;71(21):6718-27. Epub 2011 Sep 7 ([PMID: 21900401](#))
68. Han SW, Lee HJ, Bae JM, Cho NY, Lee KH, Kim TY, Oh DY, Im SA, Bang YJ, Jeong SY, Park KJ, Park JG, Kang GH, Kim TY (2012) Methylation and microsatellite status and recurrence following adjuvant FOLFOX in colorectal cancer. *Int J Cancer.* 2013 May 01;132(9):2209-16. Epub 2012 Oct 29 ([PMID: 23034738](#))
69. Hanna NH, Schneider BJ, Temin S, Baker S, Brahmer J, Ellis PM, Gaspar LE, Haddad RY, Hesketh PJ, Jain D, Jaiyesimi I, Johnson DH, Leighl NB, Phillips T, Riely GJ, Robinson AG, Rosell R, Schiller JH, Singh N, Spigel DR, Stabler JO, Tashbar J, Masters G (2020) Therapy for Stage IV Non-Small-Cell Lung Cancer Without Driver Alterations: ASCO and OH (CCO) Joint Guideline Update. *J Clin Oncol.* 2020 May 10;38(14):1608-1632. Epub 2020 Jan 28 ([PMID: 31990617](#))
70. Heinemann K (2013) Toward a molecular classification of colorectal cancer: the role of microsatellite instability status. *Front Oncol.* 2013 Oct 31;3:272 ([PMID: 24199172](#))

71. Hellmann MD, Ciuleanu TE, Pluzanski A, Lee JS, Otterson GA, Audigier-Valette C, Minenza E, Linardou H, Burgers S, Salman P, Borghaei H, Ramalingam SS, Brahmer J, Reck M, O'Byrne KJ, Geese WJ, Green G, Chang H, Szustakowski J, Bhagavatheeswaran P, Healey D, Fu Y, Nathan F, Paz-Ares L (2018) Nivolumab plus Ipilimumab in Lung Cancer with a High Tumor Mutational Burden. *N Engl J Med*. 2018 May 31;378(22):2093-2104. Epub 2018 Apr 16 ([PMID: 29658845](#))
72. Howitt BE, Shukla SA, Sholl LM, Ritterhouse LL, Watkins JC, Rodig S, Stover E, Strickland KC, D'Andrea AD, Wu CJ, Matulonis UA, Konstantinopoulos PA (2015) Association of Polymerase ϵ -Mutated and Microsatellite-Unstable Endometrial Cancers With Neoantigen Load, Number of Tumor-Infiltrating Lymphocytes, and Expression of PD-1 and PD-L1. *JAMA Oncol*. 2015 Dec;1(9):1319-23 ([PMID: 26181000](#))
73. Hung N, Chen YJ, Taha A, Olivecrona M, Boet R, Wiles A, Warr T, Shaw A, Eiholzer R, Baguley BC, Eccles MR, Braithwaite AW, Macfarlane M, Royds JA, Slatter T (2014) Increased paired box transcription factor 8 has a survival function in glioma. *BMC Cancer*. 2014 Mar 06;14:159. Epub 2014 Mar 6 ([PMID: 24602166](#))
74. Im AP, Sehgal AR, Carroll MP, Smith BD, Tefferi A, Johnson DE, Boyiadzis M (2014) DNMT3A and IDH mutations in acute myeloid leukemia and other myeloid malignancies: associations with prognosis and potential treatment strategies. *Leukemia*. 2014 Sep;28(9):1774-83. Epub 2014 Apr 4 ([PMID: 24699305](#))
75. Innocenti F, Ou FS, Qu X, Zemla TJ, Niedzwiecki D, Tam R, Mahajan S, Goldberg RM, Bertagnolli MM, Blanke CD, Sanoff H, Atkins J, Polite B, Venook AP, Lenz HJ, Kabbarah O (2019) Mutational Analysis of Patients With Colorectal Cancer in CALGB/SWOG 80405 Identifies New Roles of Microsatellite Instability and Tumor Mutational Burden for Patient Outcome. *J Clin Oncol*. 2019 May 10;37(14):1217-1227. Epub 2019 Mar 13 ([PMID: 30865548](#))
76. Italiano A, Soria JC, Toulmonde M, Michot JM, Lucchesi C, Varga A, Coindre JM, Blakemore SJ, Clawson A, Suttle B, McDonald AA, Woodruff M, Ribich S, Hedrick E, Keilhack H, Thomson B, Owa T, Copeland RA, Ho PTC, Ribrag V (2018) Tazemetostat, an EZH2 inhibitor, in relapsed or refractory B-cell non-Hodgkin lymphoma and advanced solid tumours: a first-in-human, open-label, phase 1 study. *Lancet Oncol* 2018 May;19(5):649-659 ([PMID: 29650362](#))
77. Ivetac I, Gurung R, Hakim S, Horan KA, Sheffield DA, Binge LC, Majerus PW, Tiganis T, Mitchell CA (2009) Regulation of PI(3)K/Akt signalling and cellular transformation by inositol polyphosphate 4-phosphatase-1. *EMBO Rep*. 2009 May;10(5):487-93. Epub 2009 Mar 27 ([PMID: 19325558](#))
78. Ivetac I, Munday AD, Kisseleva MV, Zhang XM, Luff S, Tiganis T, Whisstock JC, Rowe T, Majerus PW, Mitchell CA (2005) The type Ialpha inositol polyphosphate 4-phosphatase generates and terminates phosphoinositide 3-kinase signals on endosomes and the plasma membrane. *Mol Biol Cell*. 2005 May;16(5):2218-33. Epub 2005 Feb 16 ([PMID: 15716355](#))
79. Jenkins MA, Dowty JG, Ait Ouakrim D, Mathews JD, Hopper JL, Drouet Y, Lasset C, Bonadona V, Win AK (2014) Short-term risk of colorectal cancer in individuals with lynch syndrome: a meta-analysis. *J Clin Oncol* 2015 Feb 1;33(4):326-31 ([PMID: 25534380](#))
80. Johnson DB, Frampton GM, Rioth MJ, Yusko E, Xu Y, Guo X, Ennis RC, Fabrizio D, Chalmers ZR, Greenbowe J, Ali SM, Balasubramanian S, Sun JX, He Y, Frederick DT, Puzanov I, Balko JM, Cates JM, Ross JS, Sanders C, Robins H, Shyr Y, Miller VA, Stephens PJ, Sullivan RJ, Sosman JA, Lovly CM (2016) Targeted Next Generation Sequencing Identifies Markers of Response to PD-1 Blockade. *Cancer Immunol Res*. 2016 Nov;4(11):959-967. Epub 2016 Sep 26 ([PMID: 27671167](#))
81. Jones S, Li M, Parsons DW, Zhang X, Wesseling J, Kristel P, Schmidt MK, Markowitz S, Yan H, Bigner D, Hruban RH, Eshleman JR, Iacobuzio-Donahue CA, Goggins M, Maitra A, Malek SN, Powell S, Vogelstein B, Kinzler KW, Velculescu VE, Papadopoulos N (2011) Somatic mutations in the chromatin remodeling gene ARID1A occur in several tumor types. *Hum Mutat*. 2012 Jan;33(1):100-3. Epub 2011 Nov 23 ([PMID: 22009941](#))
82. Jones S, Wang TL, Shih IeM, Mao TL, Nakayama K, Roden R, Glas R, Slamon D, Diaz LA, Vogelstein B, Kinzler KW, Velculescu VE, Papadopoulos N (2010) Frequent mutations of chromatin remodeling gene ARID1A in ovarian clear cell carcinoma. *Science*. 2010 Oct 08;330(6001):228-31. Epub 2010 Sep 8 ([PMID: 20826764](#))
83. Kanaar R, Troelstra C, Swagemakers SM, Essers J, Smit B, Franssen JH, Pastink A, Bezzubova OY, Buerstedde JM, Clever B, Heyer WD, Hoeijmakers JH (1996) Human and mouse homologs of the *Saccharomyces cerevisiae* RAD54 DNA repair gene: evidence for functional conservation. *Curr Biol*. 1996 Jul 01;6(7):828-38 ([PMID: 8805304](#))
