

Correction: Oncogenomics of c-Myc transgenic mice reveal novel regulators of extracellular signaling, angiogenesis and invasion with clinical significance for human lung adenocarcinoma

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This article has been corrected: In the Materials and Methods section, regarding the Isolation of RNA, production of cRNA, array hybridization and scanning, the correct microarray GEO information is given below:

Isolation of RNA, production of cRNA, array hybridization and scanning

Total RNA was extracted using the Qiagen RNA isolation kit according to the manufacturer's instructions (Qiagen, Hilden, Germany) and copy RNA was prepared according to the Affymetrix Gene Chip[®] Expression Analysis Technical Manual (Santa Clara, CA). Further details are given in our recent report [6]. The microarray study is MIAME compliant, and the raw data have been deposited in GEO database (GSE54829; <http://www.ncbi.nlm.nih.gov/geo>).

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