

The Challenge of anti-EGFR therapy-rechallenge in management of metastatic colorectal cancer

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Treatment of metastatic colorectal cancer (mCRC) has undergone several changes thanks to the introduction of targeted therapies inhibiting several disregulatory pathways (1).

Epidermal growth factor receptor (EGFR), a tyrosine kinase receptor belonging to the ErbB family, is overexpressed in 25%-80% of CRCs and plays a major role in its pathogenesis (2) and many clinical trials proved the therapeutic efficacy of antibodies targeting EGFR (cetuximab and panitumumab) in the treatment of CRC patients (3). Therefore, the relatively recent introduction of those anti-EGFR antibodies, combined with standard chemotherapy, allowed to reach the median overall survival of 23-24 months.

Moreover, the median overall response rate to cetuximab or panitumumab based regimens is less than 30%, and several mechanisms of primary resistance to inhibition of EGFR pathway have been studied.

Retrospective and prospective analysis showed that early mutations of Kirsten Ras oncogene homologue (K-RAS) gene, occurring in 35-45% of CRC, are associated with primary resistance to anti-EGFR therapy (4-21). Although previous clinical trials have indicated that patients harboring KRAS mutations in codons 12 and 13 are non-responsive to the EGFR-targeted therapy, other analysis showed that some wild-type patients still fail to respond (22) and downstream mutations such as in BRAF, PIK3CA NRAS and rare K-Ras mutations have been investigated (23-29).

Cetuximab is currently approved in Europe as a first line treatment of KRAS WT mCRC, in combination with standard chemotherapy, as second line treatment in combination with irinotecan-based chemotherapy, or as monotherapy in irinotecan or oxaliplatin refractory patients; while panitumumab is approved for treatment of RAS WT mCRC as first line therapy in combination with FOLFOX, as a second line in combination with FOLFIRI and as a monotherapy in irinotecan and oxaliplatin-refractory patients (30).

But the improvement in management of mCRC patients due to introduction of anti-EGFR and other targeted therapies led to many outstanding issues: 1) there is a significant number of patients progressing beyond the third or fourth line of treatment still suitable for further therapy when enrolment into clinical trial is not possible; 2) Prolonged intensive treatment is burdened from the high risk of cumulative toxicity, worsening in quality of life and a not well defined possibility of early gaining of acquired resistance.

In this sense, rechallenge of an anti-EGFR, defined as reintroduction, after an intervening treatment, of the same therapy to which tumor has already proved to be resistant, could be a new approach for those heavily pretreated patients based on some important preclinical and clinical evidences.

In fact one study evaluated the variation of circulating tumor DNA (ctDNA) in serum of 24 patient receiving single-agent therapy with panitumumab. KRAS mutations were recorded in 38% of cases between 5-6 months following treatment and mathe-

mathematical modelling indicated that mutations were present in expanded subclones before the initiation of treatment. These results suggest that the selection and emergence of KRAS mutations is a possible mechanism of secondary acquired resistance to EGFR-inhibition (31). Consistently, another small study showed that point mutations of KRAS are casually associated with the onset of acquired resistance to anti-EGFR therapy. In fact analysis of metastasis from ten patients who developed resistance to cetuximab or panitumumab showed the emergence of KRAS mutant alleles were detectable in the blood months before the radiographic evidence of disease progression, and the *in vitro* model support the hypothesis of continuing mutagenesis (not only involving RAS gene, but also in B-RAF, PI3KCA genes) under the pressure of anti-EGFR therapy (32).

These studies represent an important evidence about the possibility of secondary acquisition of KRAS mutations due to the intensive anti-EGFR-based treatment and could support the strategy of rechallenge.

We can speculate that an interval therapy after first progression to the anti-EGFR therapy could restore a partial sensitivity of tumor to the rechallenge by promoting the re-expansion of RAS wild-type clones, which will constitute the major part of the tumor mass at the time of a following progression of disease. At that time, the readministration of anti-EGFR therapy may then determine a further disease response. Moreover an interval therapy based on a different treatment, which is not influenced by KRAS status or is more efficacious in KRAS mutated CRC, could facilitate the re-emersion of wt clones.

In fact, an *in vitro* model suggested that KRAS mutated cell lines are more sensitive to Oxaliplatin (33). Consistently, a retrospective study evaluating K-Ras status in 90 patients treated with FOLFOX-6 as first-line or second-line treatment showing that PFS was longer in mutated K-Ras population than in wt KRAS patients (10 vs 8 months, respectively; $p=0.001$) (34).

Another multicenter phase II prospective study evaluated the activity of a rechallenge with a cetuximab-based therapy in 39 patients who first had a clinical benefit after a line of cetuximab plus

irinotecan-based therapy, then a disease progression for which received a new line of chemotherapy and finally, after a further progression of disease, were re-treated with the same cetuximab plus irinotecan based therapy. Overall response rate (RR) was 53.8% with 19 partial responses (48.7%) and 2 complete responses (5.1%). The median time to progression (TTP) was 6.6 months, stable disease (SD) was obtained in 35.9% of patients and progression in 10.2% of cases; and 18 patients showed the same type of response (SD, partial response or complete response) during cetuximab retreatment when compared with the response obtained during the first cetuximab-based therapy. Then stable disease lasting at least 6 months and partial response during the first cetuximab-based therapy have been demonstrated to predict clinical benefit after cetuximab retreatment (35).

Conversely, a phase II prospective study including twenty patients treated with panitumumab after progression on prior cetuximab-based therapy did not show any response being stable disease (no progression for at least two cycles) the best response in 45% of patients (36). But no interval therapy or treatment holiday were permitted between cetuximab and panitumumab administration and it is not possible to know if primary refractory patients to anti-EGFR were enrolled (N-RAS and BRAF status were not evaluated) and data about response to prior cetuximab-based therapy were not available.

Therefore anti-EGFR rechallenge could be a valid therapeutic option in management of mCRC and should be investigated in randomized prospective trial enrolling a selected population of mCRC patients. This selection should be based on different reasonable factors: all-RAS and BRAF wt status, best response to prior treatment before progression (prolonged stable disease, partial response or complete response), residual toxicity and duration of treatment holiday.

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