84. Kastrinos F, Ojha RP, Leenen C, Alvero C, Mercado RC, Balmaña J, Valenzuela I, Balaguer F, Green R, Lindor NM, Thibodeau SN, Newcomb P, Win AK, Jenkins M, Buchanan DD, Bertario L, Sala P, Hampel H, Syngal S, Steyerberg EW (2015) Comparison of Prediction Models for Lynch Syndrome Among Individuals With Colorectal Cancer. *J Natl Cancer Inst* 2016 Feb;108(2) ([PMID: 26582061](#))
85. Katso RM, Pardo OE, Palamidessi A, Franz CM, Marinov M, De Laurentiis A, Downward J, Scita G, Ridley AJ, Waterfield MD, Arcaro A (2006) Phosphoinositide 3-Kinase C2beta regulates cytoskeletal organization and cell migration via Rac-dependent mechanisms. *Mol Biol Cell*. 2006 Sep;17(9):3729-44. Epub 2006 Jun 14 ([PMID: 16775008](#))
86. Kim H, Jen J, Vogelstein B, Hamilton SR (1994) Clinical and pathological characteristics of sporadic colorectal carcinomas with DNA replication errors in microsatellite sequences. *Am J Pathol* 1994 Jul;145(1):148-56 ([PMID: 8030745](#))
87. Kim J, Lee Y, Cho HJ, Lee YE, An J, Cho GH, Ko YH, Joo KM, Nam DH (2014) NTRK1 fusion in glioblastoma multiforme. *PLoS One*. 2014;9(3):e91940. Epub 2014 Mar 19 ([PMID: 24647444](#))
88. Kim JH, Kim SY, Baek JY, Cha YJ, Ahn JB, Kim HS, Lee KW, Kim JW, Kim TY, Chang WJ, Park JO, Kim J, Kim JE, Hong YS, Kim YH, Kim TW (2020) A Phase II Study of Avelumab Monotherapy in Patients with Mismatch Repair-Deficient/Microsatellite Instability-High or POLE-Mutated Metastatic or Unresectable Colorectal Cancer. *Cancer Res Treat*. 2020 Oct;52(4):1135-1144. Epub 2020 Apr 24 ([PMID: 32340084](#))
89. Kim MS, Kim YR, Yoo NJ, Lee SH (2012) Mutational analysis of DNMT3A gene in acute leukemias and common solid cancers. *APMIS*. 2013 Feb;121(2):85-94. Epub 2012 Jul 3 ([PMID: 23031157](#))
90. Klein R, Jing SQ, Nanduri V, O'Rourke E, Barbacid M (1991) The *trk* proto-oncogene encodes a receptor for nerve growth factor. *Cell*. 1991 Apr 05;65(1):189-97 ([PMID: 1849459](#))
91. Knutson SK, Kawano S, Minoshima Y, Warholc NM, Huang KC, Xiao Y, Kadowaki T, Uesugi M, Kuznetsov G, Kumar N, Wigle TJ, Klaus CR, Allain CJ, Raimondi A, Waters NJ, Smith JJ, Porter-Scott M, Chesworth R, Moyer MP, Copeland RA, Richon VM, Uenaka T, Pollock RM, Kuntz KW, Yokoi A, Keilhack H (2014) Selective inhibition of EZH2 by EPZ-6438 leads to potent antitumor activity in EZH2-mutant non-Hodgkin lymphoma. *Mol Cancer Ther*. 2014 Apr;13(4):842-54. Epub 2014 Feb 21 ([PMID: 24563539](#))

92. Korehisa S, Oki E, Iimori M, Nakaji Y, Shimokawa M, Saeki H, Okano S, Oda Y, Maehara Y (2017) Clinical significance of programmed cell death-ligand 1 expression and the immune microenvironment at the invasive front of colorectal cancers with high microsatellite instability. *Int J Cancer* 2018 Feb 15;142(4):822-832 ([PMID: 29044503](#))
93. Kristeleit R, Shapiro GI, Burris HA, Oza AM, LoRusso P, Patel MR, Domchek SM, Balmaña J, Drew Y, Chen LM, Safra T, Montes A, Giordano H, Maloney L, Goble S, Isaacson J, Xiao J, Borrow J, Rolfe L, Shapira-Frommer R (2017) A Phase I-II Study of the Oral PARP Inhibitor Rucaparib in Patients with Germline *BRCA1/2*-Mutated Ovarian Carcinoma or Other Solid Tumors. *Clin Cancer Res* 2017 Aug 1;23(15):4095-4106 ([PMID: 28264872](#))
94. Kung PP, Bingham P, Brooun A, Collins M, Deng YL, Dinh D, Fan C, Gajiwala KS, Grantner R, Gukasyan HJ, Hu W, Huang B, Kania R, Kephart SE, Krivacic C, Kumpf RA, Khamphavong P, Kraus M, Liu W, Maegley KA, Nguyen L, Ren S, Richter D, Rollins RA, Sach N, Sharma S, Sherrill J, Spangler J, Stewart AE, Sutton S, Uryu S, Verhelle D, Wang H, Wang S, Wythes M, Xin S, Yamazaki S, Zhu H, Zhu J, Zehnder L, Edwards M (2017) Optimization of Orally Bioavailable Enhancer of Zeste Homolog 2 (EZH2) Inhibitors Using Ligand and Property-Based Design Strategies: Identification of Development Candidate (R)-5,8-Dichloro-7-(methoxy(oxetan-3-yl)methyl)-2-((4-methoxy-6-methyl-2-oxo-1,2-dihydropyridin-3-yl)methyl)-3,4-dihydroisoquinolin-1(2H)-one (PF-06821497). *J Med Chem*. 2018 Feb 08;61(3):650-665. Epub 2017 Dec 27 ([PMID: 29211475](#))
95. Laury AR, Perets R, Piao H, Krane JF, Barletta JA, French C, Chirieac LR, Lis R, Loda M, Hornick JL, Drapkin R, Hirsch MS (2011) A comprehensive analysis of PAX8 expression in human epithelial tumors. *Am J Surg Pathol*. 2011 Jun;35(6):816-26 ([PMID: 21552115](#))
96. Le DT, Durham JN, Smith KN, Wang H, Bartlett BR, Aulakh LK, Lu S, Kemberling H, Wilt C, Lubner BS, Wong F, Azad NS, Rucki AA, Laheru D, Donehower R, Zaheer A, Fisher GA, Crocenzi TS, Lee JJ, Greten TF, Duffy AG, Ciombor KK, Eyring AD, Lam BH, Joe A, Kang SP, Holdhoff M, Danilova L, Cope L, Meyer C, Zhou S, Goldberg RM, Armstrong DK, Bever KM, Fader AN, Taube J, Housseau F, Spetzler D, Xiao N, Pardoll DM, Papadopoulos N, Kinzler KW, Eshleman JR, Vogelstein B, Anders RA, Diaz LA (2017) Mismatch repair deficiency predicts response of solid tumors to PD-1 blockade. *Science*. 2017 Jul 28;357(6349):409-413. Epub 2017 Jun 8 ([PMID: 28596308](#))
97. Le DT, Uram JN, Wang H, Bartlett BR, Kemberling H, Eyring AD, Skora AD, Lubner BS, Azad NS, Laheru D, Biedrzycki B, Donehower RC, Zaheer A, Fisher GA, Crocenzi TS, Lee JJ, Duffy SM, Goldberg RM, de la Chapelle A, Koshiji M, Bhajee F, Huebner T, Hruban RH, Wood LD, Cuka N, Pardoll DM, Papadopoulos N, Kinzler KW, Zhou S, Cornish TC, Taube JM, Anders RA, Eshleman JR, Vogelstein B, Diaz LA (2015) PD-1 Blockade in Tumors with Mismatch-Repair Deficiency. *N Engl J Med*. 2015 Jun 25;372(26):2509-20. Epub 2015 May 30 ([PMID: 26028255](#))
98. Lee DW, Han SW, Bae JM, Jang H, Han H, Kim H, Bang D, Jeong SY, Park KJ, Kang GH, Kim TY (2019) Tumor Mutation Burden and Prognosis in Patients with Colorectal Cancer Treated with Adjuvant Fluoropyrimidine and Oxaliplatin. *Clin Cancer Res* 2019 Oct 15;25(20):6141-6147 ([PMID: 31285374](#))
