

## ORIGINAL ARTICLE

# Prognosis of microsatellite instability and/or mismatch repair deficiency stage III colon cancer patients after disease recurrence following adjuvant treatment: results of an ACCENT pooled analysis of seven studies

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**Background:** Microsatellite instable/deficient mismatch repair (MSI/dMMR) metastatic colorectal cancers have been reported to have a poor prognosis. Frequent co-occurrence of MSI/dMMR and *BRAF*<sup>V600E</sup> complicates the association.

**Patients and methods:** Patients with resected stage III colon cancer (CC) from seven adjuvant studies with available data for disease recurrence and MMR and *BRAF*<sup>V600E</sup> status were analyzed. The primary end point was survival after recurrence (SAR). Associations of markers with SAR were analyzed using Cox proportional hazards models adjusted for age, gender, performance status, T stage, N stage, primary tumor location, grade, *KRAS* status, and timing of recurrence.

**Results:** Among 2630 patients with cancer recurrence (1491 men [56.7%], mean age, 58.5 [19–85] years), multivariable analysis revealed that patients with MSI/dMMR tumors had significantly longer SAR than did patients with microsatellite stable/proficient MMR tumors (MSS/pMMR) (adjusted hazard ratio [aHR], 0.82; 95% CI [confidence interval], 0.69–0.98; *P* = 0.029). This finding remained when looking at patients treated with standard oxaliplatin-based adjuvant chemotherapy regimens only (aHR, 0.76; 95% CI, 0.58–1.00; *P* = 0.048). Same trends for SAR were observed when analyzing MSI/dMMR versus MSS/pMMR tumor subgroups lacking *BRAF*<sup>V600E</sup> (aHR, 0.84; *P* = 0.10) or those harboring *BRAF*<sup>V600E</sup> (aHR, 0.88; *P* = 0.43), without reaching statistical significance. Furthermore, SAR was significantly shorter in tumors with *BRAF*<sup>V600E</sup> versus those lacking this mutation (aHR, 2.06; 95% CI, 1.73–2.46; *P* < 0.0001), even in the subgroup of MSI/dMMR tumors (aHR, 2.65; 95% CI, 1.67–4.21; *P* < 0.0001). Other factors associated with a shorter SAR were as follows: older age, male gender, T4/N2, proximal primary tumor location, poorly differentiated adenocarcinoma, and early recurrence.

**Conclusions:** In stage III CC patients recurring after adjuvant chemotherapy, and before the era of immunotherapy, the MSI/dMMR phenotype was associated with a better SAR compared with MSS/pMMR. *BRAF*<sup>V600E</sup> mutation was a poor prognostic factor for both MSI/dMMR and MSS/pMMR patients.

**Trial identification numbers:** NCT00079274, NCT00265811, NCT00004931, NCT00004931, NCT00026273, NCT00096278, NCT00112918.

**Key words:** colon cancer, microsatellite instability, deficient mismatch repair, recurrence, prognosis

## Introduction

Surgery alone or combined with adjuvant chemotherapy remains the cornerstone of treatment of non-metastatic colorectal cancer (CRC) [1–8]. Chromosomal instability and microsatellite instability (MSI) are distinct, well-described pathways of colorectal carcinogenesis that confer a different prognosis [9]. MSI/deficient DNA mismatch repair (MSI/dMMR) is considered a favorable prognostic factor in non-metastatic patients (stage I to III) tumors [9]; however, its prognostic role in metastatic CRC (mCRC) patients remains controversial.

A recent meta-analysis suggested that MSI/dMMR is a poor prognostic factor in patients eligible for the first-line treatment of non-resectable mCRC [10]. In this analysis, however, number of patients in each subgroup were limited which precluded definitive conclusions. Moreover, MSI/dMMR CRCs are often mutated for *BRAF*<sup>V600E</sup>, an established poor prognostic factor in mCRC that complicates the picture.

Clinical trials testing immunotherapy for MSI/dMMR mCRC patients, and targeted treatment of *BRAF*<sup>V600E</sup> mCRC patients, have yielded promising results. Having a clearer picture of the prognosis of each of these molecular subgroups is important given that approaches to their treatment is rapidly changing.

Among patients treated in seven phase III trials of adjuvant chemotherapy, we analyzed overall survival after disease recurrence in relationship to MMR and *BRAF*<sup>V600E</sup> status.

