

Anti-epidermal growth factor receptor monoclonal antibodies in cancer therapy

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Epidermal growth factor receptor

Epidermal growth factor receptor (EGFR) is a cell membrane growth factor receptor characterized by tyrosine kinase activity that plays a crucial role in the control of key cellular transduction pathways in both normal and cancerous cells. EGFR is over-expressed in a variety of human tumours, including head and neck, breast, lung, colorectal, prostate, kidney, pancreas, ovary, brain and bladder cancer [1–3].

The 170 kDa protein function depends either on the formation of EGFR – EGFR homodimers or heterodimers – that comprise the three members of the EGFR [human epidermal receptor 1 (HER1)] family of growth factor receptors (HER2, HER3 and HER4) following binding of an EGFR-selective ligand. The activating ligands include the epidermal growth factor (EGF), transforming growth factor- α (TGF- α), amphiregulin or neuregulin. The binding EGFR/ligand results in conformational changes that allow the activation of EGFR tyrosine kinase and the phosphorylation of specific tyrosine residues within the EGFR intracellular carboxyl-terminal domain. Phosphorylated tyrosine residues serve as docking sites for several signalling proteins finally stimulating cell proliferation, loss of differentiation, invasion, angiogenesis and blocking of apoptosis. Within a few hours of activation, receptors are internalized into cytoplasm, where they are either degraded or recycled back to the membrane.

Summary

The epidermal growth factor receptor (EGFR) is a transmembrane tyrosine kinase receptor involved in the proliferation and survival of cancer cells. EGFR is the first molecular target against which monoclonal antibodies (mAb) have been developed for cancer therapy. Here we review the mechanisms underlying the effects of EGFR-specific mAb in cancer therapy. The efficacy of EGFR-specific mAb in cancer occurs thanks to inhibition of EGFR-generated signalling; furthermore, the effects of antibodies on the immune system seem to play an important role in determining the overall anti-tumour response. In this review, attention is focused on cetuximab and panitumumab, two mAb introduced recently into clinical practice for treatment of metastatic colorectal and head and neck cancer which target the external part of EGFR.

Keywords: cancer, cetuximab, epidermal growth factor receptor, monoclonal antibodies, panitumumab

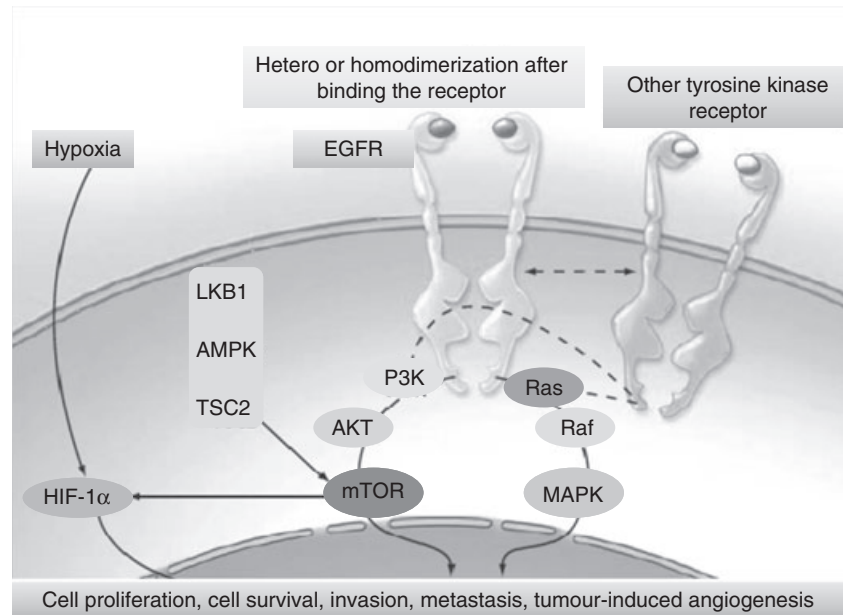
EGFR homodimers undergo degradation, whereas EGFR and HER2 heterodimerization is associated with recycling upon endocytosis, which enhances mitogenic signalling. Homodimers are weaker effectors compared with heterodimers: EGFR and HER2 is the most common heterodimer; HER2:3 plus neuregulin is the most potent combination; HER2 decelerates the internalization of HER1; HER1 requires ligand binding before dimerization, while HER2 does not require a ligand to dimerize and is often expressed at a 100-fold higher concentration compared with HER1. The complex signalling network generated by triggering EGFR includes the ras- and mitogen-activated protein kinase (MAPK) pathway that leads to cell proliferation, the phosphatidylinositol-3 kinase (PI3K) and protein kinase B (Akt) pathway driving cell cycle progression and cell survival [4]. There is also evidence that EGFR can translocate to the nucleus, where it acts as a transcription factor (Fig. 1) [5–7].