99. Lee KS, Kwak Y, Ahn S, Shin E, Oh HK, Kim DW, Kang SB, Choe G, Kim WH, Lee HS (2017) Prognostic implication of CD274 (PD-L1) protein expression in tumor-infiltrating immune cells for microsatellite unstable and stable colorectal cancer. *Cancer Immunol Immunother*. 2017 Jul;66(7):927-939. Epub 2017 Apr 12 ([PMID: 28405764](#))
100. Lee LH, Sadot E, Ivelja S, Vakiani E, Hechtman JF, Sevinsky CJ, Klimstra DS, Ginty F, Shia J (2016) ARID1A expression in early stage colorectal adenocarcinoma: an exploration of its prognostic significance. *Hum Pathol* 2016 Jul;53:97-104 ([PMID: 26980037](#))
101. Lee SY, Haq F, Kim D, Jun C, Jo HJ, Ahn SM, Lee WS (2014) Comparative genomic analysis of primary and synchronous metastatic colorectal cancers. *PLoS One*. 2014;9(3):e90459. Epub 2014 Mar 5 ([PMID: 24599305](#))
102. Lee V, Le DT (2015) Efficacy of PD-1 blockade in tumors with MMR deficiency. *Immunotherapy* 2016;8(1):1-3 ([PMID: 26643016](#))
103. Leichman L, Groshen S, O'Neil BH, Messersmith W, Berlin J, Chan E, Leichman CG, Cohen SJ, Cohen D, Lenz HJ, Gold P, Boman B, Fielding A, Locker G, Cason RC, Hamilton SR, Hochster HS (2016) Phase II Study of Olaparib (AZD-2281) After Standard Systemic Therapies for Disseminated Colorectal Cancer. *Oncologist*. 2016 Feb;21(2):172-7. Epub 2016 Jan 19 ([PMID: 26786262](#))
104. Li L, Karanika S, Yang G, Wang J, Park S, Broom BM, Manyam GC, Wu W, Luo Y, Basourakos S, Song JH, Gallick GE, Karantanos T, Korentzelos D, Azad AK, Kim J, Corn PG, Aparicio AM, Logothetis CJ, Troncoso P, Heffernan T, Toniatti C, Lee HS, Lee JS, Zuo X, Chang W, Yin J, Thompson TC (2017) Androgen receptor inhibitor-induced "BRCAness" and PARP inhibition are synthetically lethal for castration-resistant prostate cancer. *Sci Signal*. 2017 May 23;10(480) ([PMID: 28536297](#))
105. Lin EI, Tseng LH, Gocke CD, Reil S, Le DT, Azad NS, Eshleman JR (2015) Mutational profiling of colorectal cancers with microsatellite instability. *Oncotarget*. 2015 Dec 08;6(39):42334-44 ([PMID: 26517354](#))
106. Liontos M, Anastasiou I, Bamias A, Dimopoulos MA (2016) DNA damage, tumor mutational load and their impact on immune responses against cancer. *Ann Transl Med* 2016 Jul;4(14):264 ([PMID: 27563651](#))
107. Lipson EJ, Sharfman WH, Drake CG, Wollner I, Taube JM, Anders RA, Xu H, Yao S, Pons A, Chen L, Pardoll DM, Brahmer JR, Topalian SL (2012) Durable cancer regression off-treatment and effective reinduction therapy with an anti-PD-1 antibody. *Clin Cancer Res* 2013 Jan 15;19(2):462-8 ([PMID: 23169436](#))
108. Liu P, Morrison C, Wang L, Xiong D, Vedell P, Cui P, Hua X, Ding F, Lu Y, James M, Ebben JD, Xu H, Adjei AA, Head K, Andrae JW, Tschannen MR, Jacob H, Pan J, Zhang Q, Van den Bergh F, Xiao H, Lo KC, Patel J, Richmond T, Watt MA, Albert T, Selzer R, Anderson M, Wang J, Wang Y, Starnes S, Yang P, You M (2012) Identification of somatic mutations in non-small cell lung carcinomas using whole-exome sequencing. *Carcinogenesis*. 2012 Jul;33(7):1270-6. Epub 2012 Apr 17 ([PMID: 22510280](#))
109. Llosa NJ, Cruise M, Tam A, Wicks EC, Hechenbleikner EM, Taube JM, Blosser RL, Fan H, Wang H, Lubner BS, Zhang M, Papadopoulos N, Kinzler KW, Vogelstein B, Sears CL, Anders RA, Pardoll DM, Housseau F (2014) The vigorous immune microenvironment of microsatellite instable colon cancer is balanced by multiple counter-inhibitory checkpoints. *Cancer Discov* 2015 Jan;5(1):43-51 ([PMID: 25358689](#))
110. Loree JM, Pereira AAL, Lam M, Willauer AN, Raghav K, Dasari A, Morris VK, Advani S, Menter DG, Eng C, Shaw K, Broaddus R, Routbort MJ, Liu Y, Morris JS, Luthra R, Meric-Bernstam F, Overman MJ, Maru D, Kopetz S (2017) Classifying Colorectal Cancer by Tumor Location Rather than Sidedness Highlights a Continuum in Mutation Profiles and Consensus Molecular Subtypes. *Clin Cancer Res* 2018 Mar 1;24(5):1062-1072 ([PMID: 29180604](#))
111. Lynch HT, Lynch PM, Lanspa SJ, Snyder CL, Lynch JF, Boland CR (2009) Review of the Lynch syndrome: history, molecular genetics, screening, differential diagnosis, and medicolegal ramifications. *Clin Genet*. 2009 Jul;76(1):1-18 ([PMID: 19659756](#))
112. Maffucci T, Cooke FT, Foster FM, Traer CJ, Fry MJ, Falasca M (2005) Class II phosphoinositide 3-kinase defines a novel signaling pathway in cell migration. *J Cell Biol*. 2005 Jun 06;169(5):789-99. Epub 2005 May 31 ([PMID: 15928202](#))

113. Malhotra A, Wang Y, Waters J, Chen K, Meric-Bernstam F, Hall IM, Navin NE (2015) Ploidy-Seq: inferring mutational chronology by sequencing polyploid tumor subpopulations. *Genome Med.* 2015;7(1):6. Epub 2015 Jan 28 ([PMID: 25729435](#))
114. Mansouri A, Chowdhury K, Gruss P (1998) Follicular cells of the thyroid gland require Pax8 gene function. *Nat Genet.* 1998 May;19(1):87-90 ([PMID: 9590297](#))
115. Marabelle A, Fakih M, Lopez J, Shah M, Shapira-Frommer R, Nakagawa K, Chung HC, Kindler HL, Lopez-Martin JA, Miller WH Jr, Italiano A, Kao S, Piha-Paul SA, Delord JP, McWilliams RR, Fabrizio DA, Aurora-Garg D, Xu L, Jin F, Norwood K, Bang YJ (2020) Association of tumour mutational burden with outcomes in patients with advanced solid tumours treated with pembrolizumab: prospective biomarker analysis of the multicohort, open-label, phase 2 KEYNOTE-158 study. *Lancet Oncol* 2020 Oct;21(10):1353-1365 ([PMID: 32919526](#))
116. Marcus L, Lemery SJ, Keegan P, Pazdur R (2019) FDA Approval Summary: Pembrolizumab for the Treatment of Microsatellite Instability-High Solid Tumors. *Clin Cancer Res* 2019 Jul 1;25(13):3753-3758 ([PMID: 30787022](#))