## Materials and methods

### Study population and patients characteristics

Histologically proven stage III colon adenocarcinomas had been completely resected from all eligible patients in a pooled analysis of seven adjuvant trials: MOSAIC, NCCTG NO147, PETACC8, PETACC3, NSABP C07, NSABP C08, and AVANT [2–8]. Patients were randomly assigned to receive 6 months of different regimens with fluoropyrimidines alone or with CPT11, or with oxaliplatin ± targeted therapy (bevacizumab, cetuximab), with regular monitoring, as described previously. Main results of all trials have been previously published. Among the 16,120 randomized patients, 4861 recurred and patients who had consented to translational research were tested for *BRAF*<sup>V600E</sup> and MMR status.

### DNA extraction and BRAF mutation analysis

Tumor samples were prospectively banked in all trials except for MOSAIC for which tumor banking was done retrospectively.

Tumor DNA was extracted from FFPE tissues containing more than 50% of tumor cells using DNA extraction kits. Molecular analysis was carried out retrospectively. *BRAF*<sup>V600E</sup> (c.1799T>A/p.V600E) was detected by allele specific real-time PCR. All assays were alteration-

specific and robustly detect ≥10% of mutated alleles for all mutations tested.

### Microsatellite status determination

Mismatch repair (MMR) tumor status was determined by immunohistochemistry, or by MSI testing. MSI tumors were defined as showing loss of the expression of one or more MMR proteins or exhibiting high-level MSI by PCR testing. Microsatellite stable (MSS) tumors had normal MMR protein expression and/or MSS or low-level MSI status.

### Statistical analyses

The primary outcome is survival after recurrence (SAR), defined as the time from recurrence to death due to all causes. Due to potential confounding and heterogeneities across studies, all analyses were based on multivariable models which adjusted for clinicopathologic variables, time to recurrence, and stratified per treatment groups within each study. The distribution of SAR between patient subgroups by biomarkers was estimated based on direct adjusted survival curves [11, 12]. Models were adjusted for age, sex, performance score, initial T/N stage, histologic grade, time from initial treatment to recurrence, primary tumor site and biomarkers when applicable. Two-sided P values are reported; P < .05 was considered statistically significant and was not adjusted for multiple comparisons. Analyses were carried out using SAS software (version 9.4; SAS Institute Inc).

## Results

### Study population

Among the 4861 patients randomized in the seven trials with disease recurrence, 2630 had consented to translational research with available material and thus, had complete data for *BRAF*<sup>V600E</sup> and MMR status (see CONSORT diagram, [supplementary Figure S1](#), available at *Annals of Oncology* online). In the CONSORT diagram ([supplementary Figure S1](#), available at *Annals of Oncology* online) the 2230 patients excluded due to not having MSI or *BRAF* results are mixing patients for which no translational informed consent was available and patients for which *BRAF* or MSI status was not carried out due to inadequate materials or testings.

Demographic and clinical characteristics of the patients in the molecular study ( $n = 2630$ ) and the others ( $n = 2231$ ) are summarized in [supplementary Table S1](#), available at *Annals of Oncology* online.

Among the 2630 patients with MMR and *BRAF* data, 307 (11.7%) tumors harbored *BRAF*<sup>V600E</sup> and 271 (10.3%) were MSI/dMMR. Among MSI/dMMR tumors, 91 (33.6%) harbored *BRAF*<sup>V600E</sup>, and among *BRAF*<sup>V600E</sup> mutants, 91 (29.6%) were MSI/dMMR.

**Table 1. Multivariable associations between patient demographics and disease characteristics with survival after recurrence (SAR), adjusting for biomarkers (MSI/MMR and BRAF)<sup>a</sup>**

	Events/total	Hazard ratio (95% CI)	P-value <sup>b</sup>
Age, 5-year increase	1428/1987	1.04 (1.01–1.07)	0.0057
Gender			
Female	614/860	0.88 (0.79–0.98)	0.0218
Male	814/1127	Reference	
Primary tumor location			
Distal	657/1026	0.64 (0.57–0.72)	<0.0001
Proximal	771/961	Reference	
T-stage			
T1/2	60/104	Reference	
T3	1048/1469	1.20 (0.92–1.57)	0.1746
T4	320/414	1.45 (1.09–1.92)	0.0102
N-Stage			
N1	583/890	Reference	
N2	845/1097	1.36 (1.22–1.51)	<0.0001
Histologic grade			
Low grade (grade 1–2)	1022/1482	Reference	
High grade (grade 3/4)	406/505	1.36 (1.20–1.53)	<0.0001
KRAS status			
MT	597/791	1.21 (1.07–1.36)	0.0023
WT	831/1196	Reference	
Performance score			
0	1077/1519	Reference	
1	341/457	1.18 (1.04–1.34)	0.0089
2	10/11	3.54 (1.87–6.70)	0.0001
Time-to-recurrence, 1-year increase	1428/1987	0.87 (0.82–0.91)	<0.0001

<sup>a</sup>A single multivariable Cox model was fitted with variables listed in the table, plus MSI/MMR and BRAF variables. The HRs associated with MSI/MMR and BRAF are reported in Table 2 (first two sections).

