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## Afatinib efficacy against squamous cell carcinoma of the head and neck cell lines in vitro and in vivo

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## Abstract

Epidermal growth factor receptor (EGFR) inhibitors have demonstrated efficacy in squamous cell carcinoma of the head and neck (SCCHN). In addition to EGFR, other ErbB family members are expressed and activated in SCCHN. Afatinib is an ErbB family blocker that has been approved for treating patients with EGFR-mutated nonsmall cell lung cancer. We sought to determine the efficacy of afatinib in preclinical models and compare this to other EGFR-targeted agents. Afatinib efficacy was characterized in a panel of ten SCCHN cell lines and found to be most effective against cell lines amplified for *EGFR*. Afatinib had lower IC<sub>50</sub> values than did gefitinib against the same panel. Two *EGFR*-amplified cell lines that are resistant to gefitinib are sensitive to afatinib. Cetuximab was not found to have a synergistic effect with afatinib either in vitro or in vivo. Both afatinib and cetuximab were effective in tumor xenograft model. Afatinib is an effective agent in SCCHN especially in models with *EGFR* amplification.

## Keywords

Squamous cell carcinoma of the head and neck; Afatinib; Gefitinib; Cetuximab; Epidermal growth factor receptor

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## Introduction

Epidermal growth factor receptor (EGFR) and HER2 are both members of the ErbB tyrosine kinase receptor family which also comprises HER3 and HER4. When stimulated by the presence of a ligand, ErbB receptors homodimerize or heterodimerize with other members of the family, and through transphosphorylation, initiate intracellular signaling cascades [1]. Although HER2 has no known ligand, it is believed to be a preferred dimerization partner for EGFR and HER3 [2]. As members of the ErbB family are often deregulated in a number of malignancies [3], they have been an attractive option for targeted therapy. EGFR is thought to be highly expressed in as many as 90 % of the head and neck squamous cell carcinoma (SCCHN) tumors [4], while the expression of HER2 is more variable [5, 6].

One of the earliest agents to target the ErbB network system was cetuximab, a chimeric monoclonal antibody to EGFR. After demonstrating efficacy in preclinical models [7], cetuximab earned FDA approval for locally advanced as well as recurrent or metastatic SCCHN [8–11]. Another potential therapeutic target is the intracellular tyrosine kinase domain of ErbB receptors. First generation small molecule tyrosine kinase inhibitors (TKIs), such as gefitinib and erlotinib, specifically target the tyrosine kinase of EGFR and are approved for clinical use in lung cancer. Gefitinib demonstrated excellent activity in preclinical SCCHN models [12, 13]. In phase 2 trials, gefitinib had a response rate of 9–10 % in patients with recurrent or metastatic SCCHN [14]. Erlotinib had similarly promising preclinical results [15] and slightly worse response rates in clinical studies [16].

Afatinib is an irreversible TKI that targets all kinase competent ErbB family members (EGFR, HER2, and HER4; HER3 lacks essential catalytic residues and displays weak to no activity) [17]. Early in vitro studies of afatinib in human tumor cell lines demonstrated significant activity as well as an increase in the proportion of cells in the sub-G0/G1 phase

of the cell cycle [18, 19]. Effective antitumor activity of afatinib has also been shown in multiple xenograft models using human cell line-derived xenografts and transgenic mouse models [19, 20]. Phase I trials in patients with nonsmall cell lung cancer (NSCLC) and other solid tumors found afatinib to be well tolerated, with the most common adverse effects being diarrhea, rash, fatigue, and nausea [21, 22]. Although not designed to determine clinical response, there was a decrease in tumor size in 50 % of patients [22]. At this point, besides early evaluations assessing afatinib in combination with radiotherapy in one single model system (FaDu) [23], no work characterizing the effect of afatinib in a panel of SCCHN models has been reported.

## Materials and methods

### Viability assays

Cell Titer Blue assays (Promega, Madison, WI) were used to determine cell viability. Briefly, cells were seeded at 1000 cells/well on 96-well plates, incubated for 24 h in complete media, maintained in serum-free conditions for 24 h, and treated with afatinib or cetuximab for 72 h at 37 °C. Then, Cell Titer Blue reagent was added to each well and incubated for 2 h at 37 °C before 3 % SDS was added. Cell viability was quantified by scanning absorbance at 570 nm in a microplate reader (Bio-Tek, Winooski, VT).