Inhibition of the target

Two pharmacological approaches have been used successfully to inhibit EGFR functions in cancer treatment: neutralizing monoclonal antibodies and small-molecule tyrosine kinase inhibitors.

Anti-EGFR monoclonal antibodies bind to the extracellular domain of EGFR in its inactive state; they compete for receptor binding by occluding the ligand-binding region,

Fig. 1. Signal transduction pathway mediated by epidermal growth factor receptor (EGFR). The interaction with ligand that occurs in the extracellular portion of the EGFR family induces the formation of a functionally active EGFR–EGFR homodimer or of an EGFR–human epidermal receptor 2 (HER2), EGFR–HER3 or EGFR–HER4 heterodimer. As explained in the text, this results in conformational changes that allow the EGFR tyrosine kinase to be activated with the phosphorylation of specific tyrosine residues within the EGFR intracellular carboxyl-terminal domain. Phosphorylated tyrosine residues trigger a complex programme of intracellular signals to the cytoplasm and then to the nucleus that stimulates cell proliferation, loss of differentiation, invasion and angiogenesis, and blocks the apoptosis.



and thereby block ligand-induced EGFR tyrosine kinase activation [8,9]. Small-molecule EGFR tyrosine kinase inhibitors compete reversibly with Adenosine 5' triphosphate to bind to the intracellular catalytic domain of EGFR tyrosine kinase and, thus, inhibit EGFR auto-phosphorylation and downstream signalling. In addition, various small-molecule EGFR tyrosine kinase inhibitors can block different growth factor receptor tyrosine kinases, including other members of the EGFR family, or the vascular endothelial growth factor receptor. Anti-EGFR monoclonal antibodies recognize EGFR exclusively and are therefore highly selective to this receptor. Nevertheless, an intrinsic or acquired resistance to the EGFR inhibitor that limits the use of these drugs in cancer therapy has been evidenced. This could be related to constitutive activation of downstream mediators or over-expression of other tyrosine kinase receptors [10]. For example, the persistent activation of downstream signalling steps such as MAPK and PI3K/Akt could promote cell proliferation, survival, differentiation and motility [11], as illustrated in a study in which the resistant phenotype unaffected by the treatment with C225 (monoclonal antibody anti-EGFR) is due to the intrinsic activity of those pathways [12].

Moreover, the increase of angiogenesis caused by up-regulation of the vascular endothelial growth factor in human cancer cells by EGF and TGF- α could promote resistance to EGFR inhibition [13].

To date, two anti-EGFR monoclonal antibodies, panitumumab and cetuximab, are currently in widespread use in cancer treatment.

Cetuximab

Cetuximab (C225, ErbituxTM) is an immunoglobulin (Ig) G1 human–murine chimeric counterpart of the murine

monoclonal antibody M225. It binds to the EGFR with a 2-log higher affinity compared with the natural ligands TGF- α and EGF [14]. Binding of cetuximab to the EGFR promotes receptor internalization and subsequent degradation without receptor phosphorylation and activation [15]. This results in receptor down-regulation, reducing the availability of EGFR on the cell surface and preventing activation of EGFR-associated, downstream signalling pathways. Cetuximab also binds to the mutant receptor EGFRvIII, inducing internalization of 50% of antibody-receptor complexes after 3 h, and an 80% reduction in phosphorylated EGFRvIII.