<sup>b</sup>Stratified covariate Wald P-value.

MSI, microsatellite instable; MMR, mismatch repair; BRAF; CI, confidence interval; KRAS; MT, mutated; WT, wild type.

Clinical and pathological patients' characteristics per study are summarized in [supplementary Table S2](#), available at *Annals of Oncology* online.

### Demographic and clinical characteristics according to MSI/dMMR and BRAF status

Patients with tumor recurrence and MSI/dMMR were more likely to be females and to have had a proximal primary tumor that was poorly differentiated, pT4/N2,  $BRAF^{V600E}$ , had more lymph nodes examined, and were less frequently *KRAS* mutated ([supplementary Table S3](#), available at *Annals of Oncology* online).

Patients with  $BRAF^{V600E}$  tumors were more frequently females and to have had a proximal primary tumor that was poorly differentiated, pT4/N2, MSI/dMMR, and non *KRAS* mutated ([supplementary Table S4](#), available at *Annals of Oncology* online).

### Outcome in different molecular subgroups

In the overall population with recurrence ( $N=4861$ ), median follow-up was 77.3 months (95% confidence interval [CI] 75.4–80.5) and median SAR was 23.1 months (95% CI 22.3–23.9).

A single multivariable Cox model was fitted with variables listed in Tables 1 and 2 (clinic-pathologic variables plus MSI/MMR and BRAF variables). To better outline our results, the HRs associated with MSI/MMR and BRAF are reported in Table 2 (first two sections) and the median SAR and HRs are only reported for MMR and BRAF variables. In multivariable analysis adjusted for MSI/dMMR and  $BRAF^{V600E}$  status, factors associated with a poor SAR were as follows: older age, male gender, T4/N2, proximal primary tumor location, poor differentiation, and early recurrence (by 1-year increase) (Table 1).

Multivariable analysis revealed that patients with MSI/dMMR tumors had significantly better SAR than did patients with MSS/proficient (p)MMR tumors (Table 2, Figure 1). This was also observed among patients treated with standard adjuvant fluoropyrimidine+oxaliplatin only. The same trends were observed when analyzing patients with MSI/dMMR tumors lacking  $BRAF^{V600E}$  and mutant subgroups separately, without achieving statistical significance (Table 2).

As previously described, poor SAR was observed in  $BRAF^{V600E}$  versus nonmutated patients and this was also found in the subgroup of MSI/dMMR patients (Table 2).

**Table 2. Multivariable associations between patient biomarkers (MMR and BRAF status) with survival after recurrence (SAR), adjusting for demographics and disease characteristics<sup>a</sup>**

SAR	Adjusted survival			
	Event/total	Median (95% CI) <sup>KM</sup>	Hazard ratio (95% CI) <sup>Cox</sup>	P-value
MMR status				0.0290 <sup>b</sup>
dMMR	162/220	2.2 (1.9–2.7)	0.82 (0.69–0.98)	
pMMR	1266/1767	2.0 (1.9–2.2)	Reference	
BRAF status				<0.0001 <sup>b</sup>
MT	219/244	1.2 (0.9–1.4)	2.06 (1.73–2.46)	
WT	1209/1743	2.2 (2.1–2.4)	Reference	
dMMR patients only				<0.0001 <sup>b</sup>
BRAF status				
MT	64/77	0.8 (0.5–1.1)	2.65 (1.67–4.21)	
WT	98/143	1.9 (1.7–2.5)	Reference	
pMMR patients only				<0.0001 <sup>b</sup>
BRAF status				
MT	155/167	1.3 (0.9–1.5)	2.12 (1.74–2.58)	
WT	1111/1600	2.3 (2.2–2.4)	Reference	
BRAF WT patients only				0.1030 <sup>b</sup>
MMR status				
dMMR	98/143	2.4 (1.9–3.5)	0.84 (0.67–1.04)	
pMMR	1111/1600	2.3 (2.2–2.4)	Reference	
BRAF MT patients only				0.4299 <sup>b</sup>
MMR status				
dMMR	64/77	0.8 (0.7–1.1)	0.88 (0.63–1.22)	
pMMR	155/167	0.9 (0.8–1.2)	Reference	
Adjuvant FP + oxaliplatin patients only				0.0476 <sup>b</sup>
MMR status				
dMMR	66/92	2.5 (1.8–4.6)	0.76 (0.58–1.00)	
pMMR	537/771	2.0 (1.9–2.2)	Reference	

