

CORRECTION

Correction: Methylation Markers of Early-Stage Non-Small Cell Lung Cancer

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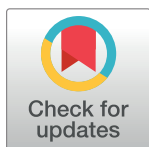
The incorrect file for [S3 Fig](#) is included in the Supporting Information. Please see the correct [S3 Fig](#) here.

Supporting Information

S3 Fig. Scatterplots showing the correlation between beta values determined by Illumina methylation arrays and Sanger bisulfite sequencing. Shown are the best-fitting line, Pearson's correlation coefficient and p-value of correlation tests. The mean correlation between two methods was 83% (range 48.9%; 97.4%).
(TIFF)

Reference

1. Lakk K, Vooder T, Kolde R, Väik K, Võsa U, Roosipuu R, et al. (2012) Methylation Markers of Early-Stage Non-Small Cell Lung Cancer. PLoS ONE 7(6): e39813. doi: [10.1371/journal.pone.0039813](https://doi.org/10.1371/journal.pone.0039813) PMID: [22768131](https://pubmed.ncbi.nlm.nih.gov/22768131/)



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