Binding of cetuximab to EGFR inhibits the progression of the cell cycle at the G0/G1 boundary, increases expression of the cell cycle regulator p27KIP1 and induces apoptosis by increasing expression of pro-apoptotic proteins (e.g. Bax and caspase-3, caspase-8 and caspase-9) [16] or by inactivation of anti-apoptotic proteins (e.g. Bcl-2) inducing decreased expression or phosphorylation [17]. Cetuximab has also been reported to inhibit the production of pro-angiogenic factors such as vascular endothelial growth factor, interleukin-8 and the basic fibroblast growth factor; inhibition of these factors is associated with a decrease in new blood vessel formation and the development of distant metastases in orthotopic cancer models [18].

Clinical overview of cetuximab

The first cetuximab Phases I and I/II trials demonstrated the safety of cetuximab alone or in combination with cytotoxic chemotherapy used as treatments for patients with metastatic squamous cell carcinoma of the head and neck, colorectal cancer and non-small-cell lung cancer [19–22].

A multi-centre, randomized Phase II trial examined the combination of cetuximab and irinotecan ($n = 218$) com-

Table 1. Up-front therapy with cetuximab and oxaliplatin or irinotecan (intention to treat population).

Authors	Patients (n)	Regimen	Response rate %	PFS (months)
Taberero 2007 [27]	43	Folfox-4	72	10.2
Bokemeyer 2007 [28]	337	Folfox-4 <i>versus</i> Folfox-4 + cmab	35.7 45.6	7.2
Borner 2006 [71]	74	Xelox <i>versus</i> Xelox + cmab	33 53	–
Folprecht 2006 [29]	21	Irinotecan/5FU/FA + cmab	67	9.9
Van Cutsem 2007 [31]	1217	Folfiri <i>versus</i> Folfiri + cmab	38.7 46.9	8 8.9
Taberero 2006 [30]	62	Folfiri + cmab every 2 weeks (escalating dose)	42	7.2

Cmab, cetuximab; PFS, progression-free survival; FU/FA, fluorouracil and leucovorin; Folfiri, fluorouracil, leucovorin and irinotecan; Folfox, oxaliplatin, leucovorin and fluorouracil.

pared to cetuximab alone ($n = 111$) in patients with EGFR-positive irinotecan-refractory metastatic colon cancer. The response rate (RR) of the combination was significantly higher, 23% *versus* 11% with cetuximab alone ($P = 0.007$), and the disease control rates were 56–32%, respectively. The time to progression was also significantly greater for the combination arm (4.1 *versus* 1.5 months, $P < 0.001$), and the median survival time was 8.6 months in the cetuximab arm ($P = 0.48$); survival in the two arms was not statistically significant. The presence of cutaneous rash correlated significantly with response, as the RR was higher in patients presenting rash compared with rash-free patients (25.8% *versus* 6.3%, $P = 0.005$) [23]. These findings were also confirmed by the MABEL (Monoclonal Antibody Erbitux in a European Pre-License study), which assessed definitively the efficacy and safety of cetuximab in the treatment of metastatic colorectal cancer (mCRC) [24].

Based on these results, cetuximab was approved for use in patients with EGFR-expressing mCRC refractory to irinotecan-based chemotherapy, in combination with irinotecan (for irinotecan-refractory patients) or as monotherapy (for irinotecan-intolerant patients).

The impact of cetuximab plus irinotecan in second-line metastatic colorectal EGFR-expressing cancer patients was examined in a multi-national Phase III trial known as the EPIC (Erbitux Plus Irinotecan for Metastatic Colorectal Cancer) study. The results reveal that treatment with cetuximab improved the progression-free survival (PFS), RR and health-related quality of life [25]. Moreover, cetuximab demonstrated activity in patients with colorectal cancer in whom other treatments have failed, improving the PFS, overall survival and quality of life over best supportive care (BSC) [26].

The role of cetuximab was also investigated in first-line treatment of mCRC. Phase II studies indicate that cetuximab combined with both irinotecan- and oxaliplatin-based chemotherapies is active with a 10–20% absolute increase in RR [27–30]. Recently, a multi-centre, randomized, Phase III study evaluated the combination of cetuximab with a standard chemotherapeutic regimen of fluorouracil, leucovorin and irinotecan (Folfiri) in previously untreated mCRC.

