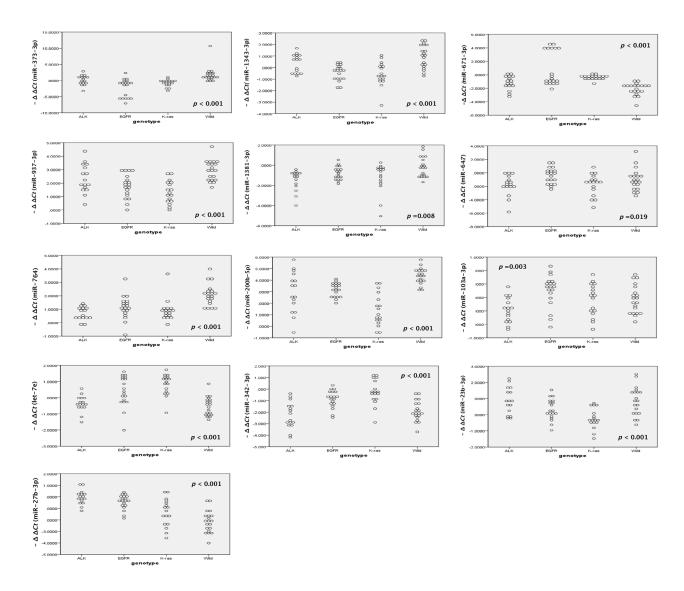
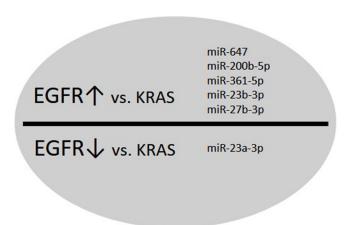
MicroRNA expression profiles and clinicopathological implications in lung adenocarcinoma according to *EGFR*, *KRAS*, and *ALK* status

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Dot plots of microRNA expressions in lung adenocarcinomas according to genetic status (ALK-rearranged, EGFR-mutated, KRAS-mutated, and triple negative).



Supplementary Figure 2: Differential expressions of microRNAs in *EGFR*-mutated lung adenocarcinomas, compared to *KRAS*-mutated lung adenocarcinomas.

Supplemetary Table 1: Association between ALK rearrangement and E-cadherin expression in lung adenocarcinomas of EGFR, KRAS and ALK groups (n = 52)

	Total N(%)	E-cadherin, N(%)		
		No loss	Loss	p value
ALK	16 (30.8)	6 (37.5)	10(62.5)	0.002*
EGFR+KRAS	36 (69.2)	29(80.6)	7 (19.4)	

^{*} Statistically significant (p < 0.05)