### Tumor xenografts treated with afatinib only or multiple ErbB-targeting agents

Mice were allowed to adjust to conditions at least for 5 days before they were used for experiments. They were housed in Macrolon type III cages in groups of ten under standardized specific pathogen-free (SPF) conditions at  $21.5 \pm 1.5$  °C temperature and  $55 \pm 10$  % humidity. Standardized diet (PROVIMI KLIBA) and autoclaved tap water were provided ad libitum. HN5 or FaDu cells were injected into the right flank of 6-week-old female BomTac:NMRI-Foxn1<sup>nu</sup> mice (Taconic, Denmark). Tumor take was monitored over time, and animals with established tumors ( $50\text{--}100$  mm<sup>3</sup>) were randomized to the treatment groups ( $n=10$ /group). Mice were treated with either vehicle or afatinib, daily or every other day or in a weekly alternating schedule (afatinib only experiment) or with vehicle or other agents daily (multiple agent experiment). Tumor volumes and body weights were recorded three times a week, and median tumor volumes as well as change in body weight were plotted over time.

### Tumor xenografts treated with afatinib and cetuximab

HN5 and SCC25 cell line xenograft experiments were conducted as previously described [24]. Briefly, female athymic nude mice (Harlan, Indianapolis, IN) were maintained in a pathogen-free animal facility in accordance with the University of Chicago Animal Care and Use Committee. Mice received standard laboratory rodent food and water as desired. All handling procedures were conducted in a laminar flow biosafety hood. At 6–7 weeks of age, mice were injected subcutaneously in the right flank with HN5 or SCC35 cells. Drug treatment was initiated when mean tumor volumes reached 200 to 250 mm<sup>3</sup>. Mice were treated with cetuximab at 30-mg/kg body weight two times per week, via i.p. injection and/or afatinib at 12.5-mg/kg body weight once per day via oral gavage, or equal volume of diluents (control). For tumor growth analysis, tumor size was measured with a vernier

caliper. Tumor volumes were calculated with the formula  $V=0.52 \times L \times W^2$ , where L and W represent the length and the width of the tumor (mm). The animals were monitored two times per week for body weight and tumor volume.

### Quantification of TGFA, EGFR, and AREG mRNA expression and EGFR copy number by real-time PCR

The real-time PCR method for quantification of *EGFR* gene expression and copy number has been reported previously [25, 26].

### FISH assays

FISH assay methods and analysis were previously described [24]. Briefly, for the evaluation of the *EGFR* gene copy number (GCN) alterations, dual-color FISH assays were conducted using an *LSIEGFR SpectrumOrange:CEP7SpectrumGreen* Probe mixture (Vysis/Abbott Molecular, Des Plaines, IL). *HER2* amplification was studied using the Vysis PathVysion *HER2* DNA Probe Kit according to manufacturer recommendations (Abbott Molecular, Des Plaines, IL). *CEP7* or *CEP17* probes were used to distinguish true gene amplification from *EGFR* or *HER2* gene copy number gain (gene polysomy) and alterations in number of chromosome 7 or 17 homologs. The absolute number of each signal, the mean copy number of signal per cell, the ratios of *EGFR* to *CEP7* or *HER2* to *CEP17*, and the percentage of cells with given copy number of each signal per cell were calculated. Cells with a gene to chromosome signal ratio  $<2$  were considered nonamplified, whereas those with a ratio greater than 2.0 (or 15 copies of *EGFR* per cells in 10 % of cells) were considered as having true amplification. Cells with ratios near cutoff points were equivocal or low amplified.

### Western blotting

Western blots on cell lysates were performed as previously described [24]. Visualization and quantification were performed with Odyssey Infrared Imaging System (Li-Cor Biosciences). Experiments were repeated at least three times. PTEN antibody was purchased from Cell Signaling Technology, Inc. (Danvers, MA). Actin antibodies were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). Secondary goat anti-rabbit IgG IRDye antibody was purchased from LI-COR Biosciences (Lincoln, NE). Secondary mouse IgM IRDye antibody was purchased from Rockland Immunochemicals Inc. (Gilbertsville, PA).