Cetuximab plus Folfiri increased RR significantly and prolonged PFS (Table 1) [31,32].

Cetuximab was also approved in 2006 for the treatment of head and neck cancer as a single agent in patients with advanced platinum resistant disease and in combination with radiotherapy for treatment of locally advanced disease.

A Phase III randomized trial of cetuximab in advanced head and neck cancer was carried out. In this study, a sample of 424 patients was divided into those treated with radiation therapy alone and those receiving supplementary treatment with cetuximab. At a median follow-up of 54 months, it was observed that the rate of survival was almost double in patients receiving cetuximab compared with patients receiving radiation therapy alone (49 *versus* 29 months; $P = 0.03$). Statistically significant increases in loco-regional control and PFS were also reported for the group receiving cetuximab. This is the first study to demonstrate a statistically significant survival benefit rate for patients treated with curative intent using an anti-EGFR antibody [33].

In addition, cetuximab plays a crucial role in the treatment of platinum-resistant squamous-cell carcinoma of head and neck cancer alone or in combination with chemotherapy (first-line setting) [34].

Panitumumab

Panitumumab (Vectibix, Amgen, Thousand Oaks, CA, USA) is a fully human IgG2 targeting the extracellular domains of EGFR monoclonal antibody. Developed by Abgenix's Xenomouse technology, which creates antibodies that do not contain murine proteins, it offers effective high affinity therapy ($K_d = 5 \times 10^{-11}$ M) with a minimum rate of allergic reactions or anaphylaxis [35].

Well tolerated – its main toxic effect is dermatological – it has never reached grade 4 in clinical trials. Because of its structure (fully human antibody), infusion-related reactions are minimal (only one of 148 patients experienced grade 3 reaction while one of 463 patients discontinued treatment due to grade 2 hypersensitivity reaction). Panitumumab could be administered weekly, fortnightly or every 3 weeks

without visible change in pharmacokinetic parameters [36].

Clinical overview of panitumumab

Panitumumab has been evaluated in clinical trials both as monotherapy and in combination with other agents for the treatment of various types, including colorectal and kidney cancer [37].

The Phase I study demonstrated that pharmacokinetic exposure showed similarities at 2.5 mg/kg per week, 6 mg/kg every 2 weeks and 9 mg/kg every 3 weeks. Grades 3- or 4-related adverse events were noted in 10% of patients, with grade 3 skin-related effects being the most frequent (7% of patients). No maximum tolerated dose was reached, and no human anti-human antibody formation or infusion-related reactions were observed [38].

In the Phase II studies, panitumumab in monotherapy and in combination with chemotherapy for the treatment of chemo-refractory colorectal cancer was active and well tolerated. A relationship between skin rash severity and survival was noted as being similar to that observed with the use of cetuximab. Moreover, its efficacy appears to be similar in patients with both low and negative EGFR levels [39–42].

A large Phase III multi-centre pivotal trial randomized patients with oxaliplatin and irinotecan-refractory EGFR-expressing mCRC between BSC and BSC plus panitumumab at a dose of 6 mg/m² every 2 weeks. The aim of this study was to show the significant difference in PFS. Panitumumab showed a 46% decrease in tumour progression rate compared with that of BSC alone (hazard ratio: 0.54; 95% confidence interval: 0.44, 0.66; $P < 0.00000001$, stratified log-rank test). The subset analyses demonstrated the consistent treatment effect of panitumumab in all subgroups of patients. The RR was significantly higher in the panitumumab arm, with an 8% partial RR and 28% achieving stable disease, compared with no partial responses and 10% stable disease with BSC alone. The time to response was 8 weeks, and the median duration of the response was 17 weeks. Panitumumab also showed activity in cross-over study patients, with 10% achieving partial response and 32% stable disease. No difference was observed in overall survival (hazard ratio: 1.00; 95% confidence interval: 0.82–1.22) due probably to the high rate of cross-over population to panitumumab.

