

Biomarkers predictive for outcome in patients with metastatic colorectal cancer treated with 1st-line FOLFOX4 ± cetuximab: Updated data from the OPUS study

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Background

- In the OPUS study patients with *KRAS* wild-type (*KRAS* wt) metastatic colorectal cancer (mCRC) receiving cetuximab plus FOLFOX4 displayed a significantly higher best overall response (OR), 61% vs 37%, (odds ratio 2.54, p=0.011) and a lower risk of disease progression (hazard ratio [HR] 0.57, p=0.0163) compared with patients receiving FOLFOX4 alone.¹
- The OPUS study confirmed earlier retrospective investigations of single-arm studies suggesting that cetuximab efficacy was confined to patients with *KRAS* wt tumors.^{2,3}
- The serine-threonine kinase BRAF is a direct downstream effector of *KRAS*. *BRAF* gene mutations have been detected in around 8% of colorectal tumors,⁴ and *BRAF* mutation status has been suggested to be predictive of cetuximab efficacy in pretreated patients.⁵
- Here we report overall survival (OS), updated progression-free survival (PFS) time and OR data for patients with *KRAS* wt tumors following an increase in the number of patients for whom *KRAS* tumor mutation status could be determined.
- The impact of *BRAF* tumor mutations on cetuximab efficacy was investigated in patients with *KRAS* wt disease.

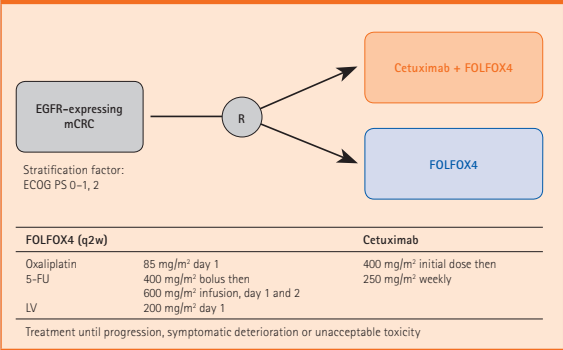
Study objectives

- Primary study objective:
 - Assess whether the OR of cetuximab plus FOLFOX4 was superior to that of FOLFOX4 alone
- Secondary objectives included:
 - Comparison of the treatment regimens with respect to PFS time and OS time and safety
- A retrospective subgroup analysis for associations between OR, PFS and OS times and the *KRAS/BRAF* mutation status of tumors.

Methods

- Study design
- This was an open label randomized multicenter phase II study (Figure 1).

Figure 1. Study design¹



5-FU, 5-fluorouracil; ECOG PS, Eastern Cooperative Oncology Group performance status; EGFR, epidermal growth factor receptor; LV, leucovorin; mCRC, metastatic colorectal cancer; R, randomization.

Patients

- Patient inclusion and exclusion criteria have been described elsewhere.¹

KRAS/BRAF tumor mutation detection

- Ascertainment of samples for *KRAS* testing was increased using tumor DNA extracted from formalin fixed paraffin embedded slide mounted sections initially prepared to evaluate tumor EGFR expression.
- *KRAS* mutations at codons 12/13 and *BRAF* (V600E) mutations were detected using the previously described polymerase chain reaction clamping and melting curve technique.¹

Data cut-offs for analyses

- OPUS study data cut-offs were: PFS (independent review committee), OR (independent review committee), 01 March 2007, and OS, 30 November 2008.

Results

Baseline characteristics (Table 1)

- Patient tumor numbers in the ITT population (n=337) for whom *KRAS* mutation status was determined increased from 233 (69%) as previously published¹ to 315 (93%).
- 309 (92%) patient tumors were evaluable for *BRAF* mutation status.
- *BRAF* mutations were detected in 11/309 (4%) evaluable tumor samples from the ITT population.
 - No patients were found with *KRAS* mt and *BRAF* mutant (*BRAF* mt) tumors
- 179/315 (57%) patients had *KRAS* wt tumors.
- 175 *KRAS* wt tumors were evaluable for *BRAF* mutations.
 - 164 (94%) were *BRAF* wt
 - 11 (6%) were *BRAF* mt