## Results

### Sensitivity to afatinib

Cell viability of ten SCCHN cell lines grown in monolayer cultures was determined over a range of afatinib concentrations (Fig. 1) and compared to  $IC_{50}$  ranges of the same cell lines to gefitinib (Table 1). In order to assess whether anti-proliferative activity could be improved upon with the addition of cetuximab, viability of cell lines with high  $IC_{50}$  values (SCC35 and Detroit 562) was tested at several doses (Fig. 2a, b). Treatment with cetuximab alone had little effect on cell viability, even at relatively high concentrations (100 nmol/L). The combination treatment resulted in CI values above 1, thus demonstrating no evidence of a synergistic or additive effect (data not shown). Treatment with afatinib and cetuximab was

tried on additional cell lines with greater sensitivity to afatinib (SQ20B and SCC61) with similar results (Fig. 2c, d).

### In vivo tumor xenografts

In vivo activity of afatinib was first characterized against FaDu cells injected into the right flank of BomTac:NMRI-Foxn1<sup>nu</sup> mice. Treatment with afatinib at 10 mg/kg followed two regimens. Both daily and intermittent treatment regimens slowed tumor growth. Continuous dosing resulted in virtually no tumor growth, while intermittent dosing saw increase in tumor volume over time (Fig. 3a). Neither treatment regimen at the doses used had a significant effect on body weight (Fig. 3b).

The anti-tumor activity of afatinib and several other ErbB targeting agents was further tested in mice bearing HN5 xenografts (Fig. 4). Daily treatment at 15 mg/kg resulted in tumor regressions in all treated animals. Four complete responses and six responses displaying 75 % or more reduction in tumor size compared to baseline were observed. Erlotinib, at 75 mg/kg, was less potent and induced six regressions and four tumors grew under treatment. Neratinib and lapatinib induced tumor growth delay, but no regressions were achieved.

The last set of in vivo experiments assessed the efficacy of afatinib in combination with cetuximab. Afatinib (12.5 mg/kg) and cetuximab (30 mg/kg), separately and in combination, were very effective at inhibiting in vivo tumor growth in both the HN5 and SCC35 mouse xenograft models (Fig. 5). At the doses tested, the combination treatment offered no advantage to single agent administration.

### PTEN expression

PTEN protein expression was examined by immunoblotting of SCCHN cell line lysates. We found that all lines express the protein at approximately equal levels (Fig. 6), eliminating PTEN loss as an explanation of primary EGFR inhibitor resistance in these lines.

### EGFR and HER2 amplification

Four of our SCCHN cell lines were *EGFR* amplified by FISH (Table 2, Fig. 7a and [26]). SCC58, HN5, and SQ20B exhibit high amplification (*EGFR/CEP7* ratio >7), and SCC25 exhibits low amplification (*EGFR/CEP7* ratio ~2). These same four lines show a gain of *EGFR* mRNA copies normalized to 18 s mRNA while the remaining cell lines do not (Table 2 and [26]). SCC28 cells do not show *EGFR* amplification (Table 2) but have high *EGFR* gene polysomy (Fig. 7b).

SCC25, HN5, SCC58, SQ20B, SCC61, and SCC28 were also tested for *HER2/CEP17 FISH*. No *HER2* amplification was found (Table 2 and Fig. 7c–h). SCC25, HN5, and SCC28 cells carried in average three copies of *HER2* per cell due to trisomy for chromosome 17.

### Discussion

In vitro, SCCHN cell lines show a range of sensitivities to afatinib. The four most sensitive cell lines, SCC58, SQ20B, SCC25, and HN5, show amplification of *EGFR* by FISH analysis and increased mRNA copy number by qPCR. This suggests that afatinib is most effective in

cell lines where EGFR is amplified and possibly acts as a driver of cell growth. EGFR gene copy numbers have not been correlated with clinical activity of EGFR inhibitors in SCCHN; however, this has not been tested in a prospective study. Our data presented here and in the past [26] indicate that a prospective trial is warranted. When afatinib IC<sub>50</sub> values are compared to those from gefitinib (Table 1), one sees that the order of increasing resistance is almost identical, the exceptions being SCC25 and SCC58, both of which are sensitive to afatinib but resistant to gefitinib. This suggests that afatinib may be a better therapeutic choice for cancers expressing high levels of EGFR and may relate to the broader ErbB inhibitory scope of afatinib as compared to gefitinib. Because gefitinib inhibits EGFR only, while afatinib targets EGFR, HER2, and HER4, one possibility could reside in differences due to HER2 signaling. However, none of the four most sensitive lines show an increased gene copy number for HER2 by FISH ruling out amplification as a possible mechanism. HER3 has been implicated as a mechanism of resistance to EGFR blockade in SCCHN in one paper [27]. As afatinib has been shown to block transphosphorylation of HER3 in vitro [19], it may be able to address this potential resistance mechanism. Indeed, afatinib activity was seen in cetuximab refractory patients in a phase II trial [28] suggesting a lack of cross-resistance in some instances.

