

Erratum

Genome-wide target interactome profiling reveals a novel *EEF1A1* epigenetic pathway for oncogenic lncRNA *MALAT1* in breast cancer: Am J Cancer Res. 2019; 9(4): 714-729

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In this article, we found an image error in **Figure 7C**. This was caused by file transfer between lab computers, resulting in duplicated images in **Figure 7C**. These two images are exactly the same, which obviously contradicts with the chart on the right. Therefore, we would like to publish this Erratum to reflect this change. We apologize for this error.

The corrected **Figure 7** is as follows.

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MALAT1 epigenetically targets *EEF1A1*

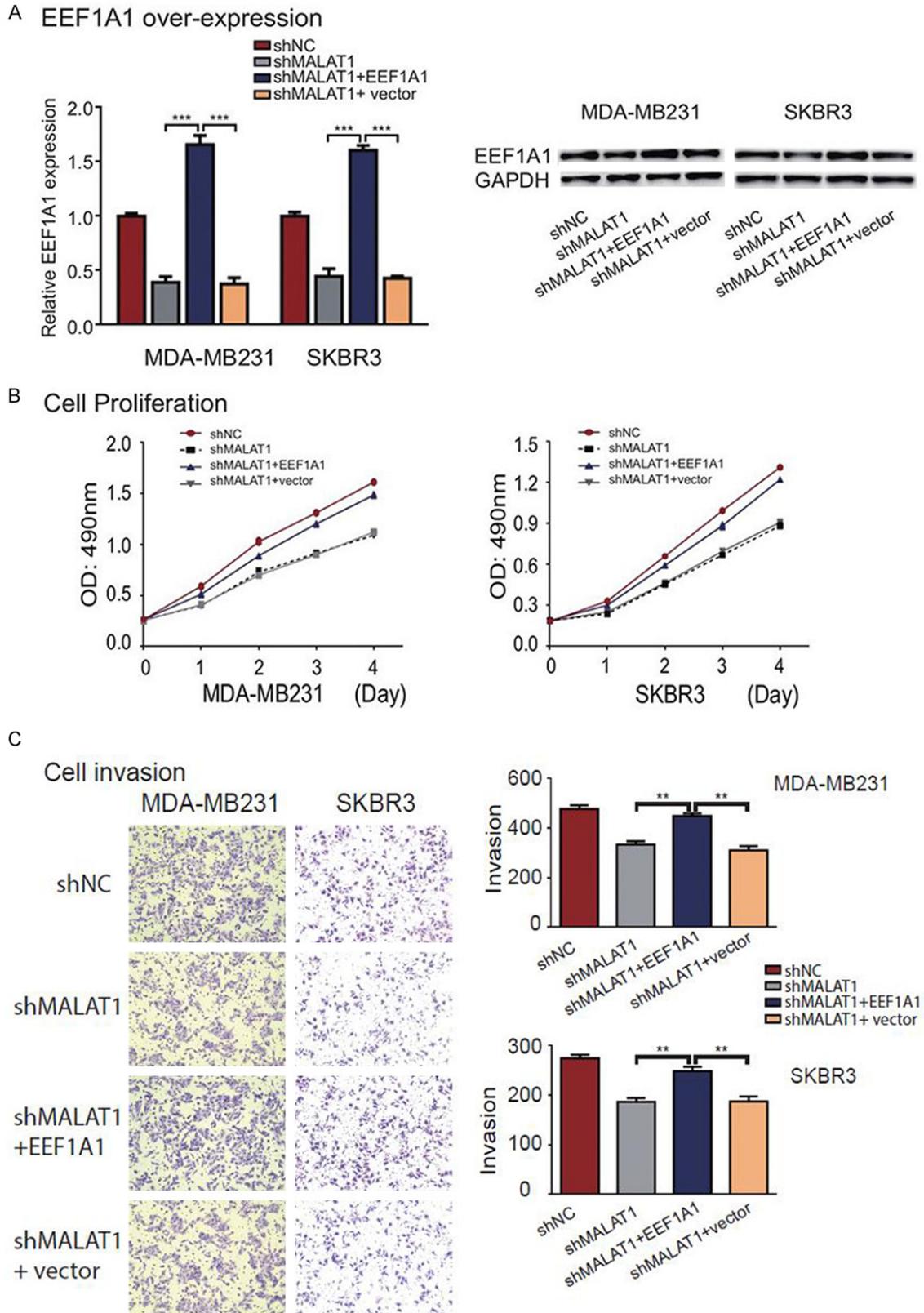


Figure 7. *EEF1A1* rescues the effect induced by *MALAT1* knockdown. A. Overexpression of *EEF1A1* in breast cancer cells. The expression of *EEF1A1* was quantitated by Q-PCR. shNC: random shRNA control; shMALAT1: Cells that were transfected with *MALAT1* shRNA-1. β -Actin was used as an internal control. *** $P < 0.001$ as compared with the control groups. Overexpression of *eEF1A1* in breast cancer cells was measured by Western blot. B. Cell growth viability as measured by CCK-8 assay. C. Cell invasion as examined by Transwell assay. Quantitation of invaded cells was shown as mean \pm SD, ** $P < 0.01$ as compared with the control groups.