Skin reactions of any grade occurred in 90% of patients receiving panitumumab, and in 9% of those receiving BSC; the incidence of grades 3–4 skin-related adverse reactions was 14% compared with 0% in patients not treated with panitumumab. The incidence of skin toxicity in the panitumumab group was dose-related; however, no correlation was observed between dose and severity [43].

A further analysis of biomarkers was conducted to determine whether the effect of panitumumab monotherapy on PFS differed in patients with tumours characterized by mutant compared with wild-type (WT) K-ras.

Amado *et al.* first demonstrated that the response to panitumumab monotherapy and the improvement in PFS was limited only to patients with WT K-ras. No patient harbouring a K-ras mutation (46%) responded to panitumumab [44].

These findings led to registration by regulatory agencies worldwide as a monotherapy for third-line treatment of colorectal cancer that is refractory to fluoropyrimidines, oxaliplatin or irinotecan. Moreover, in 2007, panitumumab was approved by the European Medicines Agency for use in patients with colorectal cancer carrying a normal, WT K-ras gene.

The role of panitumumab in combination with anti-angiogenic drugs has also been explored in a randomized Phase III study (panitumumab advanced colorectal cancer evaluation). In this trial patients with mCRC were assigned randomly for first-line treatment within each chemotherapy cohort (823 patients oxaliplatin- and 230 irinotecan-based) to bevacizumab and chemotherapy with or without panitumumab 6 mg/kg every 2 weeks. The primary end-point was PFS within the oxaliplatin cohort. The results of the study were negative, as the combination of panitumumab with bevacizumab and chemotherapy resulted in a decrease of PFS and in excessive toxicity, particularly diarrhoea, infections and pulmonary embolism. The results were consistent in both the oxaliplatin and irinotecan cohorts. Moreover, as demonstrated previously, the triple combination did not provide additional benefit in the K-ras WT population treated with panitumumab [45].

Predictive biomarkers

The early biomarker, developed in mCRC to correlate with the activity of anti-EGFR antibodies, was confined to merely expressing the target.

Chung *et al.* demonstrated that colorectal cancer patients with EGFR-negative tumours have the potential to respond to cetuximab-based therapies, registering a 25% objective RR. Consequently, the presence of the target (EGFR) does not ensure the response to anti-EGFR inhibitors. Furthermore, EGFR analysis by current immunohistochemistry techniques does not seem to have predictive value for the selection or the exclusion of patients for cetuximab, therefore EGFR immunohistochemistry is not warranted currently [46].

In addition, not even the gene copy number for EGFR determined by fluorescence *in situ* hybridization on tumour samples was associated significantly with clinical response to this targeted therapy [47].

Several studies have been carried out to define a subgroup of patients with potentially differential responses to anti-EGFR antibody therapy, and these show that benefits are confined to the subgroup with WT K-ras tumours.

K-ras is a guanosine triphosphate-binding protein with a critical role in cellular growth and survival pathways [48]. It

Table 2. Phase III study of panitumumab *versus* best supportive care (BSC).

	Panitumumab (<i>n</i> = 231)	ITT population BSC (<i>n</i> = 232)	K-ras WT panitumumab (<i>n</i> = 124)	BSC (<i>n</i> = 119)
RR, <i>n</i> (%)	22 (10)	0 (0)	21 (17)	0 (0)
mPFS (m)	1.9	1.7	2.9	1.7
PFS HR	0.54 (0.44–0.66)		0.45 (0.34–0.59)	

Amado *et al.* 2008 [44]

ITT, intention to treat; RR, response rate; PFS, progression-free survival; HR, hazard ratio; WT, wild-type.

is mutated in approximately 30–50% of colorectal cancer and results in a constitutive activation of the MAPK pathway and in lack of response with EGFR inhibitors [49–53].

Published reports so far have investigated the role of K-ras as a selection marker for EGFR inhibitor treatment on tumour samples from uncontrolled studies and include therapy with EGFR inhibitors alone or in combination with chemotherapy. In this respect, the relative effect with antibody treatment on the outcome of K-ras, as a predictive marker, is not so clear. In, the study published by Amado *et al.*, comparing panitumumab monotherapy with BSC, no clinical benefit to panitumumab at all in patients with the K-ras mutation was evidenced in any clinical end-point, thus confirming the role of the K-ras mutant as a negative predictor of response (Table 2) [44].