<sup>a</sup>Multivariable Cox models were fitted on all patients, and subgroups of patients as indicated. All models were adjusted by age, gender, PS, T-stage, N-stage, tumor location, histological grade, KRAS and years to progression. The median survival after recurrence and HRs were only reported for MMR and BRAF variables.

<sup>b</sup>Likelihood-ratio test.

<sup>KM</sup>Kaplan–Meier method; <sup>Cox</sup>Cox model.

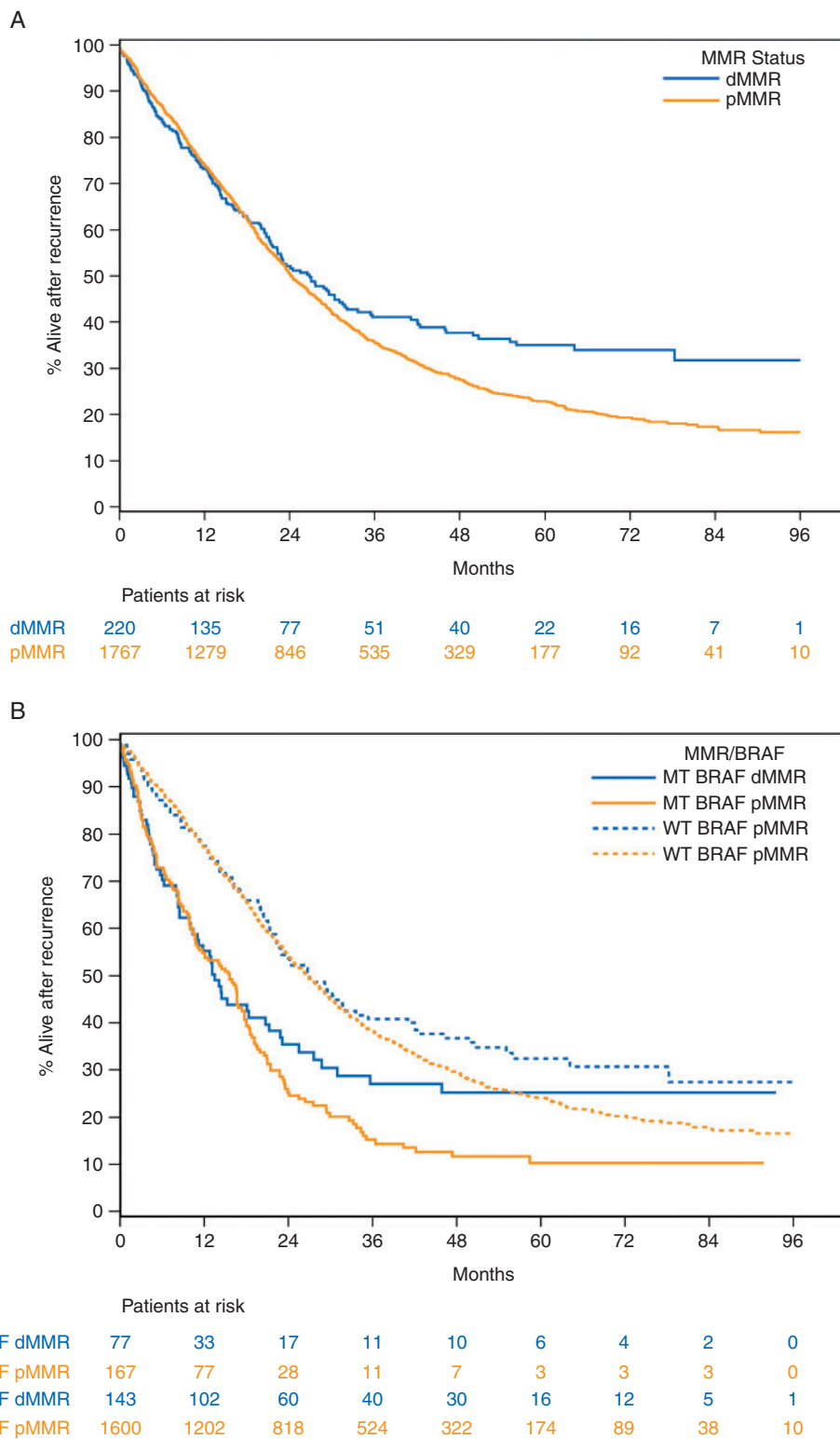
CI, confidence interval; dMMR, deficient mismatch repair; pMMR, proficient mismatch repair; FP, fluoropyrimidine; MT, mutated; WT, wild type.

