

## Microfluid cell-capture chip isolates circulating tumor cells from patients with NSCLC

Despite the dramatic responses to EGFR therapy observed in patients with non-small-cell lung cancers (NSCLC) that harbor activating mutations in *EGFR*, the majority of patients relapse—often because their tumors acquire secondary *EGFR* mutations. Maheswaran and colleagues evaluated the ability of a microfluid cell-capture chip, which contains microposts coated with antibodies against epithelial-cell adhesion molecules, to isolate circulating tumor cells (CTCs) from patients with NSCLC.

The device successfully isolated CTCs (median 74 cells/ml) from each of 27 patients with advanced NSCLC. These isolated CTCs were then used in mutational analysis of *EGFR*. The presence of the *EGFR* T790M mutation, which confers drug resistance, in pretreatment tumor specimens was associated with greatly reduced progression-free survival compared with tumor specimens that lacked this mutation (7.7 months and 16.5 months, respectively;  $P < 0.001$ ). The *EGFR* activating mutation was identified in 11 of 12 patients for whom CTCs, plasma samples and primary tumor specimens were available. CTC genotyping, therefore, had 92% sensitivity to detect such mutations, whereas plasma genotyping (which identified mutations in 4 of 12 patients) had 33% sensitivity ( $P = 0.009$ ). Moreover, increased CTC numbers were associated with clinical progression and the emergence of additional *EGFR* mutations after therapy. Conversely, a decline in CTC numbers was associated with a radiographically evident response to therapy.

The authors suggest that the cell-capture chip can reliably isolate CTCs in an acceptable quantity and with satisfactory purity for use in molecular studies.

**Original article** Maheswaran S *et al.* (2008) Detection of mutations in *EGFR* in circulating lung-cancer cells. *N Engl J Med* 359: 366–377

## Novel bispecific antibody causes tumor regression in patients with non-Hodgkin lymphoma

Clinical data indicate a role for cytotoxic T lymphocytes in the control of tumor growth. However, immunotherapy strategies that induce

antitumor T-cell responses have been characterized by low response rates. Bargou *et al.* assessed the efficacy of blinatumomab—an antibody with dual specificity for CD19 and CD3 antigens—in patients with non-Hodgkin lymphoma who relapsed after standard treatments.

This ongoing study has so far included 38 patients (with hematologic malignancies such as follicular lymphoma, mantle-cell lymphoma and chronic lymphocytic leukemia), who received blinatumomab for 4–8 weeks at doses of 0.0005–0.06 mg/m<sup>2</sup> daily. Target cells that expressed CD19 were depleted in the blood of patients who received  $\geq 0.005$  mg/m<sup>2</sup> of the drug; this decline was shown to be caused by apoptosis. Among 38 patients, 4 complete and 7 partial tumor regressions occurred in patients who received  $\geq 0.015$  mg/m<sup>2</sup> blinatumomab, which suggests that the response is dose-dependent. Tumor regression was observed in all 7 patients treated with 0.06 mg/m<sup>2</sup> blinatumomab. The majority of tumor shrinkage occurred within the first 4 weeks of therapy, and no relapse had been observed at the date of publication in patients treated with 0.03 mg/m<sup>2</sup> and 0.06 mg/m<sup>2</sup> blinatumomab. In patients who received blinatumomab doses  $\geq 0.015$  mg/m<sup>2</sup>, tumor-cell depletion was observed in the blood, lymph nodes, spleen, bone marrow and liver.

These results suggest that blinatumomab is a promising new anticancer agent. Clinical trials that investigate the efficacy of bispecific antibodies, in hematologic and solid tumors, are warranted.

**Original article** Bargou R *et al.* (2008) Tumor regression in cancer patients by very low doses of a T cell-engaging antibody. *Science* 321: 974–977

## Predictors of response to erlotinib in patients with advanced NSCLC

Previous studies in patients with non-small-cell lung cancer have shown that mutations in the genes that encode tyrosine kinase domains of *EGFR* are strongly associated with response to *EGFR* tyrosine kinase inhibitors and could be used as molecular markers.

Zhu *et al.* assessed a subset of patients from the phase III, BR.21 trial of erlotinib in patients with advanced non-small-cell lung cancer, to investigate the effect of *KRAS* mutations, *EGFR* copy number and mutation status on response