117. Martin SA, Lord CJ, Ashworth A (2010) Therapeutic targeting of the DNA mismatch repair pathway. *Clin Cancer Res.* 2010 Nov 01;16(21):5107-13. Epub 2010 Sep 7 ([PMID: 20823149](#))
118. Martin-Zanca D, Hughes SH, Barbacid M (1986) A human oncogene formed by the fusion of truncated tropomyosin and protein tyrosine kinase sequences. *Nature.* 1986 Feb 27-Mar 5;319(6056):743-8 ([PMID: 2869410](#))
119. Mathur R, Alver BH, San Roman AK, Wilson BG, Wang X, Agoston AT, Park PJ, Shivdasani RA, Roberts CW (2016) ARID1A loss impairs enhancer-mediated gene regulation and drives colon cancer in mice. *Nat Genet* 2017 Feb;49(2):296-302 ([PMID: 27941798](#))
120. Matsuda M, Miyagawa K, Takahashi M, Fukuda T, Kataoka T, Asahara T, Inui H, Watatani M, Yasutomi M, Kamada N, Dohi K, Kamiya K (1999) Mutations in the RAD54 recombination gene in primary cancers. *Oncogene.* 1999 Jun 03;18(22):3427-30 ([PMID: 10362365](#))
121. Mayle A, Yang L, Rodriguez B, Zhou T, Chang E, Curry CV, Challen GA, Li W, Wheeler D, Rebel VI, Goodell MA (2015) Dnmt3a loss predisposes murine hematopoietic stem cells to malignant transformation. *Blood.* 2015 Jan 22;125(4):629-38 ([PMID: 25416277](#))
122. Migden MR, Rischin D, Schmults CD, Guminski A, Hauschild A, Lewis KD, Chung CH, Hernandez-Aya L, Lim AM, Chang ALS, Rabinowits G, Thai AA, Dunn LA, Hughes BGM, Khushalani NI, Modi B, Schadendorf D, Gao B, Seebach F, Li S, Li J, Mathias M, Booth J, Mohan K, Stankevich E, Babiker HM, Brana I, Gil-Martin M, Homsy J, Johnson ML, Moreno V, Niu J, Owonikoko TK, Papadopoulos KP, Yancopoulos GD, Lowy I, Fury MG (2018) PD-1 Blockade with Cemiplimab in Advanced Cutaneous Squamous-Cell Carcinoma. *N Engl J Med* 2018 Jul 26;379(4):341-351 ([PMID: 29863979](#))
123. Millen R, Hendry S, Narasimhan V, Abbott R, Croxford M, Gibbs P, Tie J, Wong HL, Jones I, Kosmider S, Byrne D, Zalcberg J, Fox S, Desai J, Visvanathan K, Ramsay RG, Tran B (2020) CD8⁺ tumor-infiltrating lymphocytes within the primary tumor of patients with synchronous *de novo* metastatic colorectal carcinoma do not track with survival. *Clin Transl Immunology* 2020;9(7):e1155 ([PMID: 32953115](#))
124. Morioka S, Nigorikawa K, Sasaki J, Hazeki K, Kasuu Y, Sasaki T, Hazeki O (2016) Myeloid cell-specific inositol polyphosphate-4-phosphatase type I knockout mice impair bacteria clearance in a murine peritonitis model. *Innate Immun.* 2016 Aug;22(6):444-51. Epub 2016 Jun 1 ([PMID: 27252170](#))
125. Moslein G, Tester DJ, Lindor NM, Honchel R, Cunningham JM, French AJ, Halling KC, Schwab M, Goretzki P, Thibodeau SN (1996) Microsatellite instability and mutation analysis of hMSH2 and hMLH1 in patients with sporadic, familial and hereditary colorectal cancer. *Hum Mol Genet.* 1996 Sep; 5(9):1245-52 ([PMID: 8872463](#))
126. Noepel-Duennebacke S, Juette H, Feder IS, Kluxen L, Basara N, Hiller W, Herzog T, Klaassen-Mielke R, Mueller L, Senkal M, Engel L, Teschendorf C, Trenn G, Verdoodt B, Wolters H, Uhl W, Reinacher-Schick A, Tannappel A (2020) High microsatellite instability (MSI-H) is associated with distinct clinical and molecular characteristics and an improved survival in early Colon cancer (CC); real world data from the AIO molecular registry Colopredict Plus. *Z Gastroenterol* 2020 Jun;58(6):533-541 ([PMID: 32544965](#))
127. Nystuen A, Legare ME, Shultz LD, Frankel WN (2001) A null mutation in inositol polyphosphate 4-phosphatase type I causes selective neuronal loss in weeble mutant mice. *Neuron.* 2001 Oct 25;32(2):203-12 ([PMID: 11683991](#))
128. O'Brien EC, Brewin J, Chevassut T (2014) DNMT3A: the DioNysian MonsTer of acute myeloid leukaemia. *Ther Adv Hematol.* 2014 Dec;5(6):187-96 ([PMID: 25469209](#))
129. Overman MJ, McDermott R, Leach JL, Lonardi S, Lenz HJ, Morse MA, Desai J, Hill A, Axelson M, Moss RA, Goldberg MV, Cao ZA, Ledezne JM, Maglinte GA, Kopetz S, André T (2017) Nivolumab in patients with metastatic DNA mismatch repair-deficient or microsatellite instability-high colorectal cancer (CheckMate 142): an open-label, multicentre, phase 2 study. *Lancet Oncol.* 2017 Sep;18(9):1182-1191. Epub 2017 Jul 19 ([PMID: 28734759](#))
130. Pardoll DM (2012) The blockade of immune checkpoints in cancer immunotherapy. *Nat Rev Cancer.* 2012 Mar 22;12(4):252-64 ([PMID: 22437870](#))
131. Patel MR, Falchook GS, Hamada K, Makris L, Bendell JC (2021) A phase 2 trial of trifluridine/tipiracil plus nivolumab in patients with heavily pretreated microsatellite-stable metastatic colorectal cancer. *Cancer Med* 2021 Feb;10(4):1183-1190 ([PMID: 33544407](#))
132. Pearlman R, Frankel WL, Swanson B, Zhao W, Yilmaz A, Miller K, Bacher J, Bigley C, Nelsen L, Goodfellow PJ, Goldberg RM, Paskett E, Shields PG, Freudenheim JL, Stanich PP, Lattimer I, Arnold M, Liyanarachchi S, Kalady M, Heald B, Greenwood C, Paquette I, Prues M, Draper DJ, Lindeman C, Kuebler JP, Reynolds K, Brell JM, Shaper AA, Mahesh S, Buie N, Weeman K, Shine K, Haut M, Edwards J, Bastola S, Wickham K, Khanduja KS, Zacks R, Pritchard CC, Shirts BH, Jacobson A, Allen B, de la Chapelle A, Hampel H, Ohio Colorectal Cancer Prevention Initiative Study Group (2017) Prevalence and Spectrum of Germline Cancer Susceptibility Gene Mutations Among Patients With Early-Onset Colorectal Cancer. *JAMA Oncol.* 2017 Apr 01;3(4):464-471 ([PMID: 27978560](#))
133. Pishvaian MJ, Slack RS, Jiang W, He AR, Hwang JJ, Hankin A, Dorsch-Vogel K, Kukadiya D, Weiner LM, Marshall JL, Brody JR (2018) A phase 2 study of the PARP inhibitor veliparib plus temozolomide in patients with heavily pretreated metastatic colorectal cancer. *Cancer* 2018 Jun 1;124(11): 2337-2346 ([PMID: 29579325](#))
134. Plachov D, Chowdhury K, Walther C, Simon D, Guenet JL, Gruss P (1990) Pax8, a murine paired box gene expressed in the developing excretory system and thyroid gland. *Development.* 1990 Oct;110(2):643-51 ([PMID: 1723950](#))