Table 1. Baseline and disease characteristics in patients with <i>KRAS</i> wt tumors according to <i>BRAF</i> mutation status and treatment						
Characteristics, n (%) unless otherwise stated	<i>KRAS</i> wt (n=179)		<i>KRAS</i> wt/ <i>BRAF</i> wt (n=164)		<i>KRAS</i> wt/ <i>BRAF</i> mt (n=11)	
	FOLFOX4 (n=97)	Cetuximab + FOLFOX4 (n=82)	FOLFOX4 (n=92)	Cetuximab + FOLFOX4 (n=72)	FOLFOX4 (n=5)	Cetuximab + FOLFOX4 (n=6)
Gender						
Male	55 (57)	42 (51)	51 (55)	38 (53)	4 (80)	2 (33)
Female	42 (43)	40 (49)	41 (45)	34 (47)	1 (20)	4 (67)
Median age, years (range)	59 (36–82)	62 (24–75)	59 (37–82)	63 (24–75)	61 (36–74)	51 (40–74)
Age categories						
<65 years	63 (65)	46 (56)	60 (65)	39 (54)	3 (60)	5 (83)
≥65 years	34 (35)	36 (44)	32 (35)	33 (46)	2 (40)	1 (17)
ECOG PS 0/1	86 (89)	76 (93)	82 (89)	66 (92)	4 (80)	6 (100)
Liver metastases only	23 (24)	25 (31)	23 (25)	23 (32)	0	1 (17)
Involved disease sites ≤2	75 (77)	67 (82)	74 (80)	60 (83)	5 (100)	4 (67)
Prior adjuvant chemotherapy	21 (22)	13 (16)	21 (23)	12 (17)	0	1 (17)

ECOG PS, Eastern Cooperative Group performance status; mt, mutant; wt, wild-type.

Clinical efficacy (Table 2)

- For patients with *KRAS* wt tumors the addition of cetuximab to FOLFOX4 compared with FOLFOX4 alone led to:
 - Significant improvement in OR (Figure 2), with a mean difference in the best % change of the lesion from baseline (based on WHO criteria) of 11.6
 - Significant improvement in PFS time (Figure 3)
 - Improvement in OS time although the difference between the arms was not significant (Figure 4)
- Significant interactions between treatment outcomes and *KRAS* tumor mutation status were observed for OR and PFS (<0.001) but not for OS (p=0.12).⁶
- Significant improvements in PFS and OR were also found on adding cetuximab to FOLFOX4 for patients with *KRAS* wt/*BRAF* wt tumors; the improvement in OS was not significant (Table 2).

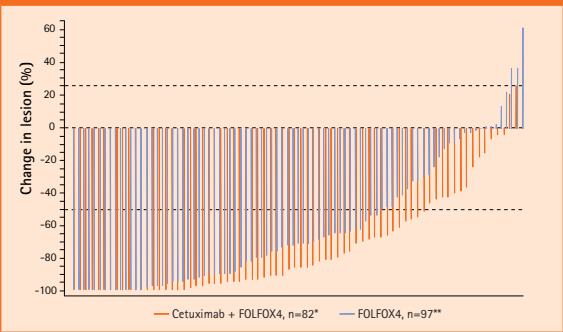
Post-study treatment

- The proportion of patients receiving post study anticancer therapy has been previously reported.⁶
- 54 (66%) patients in the cetuximab + FOLFOX4 arm and 59 (61%) in the FOLFOX4 alone arm received treatment with anticancer therapy of any kind.
 - Most frequent therapy in both arms was 5-FU/FA (41% vs 44%)
 - 10% and 16% of patients received post-study cetuximab, respectively

