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Showing 8 out of 8 references.	mutations (DEL) in exon 19 and LSSSR in exon 21 <sup>1</sup> , are reportedly correlated with
	clinical outcome in patients with non-small cell lung cancer (NSCLC) receiving the
	EGFR tyrosine kinase inhibitors gefitinib and erlotinib, suggesting that detection of
Keady	EGFR mutations would have an important role in clinical decision making We
	established and validated an easy, inexpensive, and rapid method for detecting DEL
	and LSSSR from <u>cytologic</u> material by high-resolution melting analysis (HRMA).
	Dilution for sensitivity studies revealed that DEL and L858R were detectable in the
	presence of at least 10% and 0.1% EGFR-mutant cells, respectively. We analyzed 37
i i	archived cytological slides of specimens from 29 patients with advanced NSCLC and
	compared the results with direct sequencing data obtained previously. Of 37 samples,
	34 V32N/ yierded consistent results with direct sequencing, 2 were take negative, and 1 was indetermined to The constitution of this analysis was 90% (19/21) and exactlificity.
	100% (15/15) These results surgest that HRMA of archived cutoheir specificity,
	advanced NSCI C is useful for distancing EQE mutations is clinical practice u
	and the second
	<sup>1</sup> K. Nomoto, K. Tsuta, T. Takano et al., Am J <u>Glin Pathol</u> <b>126</b> (4), 1 (2006).
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	(2006)
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