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RECEIVED 12 March 2025  
ACCEPTED 07 April 2025  
PUBLISHED 25 April 2025

CITATION  
Misir S, Yaman SO, Hepokur C, Akidan O,  
Aliyazicioglu Y, Enguita FJ and Al Zoubi MS  
(2025) Corrigendum: Identification of miR-342-  
5p/MDM4/p53 network in acute  
myeloid leukemia.  
*Front. Cell Death* 4:1592636.  
doi: 10.3389/fceld.2025.1592636

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# Corrigendum: Identification of miR-342-5p/MDM4/p53 network in acute myeloid leukemia

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## KEYWORDS

acute myeloid leukemia, biomarkers, bioinformatics, non-coding RNAs, miRNA

## A Corrigendum on Identification of miR-342-5p/MDM4/p53 network in acute myeloid leukemia

by Misir S, Ozer Yaman S, Hepokur C, Akidan O, Aliyazicioglu Y, Enguita FJ and Al Zoubi MS (2024).  
*Front. Cell. Death* 3:1503241. doi: 10.3389/fceld.2024.1503241

In the published article, there was an error in the legend for **Figures 2, 3** as published. The legends for **Figures 2, 3** were placed in the wrong order. The legend for **Figure 2** should be **Figure 3**, and the legend for **Figure 3** should be **Figure 2**. The corrected legend appears below.

“**FIGURE 2** | Prediction of miRNA targets from miRDB in HL-60 and KG-1 cells. (A) There are 119 predicted targets for hsa-miR-342-5p with expression level  $\geq 5$  in cell line HL-60. (B) There are 141 predicted targets for hsa-miR-342-5p with expression level  $\geq 5$  in cell line KG-1. In addition, those with a target score greater than 50 in all cells were selected. miR-342-5p and putative mRNAs and visualized using by Cytoscape software 3.8.”

“**FIGURE 3** | (A) Receiver operating characteristic curve analysis of miR-342-5p expression levels in control and AML groups was determined using data from GSE51908. (B) The Kaplan–Meier survival curve of AML and Control groups. Survival analysis was performed using data from **Bullinger et al. (2010)**.”

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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