PTEN expression was also not at play in afatinib resistance, as expression levels were similar in all cell lines. The main factors which could explain the differences in efficacy are the irreversible binding mode and the anticipated tighter blockade of the ErbB signaling network by afatinib. However, a measure of caution should be used in interpreting in vitro data, as SCC35, which exhibited the highest IC<sub>50</sub> to afatinib (Table 1), was still sensitive to the drug in vivo (Fig. 5).

Gefitinib does block phosphorylation of AKT and ERK in SCC25 and SCC58 [26, data not shown], possibly pointing to signaling through STAT3 downstream. Elucidation of this mechanism could point to a more specific subset of cases in which afatinib is more effective than gefitinib. De Greve et al. [29] showed an objective response to afatinib in three patients with mutations in the tyrosine kinase domain of HER2 even after other ErbB targeting treatments had failed. Afatinib may be active in the presence of multiple genetic aberrations, which render other TKIs ineffective.

Indeed, afatinib was more effective than lapatinib, erlotinib, and neratinib in the HN5 tumor xenograft experiment. These three TKIs were only tested on one cell line, limiting broad conclusions about mechanism. However the combination of irreversible binding mode and broad ErbB signaling blockage is provided by afatinib alone among the TKIs tested here.

With these experiments, we were unable to show added benefit for combination therapy with cetuximab in the most resistant cell lines to afatinib. Afatinib and cetuximab were very effective at arresting growth and decreasing tumor volume in vivo, both as single agents and in combination. In fact, these treatments worked so well that any benefit from the combination of drugs could not be seen.

## Conclusion

Afatinib is most effective at arresting growth of SCCHN cell lines in vitro that are amplified and/or overexpress EGFR. Even EGFR-amplified cell lines which are resistant to gefitinib respond well to afatinib treatment. Afatinib was well tolerated in mice at the doses tested and was more effective with continuous dosing. Afatinib was more effective than other TKIs at arresting tumor xenograft growth, perhaps due to a combination of irreversible binding and broad inhibition of ErbB network signaling. Synergy between afatinib and cetuximab could not be observed in vitro due to the poor response to cetuximab and in vivo due to the excellent response to both drugs.

