

 DRUG RESISTANCE

# Alternative routes

## DOI:

10.1038/nrc2158

## URLs

## EGFR

[http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full\\_report&list\\_uids=1956](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full_report&list_uids=1956)

## MET

[http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full\\_report&list\\_uids=4233](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full_report&list_uids=4233)

## ERBB3

[http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full\\_report&list\\_uids=2065](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene&md=Retrieve&dopt=full_report&list_uids=2065)

Two tyrosine kinase inhibitors (TKIs) — gefitinib (Iressa) and erlotinib (Tarceva) — are used to treat non-small-cell lung cancers (NSCLCs) with activating mutations of the epidermal growth factor receptor (EGFR). However, following an initial positive response most tumours become resistant to the drugs; in ~50% of cases this occurs because a secondary *EGFR* mutation arises. Now Jeffrey Engelman and colleagues have identified another route, which involves the amplification of the receptor tyrosine kinase *MET*. This bypasses EGFR inhibition and confers resistance to gefitinib, reactivating the EGFR signalling pathway through the EGFR family member *ERBB3*.

By exposing NSCLC cells, which express a gefitinib-hypersensitive EGFR mutant, to increasing drug concentrations for 6 months, the authors generated a cell line — HCC827 GR — that is resistant to gefitinib. Then they analysed the phosphorylation status of 42 receptor tyrosine kinases in response to gefitinib and showed that *MET* and *ERBB3* were both phosphorylated

in HCC827 GR cells, and their phosphorylation persisted during gefitinib treatment, whereas it was reduced in parental cells. A genome-wide copy number analysis showed amplification of the 7q31.1–7q33.3 locus, where *MET* resides, and quantitative PCR confirmed that *MET* expression was increased in the resistant cells. Sequence analysis ruled out the presence of *MET* mutations in these cells.

*MET* signalling is indeed required for the acquired gefitinib resistance of HCC827 GR cells because the use of a *MET* inhibitor, PHA-665,752, in combination with gefitinib treatment, reduced *ERBB3* and Akt phosphorylation, resulting in growth inhibition and apoptosis. Interestingly, phosphorylation of *ERBB3* in HCC827 GR cells was suppressed only in the presence of both inhibitors, indicating that *MET* can signal through *ERBB3* independently of EGFR, whereas *ERBB3* activation was previously thought only to be triggered by other *ERBB* family members.

The *ERBB3* pathway seems to be crucial in mediating *MET*-induced resistance to gefitinib because the downregulation of *ERBB3*,

using short hairpin RNA (shRNA), inhibited Akt phosphorylation and reduced the growth of

both HCC827 GR and parental cells.

Moreover, *MET* downregulation by shRNAs conferred gefitinib sensitivity to the otherwise resistant cells, restoring the ability of the drug to reduce *ERBB3* and Akt phosphorylation.

The finding that *MET* amplification leads to sustained activation of the phosphatidylinositol 3-kinase (PI3K)–Akt pathway by inducing persistent *ERBB3* phosphorylation seems common to other tumours. The authors in fact showed that two gastric cancer cell lines harbouring *MET* amplification had high levels of *ERBB3* in complex with the p85 regulatory subunit of PI3K, and these complexes were disrupted by PHA-665,752.

Does this resistance mechanism also occur in patients? Notably, 4 out of 18 NSCLCs that were resistant to gefitinib or erlotinib treatment showed *MET* amplification, and only one of these cases had a concomitant secondary mutation of EGFR.

The finding that NSCLCs can escape gefitinib inhibition through the amplification of *MET*, which is not the direct target of the drug, has several implications. It highlights the importance of searching for other genetic alterations, besides *EGFR*, that might be responsible for gefitinib and erlotinib resistance in NSCLCs. It also suggests that *MET* inhibitors, which are currently being tested in early phase clinical trials, might be effective in combination with EGFR inhibitors in treating at least a subset of resistant tumours. Finally, it raises the possibility that *MET* amplification might induce drug resistance in other EGFR-dependent tumours.

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**ORIGINAL RESEARCH PAPER** Engelman, J. A. et al. *MET* amplification leads to gefitinib resistance in lung cancer by activating *ERBB3* signaling. *Science* 26 April 2007 (doi: 10.1126/science.1141478)

**FURTHER READING** Sharma, S. V., Bell, D. W., Settleman, J. & Haber, D. A. Epidermal growth factor receptor mutations in lung cancer. *Nature Rev. Cancer* 7, 169–181 (2007)