## Discussion

We report the largest study examining the role of MMR and *BRAF*<sup>V600E</sup> status among patients ( $n = 2630$ ) with colon cancer recurrence following surgery and adjuvant chemotherapy. Among recurrent colon cancers, MSI/dMMR and *BRAF*<sup>V600E</sup> each had a low prevalence (around 10%). Whereas *BRAF*<sup>V600E</sup> indicates a poor prognosis, we found that MSI/dMMR status was associated with a better SAR. Moreover, the association of *BRAF*<sup>V600E</sup> with poor prognosis was found in both MSI/dMMR and MSS/pMMR patients. Most studies report a poor prognostic value of *BRAF*<sup>V600E</sup> in CRC, and in MSI/dMMR patients in the metastatic setting [10]. The general explanation for the poor prognosis of MSI/dMMR tumors in the metastatic setting, as compared with their favorable prognosis in the adjuvant setting, is based on possible immuneescape and immuno-editing processes with strong immune infiltration [13]. MSI/dMMR tumors

that recur appear to have escaped immune surveillance by diverse cellular and molecular processes, and thus tend to be more aggressive. Marissa et al. demonstrated recently that immune checkpoint expression cancels the prognostic relevance of tumor-infiltrating T cells in highly immunogenic colon tumors and predicts a poor outcome in MSI CRC patients [14]. Another explanation is the overlap between MSI and the poor prognostic *BRAF*<sup>V600E</sup> mutations.

This work has however some limitations including the fact that only 54% of recurring patients were analyzed due to missing patients consents of inadequate *BRAF* and MMR testing. Moreover, in contrast to prior reports, all patients had subsequent recurrent disease and were treated with previous adjuvant therapy. This may explain why our results differ compared with studies of patients with non-resectable mCRC eligible for a first-line trial [10]. However, a recent study of mCRC patients enrolled



**Figure 1.** Survival after recurrence in patients with resected stage III colon cancer according to the MMR and  $BRAF^{V600E}$  mutational status, adjusted for age, gender, PS, T-stage, N-stage, primary, grade, KRAS, recurrence years.

in a first-line study (CALGB/SWOG 80405) found that MMR status was not a poor prognostic factor [15].

In summary, our data demonstrate in a large cohort of MSI/dMMR patients with recurrence and before the era of immuno-

oncologic treatments, that MSI/dMMR (versus MSS/pMMR) status is associated with a longer survival. Furthermore,  $BRAF^{V600E}$  was observed to be a poor prognostic factor for both MSI/dMMR and MSS/pMMR patients. As both MSI and  $BRAF$

status influence recurring colon cancer patients' outcome, these factors should be used to stratify patients in future clinical trials dedicated to MSI or BRAF mutant mCRC.

## Acknowledgement

Research reported in this publication was supported by the National Cancer Institute of the National Institutes of Health under Award Number U10CA180882. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health.

## Funding

This work was supported by the National Cancer Institute at the National Institutes of Health (grant number U10CA180882).

## Disclosure

JT has received honoraria for speaker or/and advisory role from Merck KGaA, Sanofi, Roche Genentech, MSD, Lilly, Celgene, Servier, Pierre Frabre and Amgen. TA has participated in consulting or/and advisory boards for Astra-Zeneca, Amgen, BMS, MSD Oncology, HalioDx, Roche/Ventana, Sanofi, Sevier. FS has participated in consulting and/or advisory boards for Roche/Ventana Medical Systems and HalioDx. TY reports research funding from Chugai Pharmaceutical Co., Ltd., Sanofi, 423 K.K., and Sumitomo Dainippon Pharma Co., Ltd. All remaining authors have declared no conflicts of interest.

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