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| Title | Epidermal Growth Factor Receptor Mutation Analysis in Advanced Non-Small Cell Lung Cancer: A Review of the Clinical Effectiveness and Guidelines |
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| Reference | 2010 |

Aim

To assess the clinical effectiveness of epidermal growth factor receptor (EGFR) mutation analysis using polymerase chain reaction (PCR) to identify patients with advanced non-small-cell lung cancer (NSCLC) who are likely to respond to treatment with tyrosine kinase inhibitors (TKI); to assess the diagnostic performance of polymerase chain reaction-based methods that are used to evaluate EGFR mutations in patients with advanced NSCLC; and to assess the quality of guidelines on testing for EGFR mutations in patients with NSCLC.

Conclusions and results

Evidence from observational studies suggests that PCR-based approaches can be used to identify mutations in the EGFR gene with a similar sensitivity to that of direct sequencing. PCR-based tests are likely useful for identifying patients with NSCLC who are likely to respond to treatment with a TKI. In several studies, genetic material was microdissected from tumor-rich areas. This may require additional expertise and incur cost. Most of the studies did not use a commercially available reagent kit. Commercially available kits may have quality-control advantages over assays developed “in-house” as regards quality control in a clinical laboratory setting.

Methods

A literature search encompassed key health technology assessment resources, international health technology agencies, and a focused Internet search. The search was limited to articles published in English. Filters were applied to limit the retrieval to health technology assessments, systematic reviews, meta-analyses, randomized controlled trials, controlled clinical trials, observational studies, and guidelines. Two independent reviewers screened articles using predefined criteria. Any disagreements were resolved through discussion until consensus was achieved.