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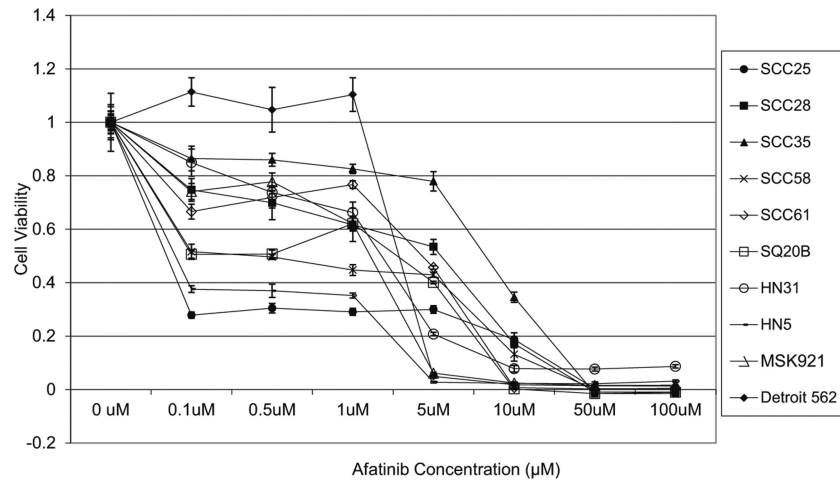
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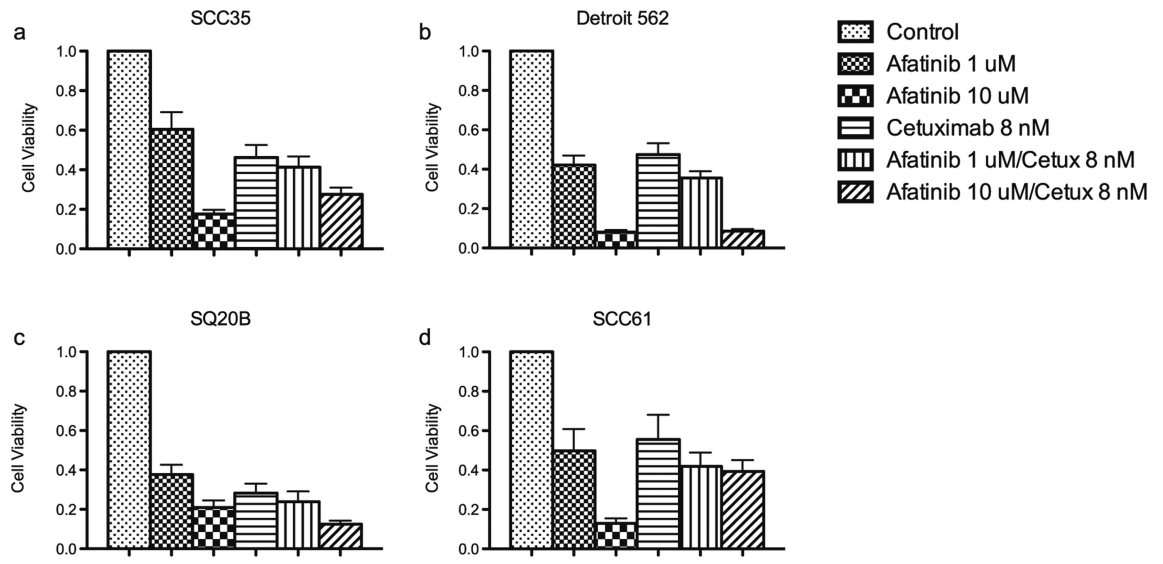
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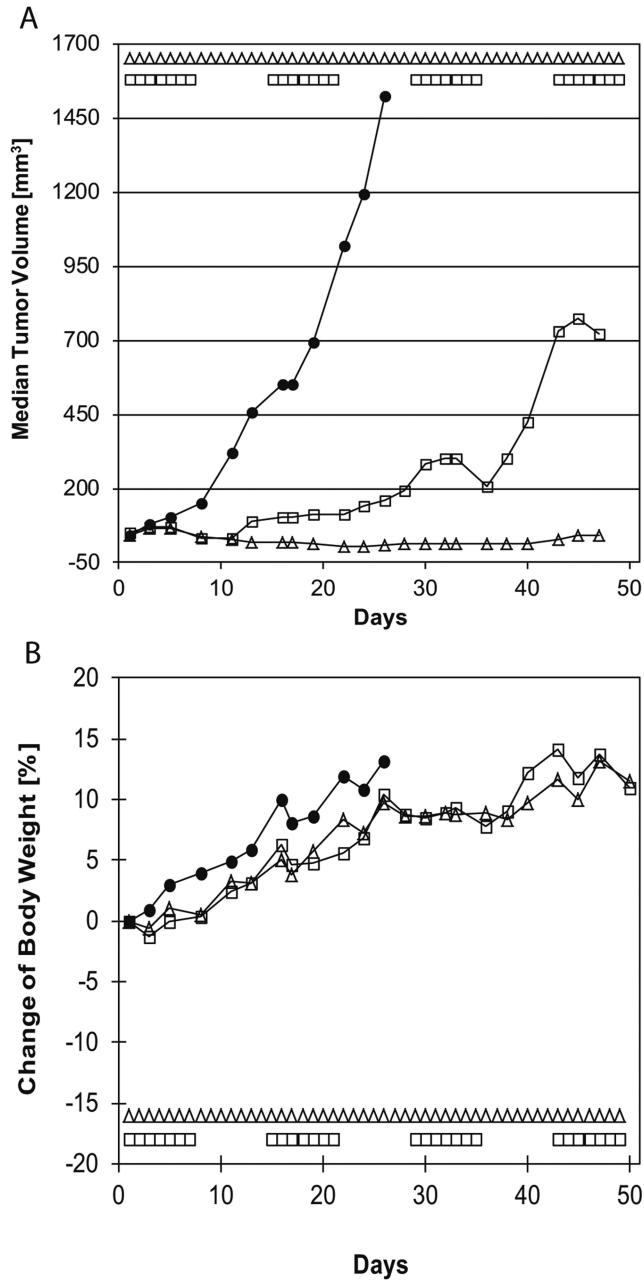
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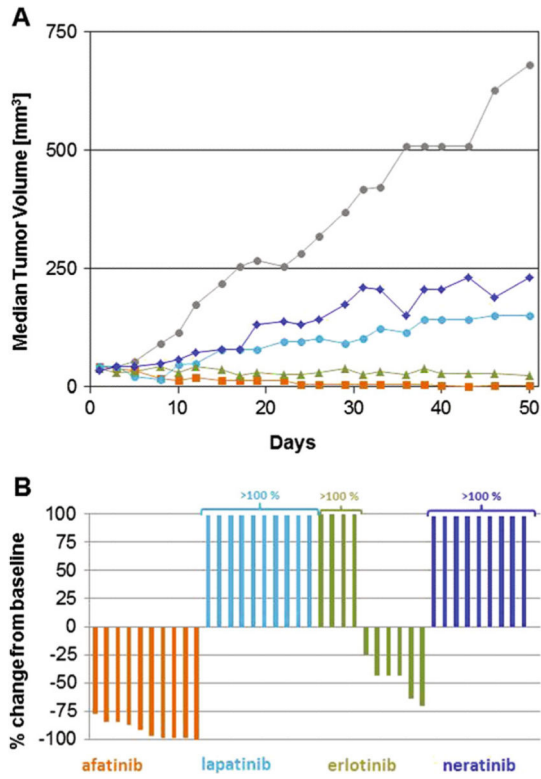
**Fig. 1.** Viability of ten SCCHN cell lines treated with a range of concentrations of afatinib. Results from Cell Titer Blue assays