135. Planchard D, Popat S, Kerr K, Novello S, Smit EF, Faivre-Finn C, Mok TS, Reck M, Van Schil PE, Hellmann MD, Peters S, ESMO Guidelines Committee (2018) Metastatic non-small cell lung cancer: ESMO Clinical Practice Guidelines for diagnosis, treatment and follow-up. *Ann Oncol.* 2018 Oct 01;29(Suppl 4):iv192-iv237 ([PMID: 30285222](#))

136. Popat S, Hubner R, Houlston RS (2005) Systematic review of microsatellite instability and colorectal cancer prognosis. *J Clin Oncol* 2005 Jan 20;23(3):609-18 ([PMID: 15659508](#))
137. Poulogiannis G, Frayling IM, Arends MJ (2010) DNA mismatch repair deficiency in sporadic colorectal cancer and Lynch syndrome. *Histopathology*. 2010 Jan;56(2):167-79 ([PMID: 20102395](#))
138. Rasio D, Murakumo Y, Robbins D, Roth T, Silver A, Negrini M, Schmidt C, Burczak J, Fishel R, Croce CM (1997) Characterization of the human homologue of RAD54: a gene located on chromosome 1p32 at a region of high loss of heterozygosity in breast tumors. *Cancer Res*. 1997 Jun 15;57(12):2378-83 ([PMID: 9192813](#))
139. Ribic CM, Sargent DJ, Moore MJ, Thibodeau SN, French AJ, Goldberg RM, Hamilton SR, Laurent-Puig P, Gryfe R, Shepherd LE, Tu D, Redston M, Gallinger S (2003) Tumor microsatellite-instability status as a predictor of benefit from fluorouracil-based adjuvant chemotherapy for colon cancer. *N Engl J Med* 2003 Jul 17;349(3):247-57 ([PMID: 12867608](#))
140. Richard C, Fumet JD, Chevrier S, Derangère V, Ledys F, Lagrange A, Favier L, Coudert B, Arnould L, Truntzer C, Boidot R, Ghiringhelli F (2018) Exome Analysis Reveals Genomic Markers Associated with Better Efficacy of Nivolumab in Lung Cancer Patients. *Clin Cancer Res* 2019 Feb 1;25(3):957-966 ([PMID: 30154227](#))
141. Rizvi NA, Hellmann MD, Snyder A, Kvistborg P, Makarov V, Havel JJ, Lee W, Yuan J, Wong P, Ho TS, Miller ML, Rekhtman N, Moreira AL, Ibrahim F, Bruggeman C, Gasmi B, Zappasodi R, Maeda Y, Sander C, Garon EB, Merghoub T, Wolchok JD, Schumacher TN, Chan TA (2015) Cancer immunology. Mutational landscape determines sensitivity to PD-1 blockade in non-small cell lung cancer. *Science*. 2015 Apr 03;348(6230):124-8. Epub 2015 Mar 12 ([PMID: 25765070](#))
142. Rodgers LH, Ó hAinmhire E, Young AN, Burdette JE (2016) Loss of PAX8 in high-grade serous ovarian cancer reduces cell survival despite unique modes of action in the fallopian tube and ovarian surface epithelium. *Oncotarget*. 2016 May 31;7(22):32785-95 ([PMID: 27129161](#))
143. Rooney MS, Shukla SA, Wu CJ, Getz G, Hacohen N (2015) Molecular and genetic properties of tumors associated with local immune cytolytic activity. *Cell*. 2015 Jan 15;160(1-2):48-61 ([PMID: 25594174](#))
144. Rosenberg JE, Hoffman-Censits J, Powles T, van der Heijden MS, Balar AV, Necchi A, Dawson N, O'Donnell PH, Balmanoukian A, Loriot Y, Srinivas S, Retz MM, Grivas P, Joseph RW, Galsky MD, Fleming MT, Petrylak DP, Perez-Gracia JL, Burris HA, Castellano D, Canil C, Bellmunt J, Bajorin D, Nickles D, Bourgon R, Frampton GM, Cui N, Mariathasan S, Abidoye O, Fine GD, Dreicer R (2016) Atezolizumab in patients with locally advanced and metastatic urothelial carcinoma who have progressed following treatment with platinum-based chemotherapy: a single-arm, multicentre, phase 2 trial. *Lancet*. 2016 May 07;387(10031):1909-20. Epub 2016 Mar 4 ([PMID: 26952546](#))
145. Russo A, O'Bryan JP (2012) Intersectin 1 is required for neuroblastoma tumorigenesis. *Oncogene*. 2012 Nov 15;31(46):4828-34. Epub 2012 Jan 23 ([PMID: 22266851](#))
146. Salem ME, Battaglin F, Goldberg RM, Puccini A, Shields AF, Arguello D, Korn WM, Marshall JL, Grothey A, Lenz HJ (2019) Molecular Analyses of Left- and Right-Sided Tumors in Adolescents and Young Adults with Colorectal Cancer. *Oncologist*. 2020 May;25(5):404-413. Epub 2019 Dec 17 ([PMID: 31848314](#))
147. Salem ME, Puccini A, Grothey A, Raghavan D, Goldberg RM, Xiu J, Korn WM, Weinberg BA, Hwang JJ, Shields AF, Marshall JL, Philip PA, Lenz HJ (2018) Landscape of Tumor Mutation Load, Mismatch Repair Deficiency, and PD-L1 Expression in a Large Patient Cohort of Gastrointestinal Cancers. *Mol Cancer Res* 2018 May;16(5):805-812 ([PMID: 29523759](#))
148. Samstein RM, Lee CH, Shoushtari AN, Hellmann MD, Shen R, Janjigian YY, Barron DA, Zehir A, Jordan EJ, Omuro A, Kaley TJ, Kendall SM, Motzer RJ, Hakimi AA, Voss MH, Russo P, Rosenberg J, Iyer G, Bochner BH, Bajorin DF, Al-Ahmadie HA, Chaft JE, Rudin CM, Riely GJ, Baxi S, Ho AL, Wong RJ, Pfister DG, Wolchok JD, Barker CA, Gutin PH, Brennan CW, Tabar V, Mellinghoff IK, DeAngelis LM, Ariyan CE, Lee N, Tap WD, Gounder MM, D'Angelo SP, Saltz L, Stadler ZK, Scher HI, Baselga J, Razavi P, Klebanoff CA, Yaeger R, Segal NH, Ku GY, DeMatteo RP, Ladanyi M, Rizvi NA, Berger MF, Riaz N, Solit DB, Chan TA, Morris LGT (2019) Tumor mutational load predicts survival after immunotherapy across multiple cancer types. *Nat Genet* 2019 Feb;51(2):202-206 ([PMID: 30643254](#))
149. Sandhu SK, Schelman WR, Wilding G, Moreno V, Baird RD, Miranda S, Hylands L, Riisnaes R, Forster M, Omlin A, Kreischer N, Thway K, Gevensleben H, Sun L, Loughney J, Chatterjee M, Toniatti C, Carpenter CL, Iannone R, Kaye SB, de Bono JS, Wenham RM (2013) The poly(ADP-ribose) polymerase inhibitor niraparib (MK4827) in BRCA mutation carriers and patients with sporadic cancer: a phase 1 dose-escalation trial. *Lancet Oncol* 2013 Aug;14(9):882-92 ([PMID: 23810788](#))
150. Sargent DJ, Marsoni S, Monges G, Thibodeau SN, Labianca R, Hamilton SR, French AJ, Kabat B, Foster NR, Torri V, Ribic C, Grothey A, Moore M, Zaniboni A, Seitz JF, Sinicrope F, Gallinger S (2010) Defective mismatch repair as a predictive marker for lack of efficacy of fluorouracil-based adjuvant therapy in colon cancer. *J Clin Oncol* 2010 Jul 10;28(20):3219-26 ([PMID: 20498393](#))
