

CORRIGENDUM

Specific microRNAs are downregulated in human thyroid anaplastic carcinomas

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The authors wish to replace Figure 1a with the following figure, which shows an alternative experiment. Representative ATC (anaplastic thyroid carcinoma) ($n=6$) and PTC (papillary thyroid carcinoma) ($n=6$) samples were analyzed for the expression of the miR-30d, miR-125b, miR-26a, miR30a-5p. As control, three normal thyroid samples were used. The results confirm the strong downregulation of the examined miRs in ATC but not in PTC samples. For this experiment, qRT-PCR analysis was performed in 6 papillary (PTC1-6) and 6 anaplastic (ATC1-6) thyroid carcinoma samples. cDNAs were generated by using

miScript Kit (Qiagen, Valencia, CA, USA), and amplifications were performed by using miScript SYBR Green PCR Kit (Qiagen) and miScript Primer Assays for miR-30d, miR-125b, miR-26a, miR30a-5p and U6 for normalization (Qiagen). Reactions were performed in duplicate in a real-time CFX96 thermocycler (Bio-Rad, Hercules, CA, USA). The relative quantity of each microRNA was normalized to the U6 RNA amount by using the 2- $DDCt$ formula and the values indicate relative expression levels in tumor samples \pm s.d. compared with average expression levels of three normal thyroid samples.

In Figures a–d, bar graphs indicate expression levels of (a) miR-30d, (b) miR-125b, (c) miR-26a and (d) miR30a-5p.

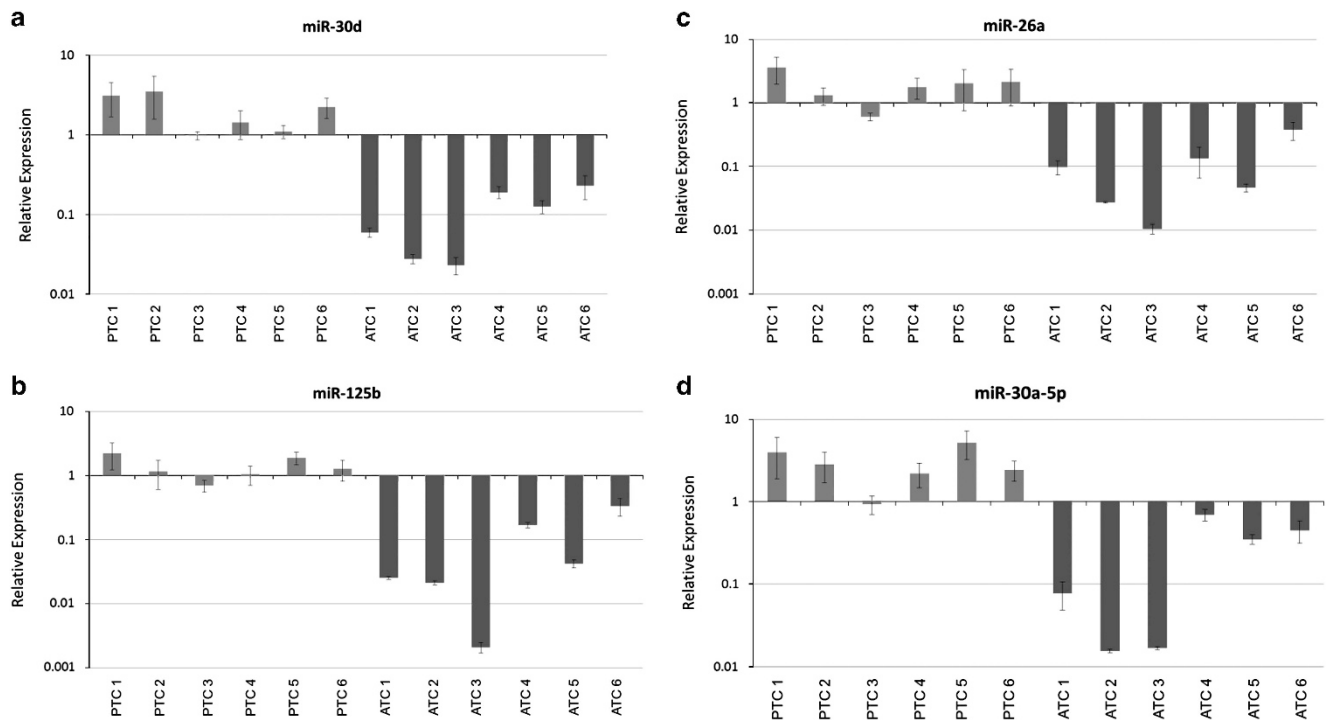


Figure 1.