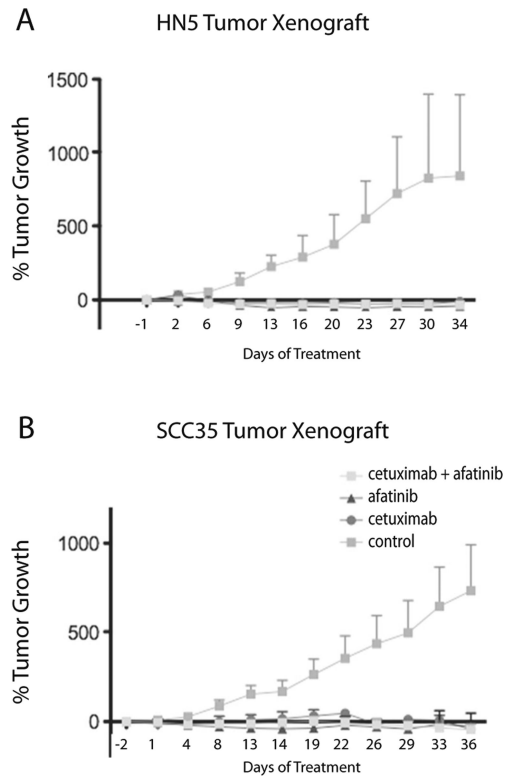
**Fig. 2.** Viability of SCCHN cell lines treated with afatinib and cetuximab. Results from Cell Titer Blue assays. **a** SCC35, **b** Detroit 562, **c** SQ20B, **d** SCC61



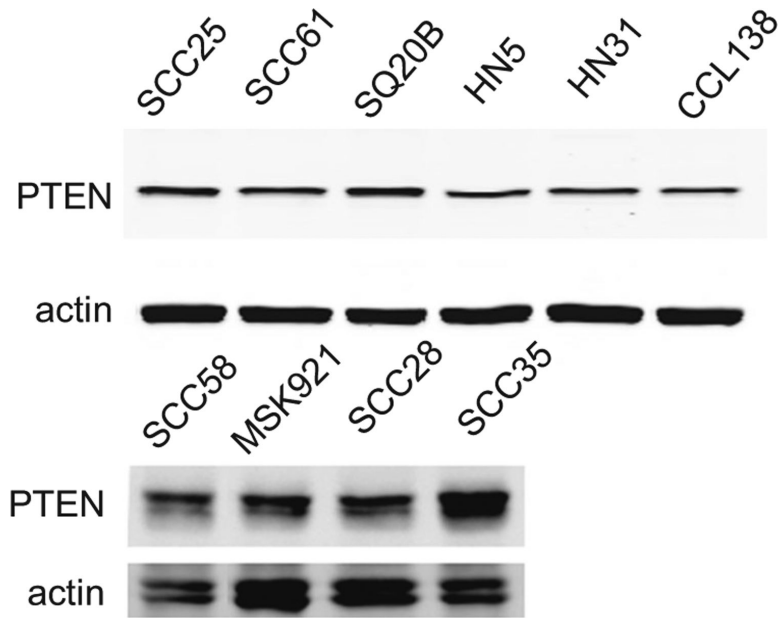
**Fig. 3.** In vivo activity of afatinib in FaDu cell line-derived xenografts. Mice were treated daily with either vehicle (*black circle*) or afatinib as described in (*white triangle*) or in a weekly alternating schedule (*white square*). Tumor volumes and body weights were recorded three times a week, and median tumor volumes (**a**) as well as change in body weight (**b**) were plotted over time