Table 2. Efficacy data in patients with <i>KRAS</i> wt tumors according to <i>BRAF</i> mutation status and treatment						
	<i>KRAS</i> wt (n=179)		<i>KRAS</i> wt/ <i>BRAF</i> wt (n=164)		<i>KRAS</i> wt/ <i>BRAF</i> mt (n=11)	
	FOLFOX4 (n=97)	Cetuximab + FOLFOX4 (n=82)	FOLFOX4 (n=92)	Cetuximab + FOLFOX4 (n=72)	FOLFOX4 (n=5)	Cetuximab + FOLFOX4 (n=6)
Tumor response						
OR rate (%)	34.0	57.3	35.9	59.7	0	33.3
[95%CI]	[24.7–44.3]	[45.9–68.2]	[26.1–46.5]	[47.5–71.1]	[0.0–52.2]	[4.3–77.7]
Odds ratio		2.5512		2.6491		NA
[95% CI]		[1.3799–4.7169]		[1.3901–5.0483]		NA
p-value*		0.0027		0.0029		0.2207
PFS time						
Median PFS, months	7.2	8.3	7.2	8.3	1.7	7.1
[95% CI]	[5.6–7.4]	[7.2–12.0]	[5.6–7.4]	[7.3–12.7]	[0.9–7.9]	[4.2–NA]
Hazard ratio		0.567		0.556		0.449
[95% CI]		[0.375–0.856]		[0.358–0.864]		[0.087–2.303]
p-value*		0.0064		0.0083		0.3255
OS time						
Median OS, months	18.5	22.8	19.5	22.8	4.4	20.7
[95%CI]	[16.4–22.6]	[19.3–25.9]	[17.0–23.8]	[19.3–25.8]	[0.9–10.1]	[10.3–30.4]
Hazard ratio		0.855		0.894		0.104
[95% CI]		[0.599–1.219]		[0.615–1.301]		[0.011–0.964]
p-value*		0.3854		0.5582		0.0167

*Stratified Cochrane-Mantel-Haenszel test; *Stratified log-rank test.
CI, confidence interval; mt, mutant; NA, not available; OR, best overall response; OS, overall survival; PFS, progression-free survival; wt, wild-type.

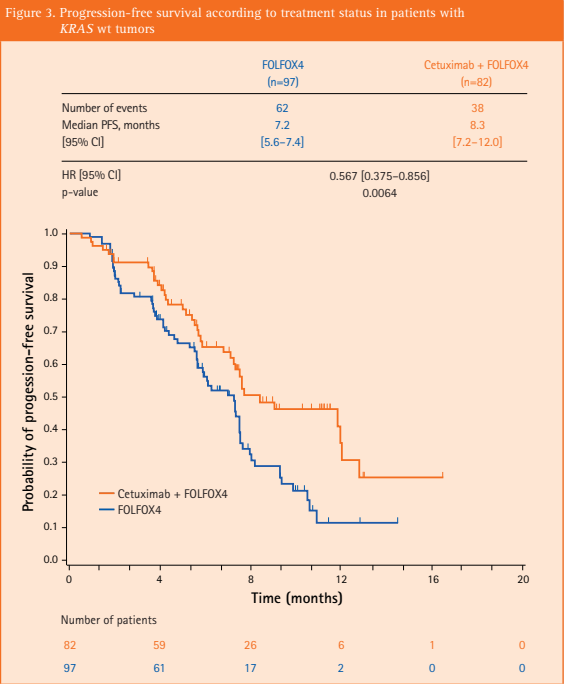
Figure 2. Tumor regression according to treatment status in patients with *KRAS* wt tumors



*Data are missing for 5 patients. **Data are missing for 4 patients.
wt, wild-type.

Safety data for patients with *KRAS* wt/*BRAF* wt tumor mutation status (Table 3)