151. Sasaki J, Kofuji S, Itoh R, Momiyama T, Takayama K, Murakami H, Chida S, Tsuya Y, Takasuga S, Eguchi S, Asanuma K, Horie Y, Miura K, Davies EM, Mitchell C, Yamazaki M, Hirai H, Takenawa T, Suzuki A, Sasaki T (2010) The PtdIns(3,4)P(2) phosphatase INPP4A is a suppressor of excitotoxic neuronal death. *Nature*. 2010 May 27;465(7297):497-501. Epub 2010 May 12 ([PMID: 20463662](#))
152. Schulten HJ, Bangash M, Karim S, Dallol A, Hussein D, Merdad A, Al-Thoubaity FK, Al-Maghrabi J, Jamal A, Al-Ghamdi F, Choudhry H, Baeesa SS, Chaudhary AG, Al-Qahtani MH (2017) Comprehensive molecular biomarker identification in breast cancer brain metastases. *J Transl Med*. 2017 Dec 29;15(1):269 ([PMID: 29287594](#))
153. Schumacher TN, Hacohen N (2016) Neoantigens encoded in the cancer genome. *Curr Opin Immunol* 2016 Aug;41:98-103 ([PMID: 27518850](#))
154. Sen M, Wang X, Hamdan FH, Rapp J, Eggert J, Kosinsky RL, Wegwitz F, Kutschat AP, Younesi FS, Gaedcke J, Grade M, Hessmann E, Papanonis A, Ströbel P, Johnsen SA (2019) ARID1A facilitates KRAS signaling-regulated enhancer activity in an AP1-dependent manner in colorectal cancer cells. *Clin Epigenetics*. 2019 Jun 19;11(1):92 ([PMID: 31217031](#))
155. Shah N, Lankarovich M, Lee H, Yoon JG, Schroeder B, Foltz G (2013) Exploration of the gene fusion landscape of glioblastoma using transcriptome sequencing and copy number data. *BMC Genomics*. 2013 Nov 22;14:818 ([PMID: 24261984](#))
156. Shamseddine A, Zeidan YH, El Husseini Z, Kreidieh M, Al Darazi M, Turfa R, Kattan J, Khalifeh I, Mukherji D, Temraz S, Alqasem K, Amarin R, Al Awabdeh T, Deeba S, Jamali F, Mohamad I, Elkhaldi M, Daoud F, Al Masri M, Dabous A, Hushki A, Jaber O, Charafeddine M, Geara F (2020) Efficacy and safety-in analysis of short-course radiation followed by mFOLFOX-6 plus avelumab for locally advanced rectal adenocarcinoma. *Radiat Oncol* 2020 Oct 7;15(1):233 ([PMID: 33028346](#))

157. Sharma R, Sanchez-Ferras O, Bouchard M (2015) Pax genes in renal development, disease and regeneration. *Semin Cell Dev Biol.* 2015 Aug;44:97-106. Epub 2015 Sep 26 ([PMID: 26410163](#))
158. Shen J, Ju Z, Zhao W, Wang L, Peng Y, Ge Z, Nagel ZD, Zou J, Wang C, Kapoor P, Ma X, Ma D, Liang J, Song S, Liu J, Samson LD, Ajani JA, Li GM, Liang H, Shen X, Mills GB, Peng G (2018) ARID1A deficiency promotes mutability and potentiates therapeutic antitumor immunity unleashed by immune checkpoint blockade. *Nat Med.* 2018 May;24(5):556-562. Epub 2018 May 7 ([PMID: 29736026](#))
159. Shen J, Peng Y, Wei L, Zhang W, Yang L, Lan L, Kapoor P, Ju Z, Mo Q, Shih IeM, Uray IP, Wu X, Brown PH, Shen X, Mills GB, Peng G (2015) ARID1A Deficiency Impairs the DNA Damage Checkpoint and Sensitizes Cells to PARP Inhibitors. *Cancer Discov.* 2015 Jul;5(7):752-67. Epub 2015 Jun 11 ([PMID: 26069190](#))
160. Sigurdsson S, Van Komen S, Petukhova G, Sung P (2002) Homologous DNA pairing by human recombination factors Rad51 and Rad54. *J Biol Chem.* 2002 Nov 08;277(45):42790-4. Epub 2002 Aug 29 ([PMID: 12205100](#))
161. Sindić A, Crljen V, Matković K, Lukinović-Skudar V, Visnjić D, Banfić H (2006) Activation of phosphoinositide 3-kinase C2 beta in the nuclear matrix during compensatory liver growth. *Adv Enzyme Regul.* 2006;46:280-7. Epub 2006 Jul 18 ([PMID: 16857245](#))
162. Smeby J, Kryeziu K, Berg KCG, Eilertsen IA, Eide PW, Johannessen B, Guren MG, Nesbakken A, Bruun J, Lothe RA, Sveen A (2020) Molecular correlates of sensitivity to PARP inhibition beyond homologous recombination deficiency in pre-clinical models of colorectal cancer point to wild-type TP53 activity. *EBioMedicine.* 2020 Sep;59:102923. Epub 2020 Aug 13 ([PMID: 32799124](#))
163. Smirnova M, Van Komen S, Sung P, Klein HL (2004) Effects of tumor-associated mutations on Rad54 functions. *J Biol Chem.* 2004 Jun 04;279(23):24081-8. Epub 2004 Mar 31 ([PMID: 15056673](#))
164. Song Y, Wang L, Ran W, Li G, Xiao Y, Wang X, Zhang L, Xing X (2020) Effect of Tumor Location on Clinicopathological and Molecular Markers in Colorectal Cancer in Eastern China Patients: An Analysis of 2,356 Cases. *Front Genet.* 2020;11:96. Epub 2020 Feb 25 ([PMID: 32161617](#))
165. Stadler ZK, Battaglioli F, Middha S, Hechtman JF, Tran C, Cercek A, Yaeger R, Segal NH, Varghese AM, Reidy-Lagunes DL, Kemeny NE, Salo-Mullen EE, Ashraf A, Weiser MR, Garcia-Aguilar J, Robson ME, Offit K, Arcila ME, Berger MF, Shia J, Solit DB, Saltz LB (2016) Reliable Detection of Mismatch Repair Deficiency in Colorectal Cancers Using Mutational Load in Next-Generation Sequencing Panels. *J Clin Oncol.* 2016 Jun 20;34(18):2141-7. Epub 2016 Mar 28 ([PMID: 27022117](#))
166. Stephens RM, Loeb DM, Copeland TD, Pawson T, Greene LA, Kaplan DR (1994) Trk receptors use redundant signal transduction pathways involving SHC and PLC-gamma 1 to mediate NGF responses. *Neuron.* 1994 Mar;12(3):691-705 ([PMID: 8155326](#))
167. Tacconelli A, Farina AR, Cappabianca L, Desantis G, Tessitore A, Vetuschi A, Sferra R, Rucci N, Argenti B, Screpanti I, Gulino A, Mackay AR (2004) TrkA alternative splicing: a regulated tumor-promoting switch in human neuroblastoma. *Cancer Cell.* 2004 Oct;6(4):347-60 ([PMID: 15488758](#))
168. Tacha D, Zhou D, Cheng L (2011) Expression of PAX8 in normal and neoplastic tissues: a comprehensive immunohistochemical study. *Appl Immunohistochem Mol Morphol.* 2011 Jul;19(4):293-9 ([PMID: 21285870](#))
169. Taieb J, Kourie HR, Emile JF, Le Malicot K, Balogoun R, Taberero J, Mini E, Folprecht G, Van Laethem JL, Mulot C, Bouché O, Aparicio T, Michel P, Thaler J, Bridgewater J, Van Cutsem E, Perkins G, Lepage C, Salazar R, Laurent-Puig P, Pan-European Trials in Alimentary Tract Cancer (PETACC)-8 Investigators (2018) Association of Prognostic Value of Primary Tumor Location in Stage III Colon Cancer With RAS and BRAF Mutational Status. *JAMA Oncol.* 2018 Jul 12;4(7):e173695 ([PMID: 29167892](#))