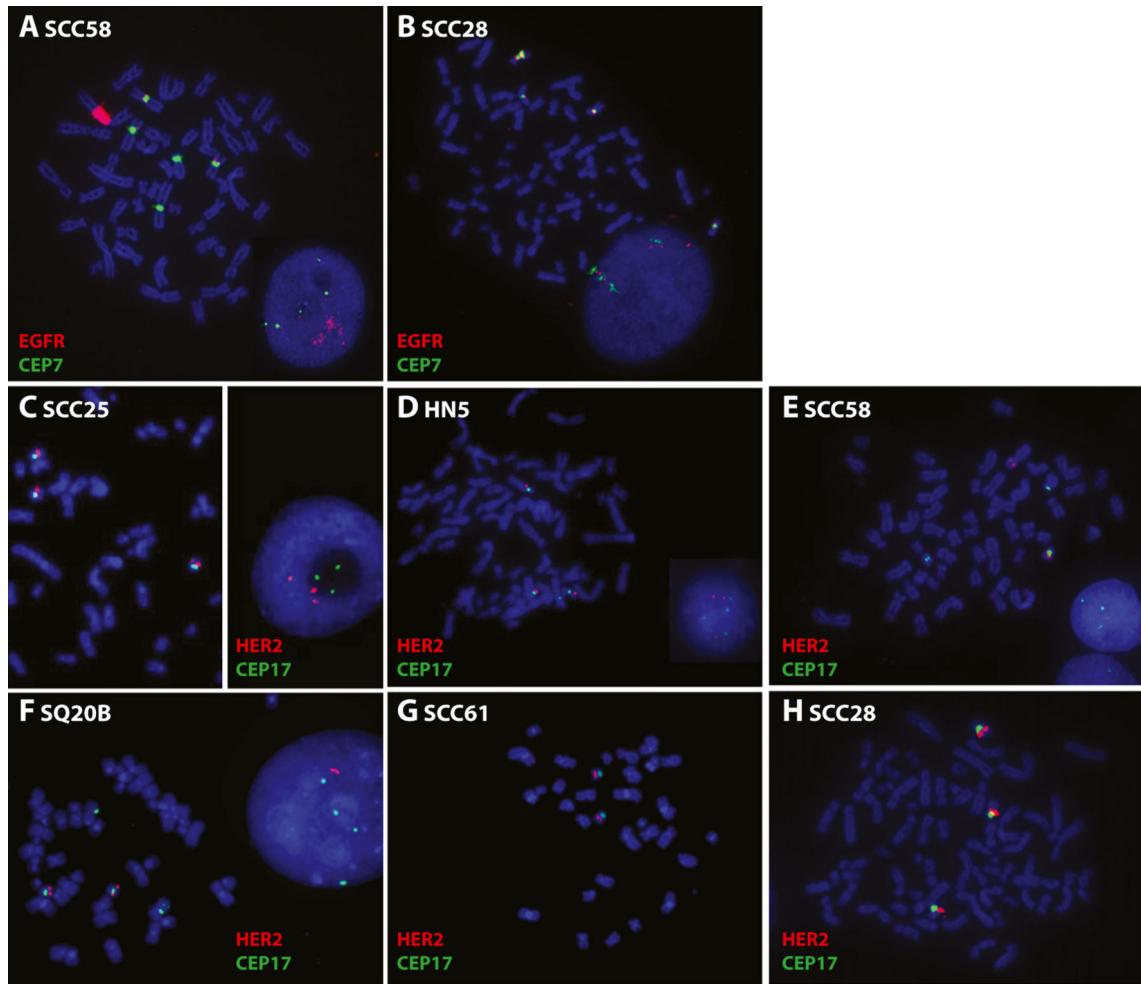
**Fig. 4.** In vivo activity of afatinib in HN5 cell line-derived xenografts. Mice were treated daily with either vehicle (*gray*), 2×100 mg/kg/day lapatinib (*turquoise*), 75 mg/kg/day erlotinib (*green*), 50 mg/kg/day neratinib (*blue*), or 15 mg/kg/day afatinib (*orange*) as described in “Materials and methods.” Tumor volumes were recorded three times a week. Median tumor volumes were plotted over time (**a**). On day 50, relative tumor volume changes were recorded **b**