Analyses of K-ras status and response to cetuximab have provided similar results.

The retrospective correlative analysis of patients enrolled in the CO.17 trial performed by Karapetis *et al.* show that the benefit of cetuximab treatment was confined to patients who had a tumour with no K-ras mutations, with few or no effects in the presence of a K-ras mutation (Table 3) [54].

Moreover, in patients with mCRC treated with first-line infused fluorouracil, folinic acid and oxaliplatin with or without cetuximab the improved RR and PFS associated with cetuximab was confined to those patients having a K-ras WT tumour [55–57].

Recently, another breakthrough has been achieved focusing on the role of v-raf murine sarcoma viral oncogene homologue B1 (BRAF) as a potential biomarker for anti-EGFR antibody treatment. Di Nicolantonio *et al.* showed that in the presence of BRAF mutations (BRAF V600E allele) there was no response to either cetuximab or panitumumab [58].

Immunological mechanisms

In recent years, it has been shown that the anti-tumoral effects of mAb may be due to their ability to act on the immune system [59]. In general terms, the use of mouse chimeric antibodies may elicit immune responses specifically for the mouse portion of the molecule leading to a destruction of the antibody; in some cases, however, also to the destruction of targeted tumoral cells [60]. In addition, several reports have described, both *in vitro* and *in vivo*, how these antibodies are able to elicit antibody-dependent cellular cytotoxicity (ADCC), complement-mediated cytotoxicity, or both. These effector responses are due to the binding of the Fc portion of antibodies to the Fc receptors expressed on the surface of different cell types. This binding leads to a wide array of effects, from uptake to killing. It should be noted that macrophages, dendritic cells, neutrophils, eosinophils, B cells, mast cells, natural killer (NK) cells, platelets and Langerhans cells express Fc receptors capable of discriminating different Ig classes. The effects of ligation of the Fc portion of the antibody with the Fc receptors on the cells depend upon the specificity of the Fc receptors for a given Ig class and on the cell types. For instance, cetuximab is an IgG1, therefore it is able to bind several Fc receptors: FcγRI (CD64), FcγRII-A (CD32), FcγRII-B1 (CD32), FcγRII-B2 (CD32) and FcγRIII (CD16) [61]. For this reason, cetuximab is able to mediate ADCC induced by NK activity through binding to FcγRIII, but it is also able to engage Fc receptors on the surface of other cells such as eosinophils, mast cells, dendritic cells, B cells and other cell types [62]. Therefore, an obvious scenario can be envisaged in which the overall effects of these antibodies are also due to complex mechanisms other than those of ADCC and complement-mediated cytotoxicity. Recently, another intriguing mechanism has

Table 3. Phase III study of cetuximab *versus* best supportive care (BSC).

	Cetuximab (<i>n</i> = 287)	ITT population BSC (<i>n</i> = 285)	K-ras WT cetuximab (<i>n</i> = 124)	BSC (<i>n</i> = 119)
RR, <i>n</i> (%)	–	–	13 (13)	0 (0)
mPFS (m)	1.9	1.8	3.8	1.9
PFS HR		0.68 (0.57–0.80)	0.40 (0.30–0.54)	
mOS (m)	6.1	4.6	9.5	4.8
OS HR		0.77 (0.64–0.92)	0.55 (0.41–0.74)	

Karapetis *et al.* 2008 [54]

ITT, intention to treat; PFS, progression-free survival; RR, response rate; HR, hazard ratio; OS, overall survival; WT, wild-type.