170. Taieb J, Le Malicot K, Shi Q, Penault-Llorca F, Bouché O, Taberero J, Mini E, Goldberg RM, Folprecht G, Luc Van Laethem J, Sargent DJ, Alberts SR, Emile JF, Laurent Puig P, Sinicrope FA (2016) Prognostic Value of BRAF and KRAS Mutations in MSI and MSS Stage III Colon Cancer. *J Natl Cancer Inst.* 2017 May;109(5). Epub 2016 Dec 31 ([PMID: 28040692](#))
171. Tan WJ, Hamzah JL, Acharyya S, Foo FJ, Lim KH, Tan IBH, Tang CL, Chew MH (2017) Evaluation of Long-Term Outcomes of Microsatellite Instability Status in an Asian Cohort of Sporadic Colorectal Cancers. *J Gastrointest Cancer* 2018 Sep;49(3):311-318 ([PMID: 28550452](#))
172. Taube JM, Klein A, Brahmer JR, Xu H, Pan X, Kim JH, Chen L, Pardoll DM, Topalian SL, Anders RA (2014) Association of PD-1, PD-1 ligands, and other features of the tumor immune microenvironment with response to anti-PD-1 therapy. *Clin Cancer Res* 2014 Oct 1;20(19):5064-74 ([PMID: 24714771](#))
173. Thibodeau SN, Bren G, Schaid D (1993) Microsatellite instability in cancer of the proximal colon. *Science* 1993 May 7;260(5109):816-9 ([PMID: 8484122](#))
174. Timmermann B, Kerick M, Roehr C, Fischer A, Isau M, Boerno ST, Wunderlich A, Barmeyer C, Seemann P, Koenig J, Lappe M, Kuss AW, Garshasbi M, Bertram L, Trappe K, Werber M, Herrmann BG, Zatlouk K, Lehrach H, Schweiger MR (2010) Somatic mutation profiles of MSI and MSS colorectal cancer identified by whole exome next generation sequencing and bioinformatics analysis. *PLoS One.* 2010 Dec 22;5(12):e15661 ([PMID: 21203531](#))
175. Tong Y, Merino D, Nimmervoll B, Gupta K, Wang YD, Finkelstein D, Dalton J, Ellison DW, Ma X, Zhang J, Malkin D, Gilbertson RJ (2015) Cross-Species Genomics Identifies TAF12, NFYC, and RAD54L as Choroid Plexus Carcinoma Oncogenes. *Cancer Cell.* 2015 May 11;27(5):712-27 ([PMID: 25965574](#))
176. Topalian SL, Drake CG, Pardoll DM (2012) Targeting the PD-1/B7-H1(PD-L1) pathway to activate anti-tumor immunity. *Curr Opin Immunol.* 2012 Apr;24(2):207-12. Epub 2012 Jan 9 ([PMID: 22236695](#))
177. Trowbridge JJ, Orkin SH (2011) Dnmt3a silences hematopoietic stem cell self-renewal. *Nat Genet.* 2011 Dec 27;44(1):13-4 ([PMID: 22200773](#))
178. Uhlig J, Cecchini M, Sheth A, Stein S, Lacy J, Kim HS (2021) Microsatellite Instability and KRAS Mutation in Stage IV Colorectal Cancer: Prevalence, Geographic Discrepancies, and Outcomes From the National Cancer Database. *J Natl Compr Canc Netw* 2021 Feb 2;:1-12 ([PMID: 33530058](#))
179. Umar A, Boland CR, Terdiman JP, Syngal S, de la Chapelle A, Rüschoff J, Fishel R, Lindor NM, Burgart LJ, Hamelin R, Hamilton SR, Hiatt RA, Jass J, Lindblom A, Lynch HT, Peltomaki P, Ramsey SD, Rodriguez-Bigas MA, Vasen HF, Hawk ET, Barrett JC, Freedman AN, Srivastava S (2004) Revised Bethesda Guidelines for hereditary nonpolyposis colorectal cancer (Lynch syndrome) and microsatellite instability. *J Natl Cancer Inst* 2004 Feb 18;96(4):261-8 ([PMID: 14970275](#))
180. Vaishnavi A, Capelletti M, Le AT, Kako S, Butaney M, Ercan D, Mahale S, Davies KD, Aisner DL, Pilling AB, Berge EM, Kim J, Sasaki H, Park S, Kryukov G, Garraway LA, Hammerman PS, Haas J, Andrews SW, Lipson D, Stephens PJ, Miller VA, Varella-Garcia M, Jänne PA, Doebele RC (2013) Oncogenic and drug-sensitive NTRK1 rearrangements in lung cancer. *Nat Med.* 2013 Nov;19(11):1469-1472. Epub 2013 Oct 27 ([PMID: 24162815](#))

181. Valentini AM, Di Pinto F, Cariola F, Guerra V, Giannelli G, Caruso ML, Pirrelli M (2018) PD-L1 expression in colorectal cancer defines three subsets of tumor immune microenvironments. *Oncotarget* 2018 Feb 2;9(9):8584-8596 ([PMID: 29492219](#))
182. Vallböhmer D, Brabender J, Yang D, Schneider PM, Metzger R, Danenberg KD, Hölscher AH, Danenberg PV (2006) DNA methyltransferases messenger RNA expression and aberrant methylation of CpG islands in non-small-cell lung cancer: association and prognostic value. *Clin Lung Cancer*. 2006 Jul;8(1):39-44 ([PMID: 16870044](#))
183. Vilar E, Gruber SB (2010) Microsatellite instability in colorectal cancer-the stable evidence. *Nat Rev Clin Oncol*. 2010 Mar;7(3):153-62. Epub 2010 Feb 9 ([PMID: 20142816](#))
184. Vogt N, Gibaud A, Almeida A, Ourliac-Garnier I, Debatisse M, Malfoy B (2010) Relationships linking amplification level to gene over-expression in gliomas. *PLoS One*. 2010 Dec 08;5(12):e14249. Epub 2010 Dec 8 ([PMID: 21170331](#))
185. Wang C, Feng Z, Jiang K, Zuo X (2016) [ARTICLE WITHDRAWN] Upregulation of MicroRNA-935 Promotes the Malignant Behaviors of Pancreatic Carcinoma PANC-1 Cells via Targeting Inositol Polyphosphate 4-Phosphatase Type I Gene (INPP4A). *Oncol Res*. 2017 Apr 14;25(4):559-569. Epub 2016 Oct 11 ([PMID: 27733216](#))
186. Wang R, Wu Y, Huang W, Chen W (2017) MicroRNA-940 Targets INPP4A or GSK3 β and Activates the Wnt/ β -Catenin Pathway to Regulate the Malignant Behavior of Bladder Cancer Cells. *Oncol Res*. 2018 Jan 19;26(1):145-155. Epub 2017 Mar 23 ([PMID: 28337959](#))
187. Wang RF, Wang HY (2016) Immune targets and neoantigens for cancer immunotherapy and precision medicine. *Cell Res* 2017 Jan;27(1):11-37 ([PMID: 28025978](#))
188. Watanabe T, Wu TT, Catalano PJ, Ueki T, Satriano R, Haller DG, Benson AB 3rd, Hamilton SR (2001) Molecular predictors of survival after adjuvant chemotherapy for colon cancer. *N Engl J Med* 2001 Apr 19;344(16):1196-206 ([PMID: 11309634](#))
189. Wei XL, Wang DS, Xi SY, Wu WJ, Chen DL, Zeng ZL, Wang RY, Huang YX, Jin Y, Wang F, Qiu MZ, Luo HY, Zhang DS, Xu RH (2015) Clinicopathologic and prognostic relevance of ARID1A protein loss in colorectal cancer. *World J Gastroenterol* 2014 Dec 28;20(48):18404-12 ([PMID: 25561809](#))