**Fig. 5.** SCCHN cell line tumor xenografts respond to treatment by afatinib alone, or in combination with cetuximab. % tumor growth is shown. **a** HN5, **b** SCC35



**Fig. 6.** PTEN expression in SCCHN cell lines. Western blotting shows that PTEN expression is consistent across cell lines. Actin serves as loading control



**Fig. 7.** *EGFR* and *HER2* FISH. Images of metaphase and interphase nuclei after FISH are presented. The *EGFR* and *HER2* genes are localized by red fluorescent signals, and chromosome 7 and 17 centromeres (*CEP7* and *CEP17*) are localized by green fluorescent signals. The cells were counterstained with DAPI (blue). *EGFR/CEP7* staining of **a** SCC58 and **b** SCC28 cells. *HER2/CEP17* staining of **c** SCC25, **d** HN5, **e** SCC58, **f** SQ20B, **g** SCC61, and **h** SCC28



**Table 1**Afatinib IC<sub>50</sub>s compared to gefitinib IC<sub>50</sub>s of SCCHN cell lines

Cell line	Afatinib IC <sub>50</sub> (μM)	Confidence interval	Gefitinib IC <sub>50</sub> (μM)
SCC25	0.0102	9.796e-005 to 1.062	5-10
HN5	0.06412	0.009348 to 0.4398	0.05-0.5
SCC58	0.3405	0.07144 to 1.623	>100
SQ20B	0.5176	0.08746 to 3.064	1-5
MSK921	1.174	0.6654 to 2.071	1-5
HN31	1.225	0.8320 to 1.805	1-5
SCC61	1.903	0.6092 to 5.942	5
SCC28	1.951	0.7886 to 4.827	>5
Detroit 562	2.485	0.9399 to 6.569	5-50
SCC35	8.008	5.645 to 11.36	100

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**Table 2***EGFR* and *HER2* gene copy number alterations and mRNA expression levels of SCCHN cell lines

Cell line	FISH				qPCR			
	<i>EGFR</i> GCN <sup>1</sup>	<i>CEP7</i> GCN <sup>2</sup>	<i>EGFR/CEP7</i> <sup>3</sup>	Interpretation	TGFa/18 s	AREG/18 s	<i>EGFR</i> /18 s	Interpretation
SCC58	32.94	4.3	7.7	High Ampl	2.02	9.67	17.08	Gain
SCC28	5.02	5.33	0.94	No Ampl,	0.39	1.86	1.77	Normal
	<i>HER2</i> GCN <sup>1</sup>	<i>CEP17</i> GCN <sup>2</sup>	<i>HER2/CEP17</i> <sup>3</sup>	Interpretation				
SCC25	3.25	3.05	1.07	No Ampl,				
HN5	2.88	3.9	0.74	No Ampl,				
SCC58	1.87	2.87	0.65	No Ampl				
SQ20B	2.06	4.20	0.49	No Ampl				
SCC61	2.3	2.04	1.13	No Ampl				
SCC28	2.85	2.93	0.97	No Ampl,				

FISH:

qPCR: Mean values from three experiments were calculated relative to 18 s as control, arbitrarily set at 1. Gain, &gt; 7 copies of gene/18 s

<sup>1</sup> Mean copy number *EGFR* or *HER2* per cell<sup>2</sup> mean copy number of centromere enumeration probe (*CEP7* or *CEP17*) per cell<sup>3</sup> mean gene to centromere ratio; Ampl, amplification, > 15 copies of gene per cell in > 10 % of cells or gene to CEP ratio > 2.0. An average of 80 (range, 50 to 180) well-defined malignant nuclei were scored