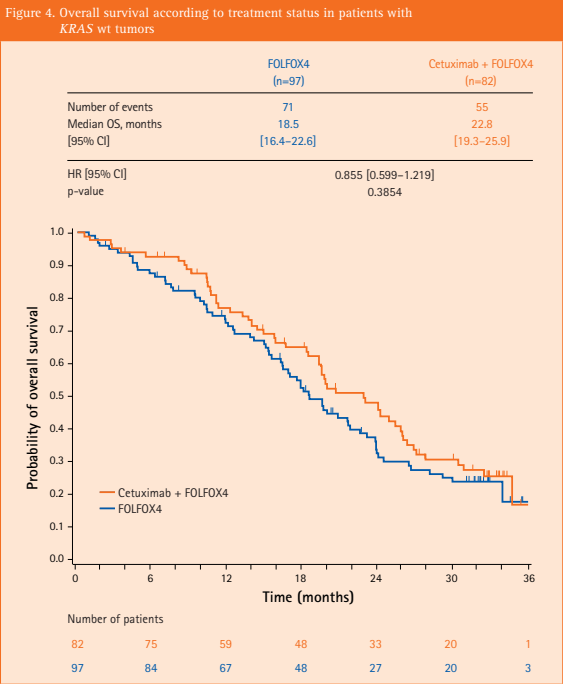
- Safety data have been reported previously for the ITT¹ and *KRAS* wt⁶ populations and were found to be generally similar for patients with *KRAS* wt/*BRAF* wt tumors.
- In the cetuximab + FOLFOX4 arm
 - 5 patients (7%) discontinued cetuximab only
 - 17 patients (24%) discontinued chemotherapy only
 - 8 patients (11%) discontinued both cetuximab and chemotherapy together
- In the FOLFOX4 alone arm 26 patients (28%) discontinued chemotherapy.
- 46 patients (64%) died of disease progression in the cetuximab-containing arm and 63 (68%) in the chemotherapy-alone arm. There were no cetuximab-related deaths.



CI, confidence interval; HR, hazard ratio; PFS, progression-free survival; wt, wild-type.

Table 3. Grade 3 or 4 adverse events and special adverse events occurring in patients with <i>KRAS</i> wt tumors according to <i>BRAF</i> mutation status and treatment				
Adverse events*, n (%)	<i>KRAS</i> wt (n=179)		<i>KRAS</i> wt/ <i>BRAF</i> wt (n=164)	
	FOLFOX4 (n=97)	Cetuximab + FOLFOX4 (n=82)	FOLFOX4 (n=92)	Cetuximab + FOLFOX4 (n=72)
Any grade 3/4 event	62 (64)	67 (82)	58 (63)	72 (81)
Neutropenia	31 (32)	29 (35)	30 (33)	26 (36)
Rash**	0	9 (11)	0	8 (11)
Diarrhea**	5 (5)	7 (9)	5 (5)	5 (7)
Peripheral sensory neuropathy**	8 (8)	3 (4)	8 (9)	3 (4)
Leukopenia	5 (5)	6 (7)	5 (5)	6 (8)
Pulmonary embolism	1 (1)	4 (5)	1 (1)	4 (6)
Composite categories*, n (%)				
Skin reactions**	0	15 (18)	0	12 (17)
Neurotoxicity associated events**	0	6 (7)	14 (15)	6 (8)
Cardiac events	0	6 (7)	0	3 (4)

*Reported in >5% of patients with *KRAS* wt tumors in either treatment arm; **No grade 4 events recorded;
*Including terms as defined in the Medical Dictionary for Regulatory Activities.
wt, wild-type.



CI, confidence interval; HR, hazard ratio; OS, overall survival; wt, wild-type.

Conclusions

- The addition of cetuximab to FOLFOX4 in patients with *KRAS* wt tumors led to significantly longer PFS and a significantly higher OR than in patients receiving FOLFOX4 alone.
- In patients with *KRAS* wt tumors, the addition of cetuximab to FOLFOX4 in the 1st-line treatment of mCRC led to an improvement in OS compared with patients receiving FOLFOX4 alone (18.5 to 22.8 months); this difference was not statistically significant which may be due to the small sample size.
- This analysis confirms *KRAS* tumor mutation status to be a predictive factor for the efficacy of cetuximab treatment in terms of OR and PFS.
- In the OPUS study the prevalence of *BRAF* mutations in patients with *KRAS* wt tumors (11 patients) prevents conclusions being drawn on its use as a predictive or prognostic biomarker.

References

1. Bokemeyer C, et al. J Clin Oncol 2009;27:663-71.
2. De Roock W, et al. Ann Oncol 2008;19:508-15.
3. Lièvre A, et al. J Clin Oncol 2008;26:374-9.
4. Roth A, et al. J Clin Oncol 2009 E-pub doi/10.1200/JCO.2009.23.3452.
5. Di Nicolantonio F, et al. J Clin Oncol 2008;26:5705-12.
6. Bokemeyer C, et al. Eur J Cancer Suppl 2009;7(2):346. Abstract P-6079.

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