190. Wiegand KC, Shah SP, Al-Agha OM, Zhao Y, Tse K, Zeng T, Senz J, McConechy MK, Anglesio MS, Kalloger SE, Yang W, Heravi-Moussavi A, Giuliany R, Chow C, Fee J, Zayed A, Prentice L, Melnyk N, Turashvili G, Delaney AD, Madore J, Yip S, McPherson AW, Ha G, Bell L, Fereday S, Tam A, Galletta L, Tonin PN, Provencher D, Miller D, Jones SJ, Moore RA, Morin GB, Oloumi A, Boyd N, Aparicio SA, Shih IeM, Mes-Masson AM, Bowtell DD, Hirst M, Gilks B, Marra MA, Huntsman DG (2010) ARID1A mutations in endometriosis-associated ovarian carcinomas. *N Engl J Med*. 2010 Oct 14;363(16):1532-43. Epub 2010 Sep 8 ([PMID: 20942669](#))
191. Wielders E, Delzenne-Goette E, Dekker R, van der Valk M, Te Riele H (2017) Truncation of the MSH2 C-terminal 60 amino acids disrupts effective DNA mismatch repair and is causative for Lynch syndrome. *Fam Cancer*. 2017 Apr;16(2):221-229 ([PMID: 27873144](#))
192. Williamson CT, Miller R, Pemberton HN, Jones SE, Campbell J, Konde A, Badham N, Rafiq R, Brough R, Gulati A, Ryan CJ, Francis J, Vermulen PB, Reynolds AR, Reaper PM, Pollard JR, Ashworth A, Lord CJ (2016) ATR inhibitors as a synthetic lethal therapy for tumours deficient in ARID1A. *Nat Commun* 2016 Dec 13;7:13837 ([PMID: 27958275](#))
193. Wooten MW, Seibenhener ML, Mamidipudi V, Diaz-Meco MT, Barker PA, Moscat J (2001) The atypical protein kinase C-interacting protein p62 is a scaffold for NF-kappaB activation by nerve growth factor. *J Biol Chem*. 2001 Mar 16;276(11):7709-12. Epub 2001 Jan 22 ([PMID: 11244088](#))
194. Wu YL, Planchard D, Lu S, Sun H, Yamamoto N, Kim DW, Tan DSW, Yang JC, Azrif M, Mitsudomi T, Park K, Soo RA, Chang JWC, Alip A, Peters S, Douillard JY (2019) Pan-Asian adapted Clinical Practice Guidelines for the management of patients with metastatic non-small-cell lung cancer: a CSCO-ESMO initiative endorsed by JSMO, KSMO, MOS, SSO and TOS. *Ann Oncol*. 2019 Feb 01;30(2):171-210 ([PMID: 30596843](#))
195. Xie C, Fu L, Han Y, Li Q, Wang E (2014) Decreased ARID1A expression facilitates cell proliferation and inhibits 5-fluorouracil-induced apoptosis in colorectal carcinoma. *Tumour Biol* 2014 Aug;35(8):7921-7 ([PMID: 24833095](#))
196. Yang J, Wei X, Wu Q, Xu Z, Gu D, Jin Y, Shen Y, Huang H, Fan H, Chen J (2011) Clinical significance of the expression of DNA methyltransferase proteins in gastric cancer. *Mol Med Rep*. 2011 Nov-Dec;4(6):1139-43. Epub 2011 Aug 31 ([PMID: 21887466](#))
197. Yarchoan M, Hopkins A, Jaffee EM (2017) Tumor Mutational Burden and Response Rate to PD-1 Inhibition. *N Engl J Med*. 2017 Dec 21;377(25):2500-2501 ([PMID: 29262275](#))
198. Yarchoan M, Huang CY, Zhu Q, Ferguson AK, Durham JN, Anders RA, Thompson ED, Rozich NS, Thomas DL 2nd, Nauroth JM, Rodriguez C, Osipov A, De Jesus-Acosta A, Le DT, Murphy AG, Laheru D, Donehower RC, Jaffee EM, Zheng L, Azad NS (2019) A phase 2 study of GVAX colon vaccine with cyclophosphamide and pembrolizumab in patients with mismatch repair proficient advanced colorectal cancer. *Cancer Med* 2020 Feb;9(4):1485-1494 ([PMID: 31876399](#))
199. Ye J, Hameed O, Findeis-Hosey JJ, Fan L, Li F, McMahon LA, Yang Q, Wang HL, Xu H (2011) Diagnostic utility of PAX8, TTF-1 and napsin A for discriminating metastatic carcinoma from primary adenocarcinoma of the lung. *Biotech Histochem*. 2012 Jan;87(1):30-4. Epub 2011 Aug 15 ([PMID: 21838611](#))
200. Ye J, Zhou Y, Weiser MR, Gönen M, Zhang L, Samdani T, Bacares R, DeLair D, Ivelja S, Vakiani E, Klimstra DS, Soslow RA, Shia J (2014) Immunohistochemical detection of ARID1A in colorectal carcinoma: loss of staining is associated with sporadic microsatellite unstable tumors with medullary histology and high TNM stage. *Hum Pathol* 2014 Dec;45(12):2430-6 ([PMID: 25311944](#))
201. Yoshino T, Petheroudakis G, Mishima S, Overman MJ, Yeh KH, Baba E, Naito Y, Calvo F, Saxena A, Chen LT, Takeda M, Cervantes A, Taniguchi H, Yoshida K, Kodera Y, Kitagawa Y, Taberner J, Burris H, Douillard JY (2020) JSCO-ESMO-ASCO-JSMO-TOS: international expert consensus recommendations for tumour-agnostic treatments in patients with solid tumours with microsatellite instability or NTRK fusions. *Ann Oncol*. 2020 Jul;31(7):861-872. Epub 2020 Apr 6 ([PMID: 32272210](#))
202. Zhang R, Qin W, Xu GL, Zeng FF, Li CX (2011) A meta-analysis of the prevalence of somatic mutations in the hMLH1 and hMSH2 genes in colorectal cancer. *Colorectal Dis* 2012 Mar;14(3):e80-9 ([PMID: 21988782](#))
203. Zhang Z, Fan HY, Goldman JA, Kingston RE (2007) Homology-driven chromatin remodeling by human RAD54. *Nat Struct Mol Biol*. 2007 May;14(5):397-405. Epub 2007 Apr 8 ([PMID: 17417655](#))
204. de Bono J, Ramanathan RK, Mina L, Chugh R, Glaspy J, Rafii S, Kaye S, Sachdev J, Heymach J, Smith DC, Henshaw JW, Herriott A, Patterson M, Curtin NJ, Byers LA, Wainberg ZA (2017) Phase I, Dose-Escalation, Two-Part Trial of the PARP Inhibitor Talazoparib in Patients with Advanced Germline BRCA1/2 Mutations and Selected Sporadic Cancers. *Cancer Discov*. 2017 Jun;7(6):620-629. Epub 2017 Feb 27 ([PMID: 28242752](#))

205. U.S. Food and Drug Administration. Ipilimumab. https://www.accessdata.fda.gov/drugsatfda_docs/label/2020/125377s119lbl.pdf
206. U.S. Food and Drug Administration. Nivolumab. https://www.accessdata.fda.gov/drugsatfda_docs/label/2021/125554s090lbl.pdf
207. U.S. Food and Drug Administration. Pembrolizumab. https://www.accessdata.fda.gov/drugsatfda_docs/label/2020/125514s088lbl.pdf
208. (2020) Non-Small Cell Lung Cancer NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines) for Non-Small Cell Lung Cancer V.8.2020 https://www.nccn.org/professionals/physician_gls/pdf/nscl.pdf