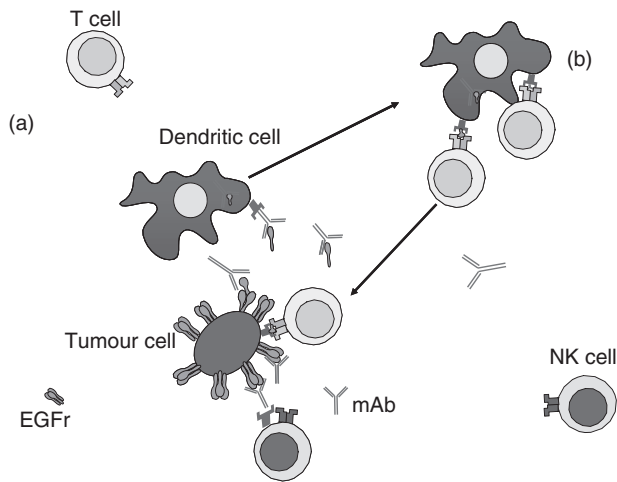


Fig. 2. Effects of monoclonal antibody (mAb) therapies on the immune system. (a) The mAb bind the epidermal growth factor receptor (EGFR) expressed on the surface of the tumour cell; this binding activates antibody dependent cellular cytotoxicity by engaging Fc receptors on the surface of natural killer (NK) cells. Some T cells may also be activated by specific recognition of the Fc portion of the mAb. Finally, the binding mAb/EGFR can allow, by trogocytosis, the internalization of part of EGFR by antigen-presenting cells as the dendritic cells. (b) Dendritic cells present peptides derived from the internalized portion of EGFR to T cells, priming an immune response more efficient in recognition of EGFR peptides associated with major histocompatibility complex expressed by tumour cells. As reported in the text, Fc receptors are expressed on the surface of many cells (mast cells, eosinophils, etc.) that can secrete a variety of different cytokines and chemokines, thus amplifying the overall immune response induced by mAb.

been highlighted that can help to exploit the potential effect of mAb in solid cancer. Upon recognition by antibodies of ligands on target cells, the components of this immune complex (antibody and ligand) can be removed (shaved) from these cells and internalized by cells expressing Fc γ receptors [63–66].

The mechanism has been called trogocytosis, or nibbling. Trogocytosis triggers a cascade of complex events, depending upon the ligand size and IgG class involved, which affect the immune response in different ways, sometimes even depressing the immune response or the efficacy of the binding antibody/target due to antigen shedding in blood [67]. It has also been shown to be effective when IgG2 antibodies, such as panitumumab, are used (Fig. 2) [68].

Finally, it should be remembered that the ability to interfere with the signalling initiated by EGFR affects many other receptor systems, for instance chemokine and cytokine receptors Toll-like receptors, that are critical for immune responses [69]. Another interesting perspective worthy of note is that activation of the immunological mechanisms described above corresponds with the observation that increased efficacy in patients treated with EGFR mAb is correlated with the presence of cutaneous rash. Indeed, the links

between complement activation, ADCC and cutaneous rash are well known to clinical immunologists.

On the basis of such premises, it can be suggested that the beneficial effects of anti-EGFR antibodies do not depend only upon their ability to interfere with the signalling generated by the receptor and this could explain, at least in part, the different response observed in treated patients. Furthermore, a better understanding of the interactions between the mAb used as therapy for solid tumours and the immune system could be critical in designing new approaches for immunotherapy in cancer.

Conclusion

The use of anti-EGFR monoclonal antibodies is diffuse in cancer therapy; however, clinical responses have been observed in only 15% of patients treated. Moreover, the use of these drugs has contributed only a modest overall survival benefit in comparison to commonly practised BSC. Although these results could be considered interesting, it should be taken into account that in several countries the analysis of cost-effectiveness is now proportionally extremely limiting, given the economic crisis worldwide and its effects on the budgets of national health systems. As a result, these data would not be sufficient to influence or avoid a progressive reduction in the use of mAb therapy in clinical practice.

The challenge for the near future is to identify biomarkers that are capable of predicting and targeting eligible patients for anti-EGFR antibody treatment on a one-to-one scale. On the basis of reported trials and current debate [70], complex efforts are needed that take into consideration not only the mechanistic effect of blocking EGFR, but also the further exploiting of mAb effects on the immune system. Clearly, this is no simple task and, although there is now an impressive quantity of data in the field, we are still far from having a thorough understanding of the scenario as a whole. In this respect, the consideration that EGFR-related signalling is involved in many steps of immune responses provides an inkling of just how complex is the road ahead. However, in the long term the final destination will be well worth all our painstaking and strenuous efforts.

Disclosure

The authors declare no conflict of interest.

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