Clinical Epigenetics

CORRECTION

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Correction: miR-29c plays a suppressive role in breast cancer by targeting the TIMP3/STAT1/FOXO1 pathway

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In the original publication of this article [1], there was an error in the Fig. 6. The MDA-MB-231 Invasion Cntl group in Fig. 6g and the MDA-MB-436 Migration Cntl group in Fig. 6h were wrongly chosen. Theincorrect and correct Fig. 6 are shown in this correction article.

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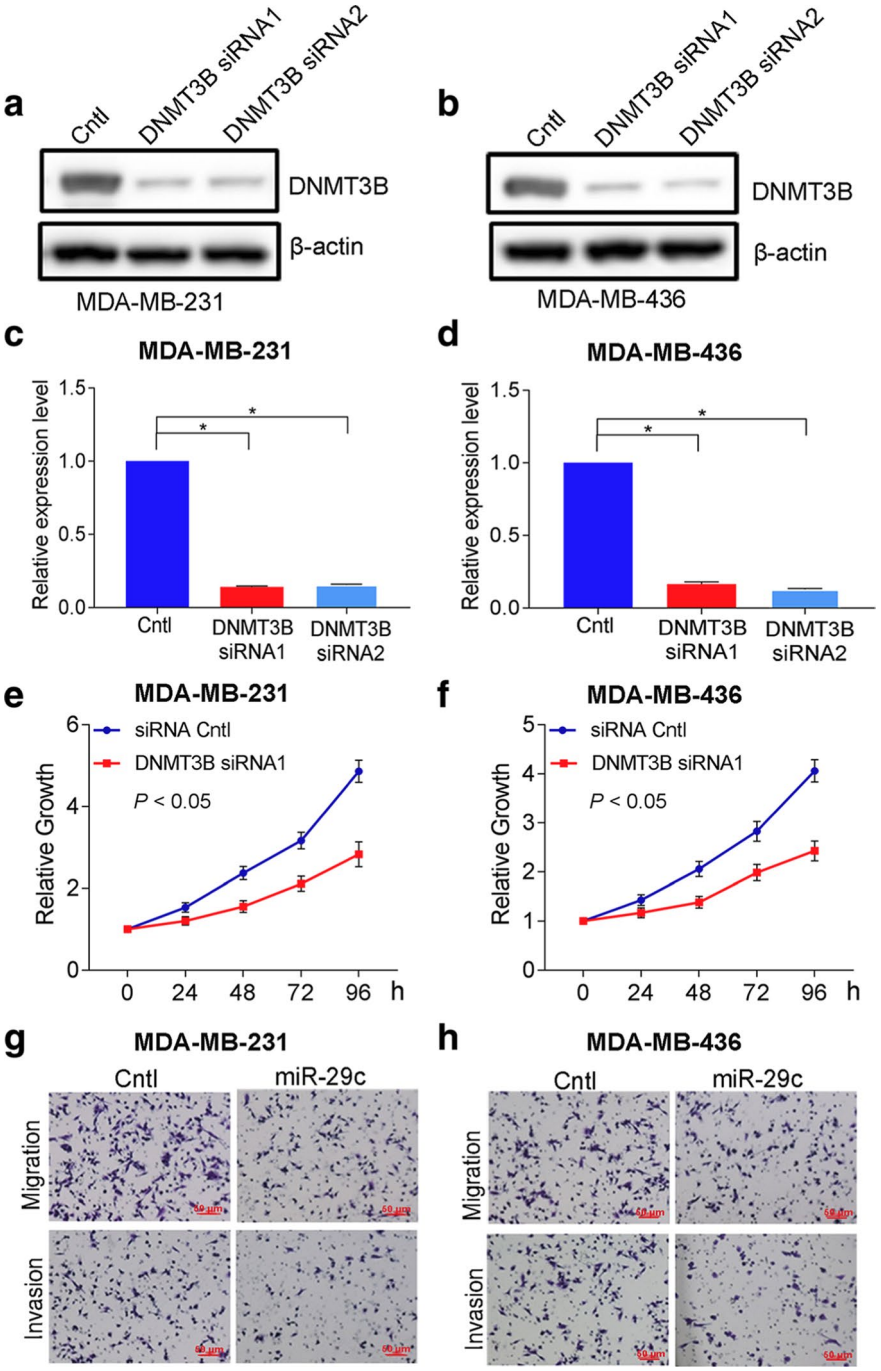
Incorrect Fig. 6.



(See figure on next page.)

Fig. 6 Knockdown of DNMT3B inhibited the proliferation, migration, and invasion in MDA-MB-231 and MDA-MB-436 cells. **a** Protein levels of DNMT3B detected by Western blotting in MDA-MB-231 cells after the transfection of DNMT3B siRNA 1 and 2. **b** Protein levels of DNMT3B detected by Western blotting in MDA-MB-436 cells after the transfection of DNMT3B siRNA 1 and 2. **c** Quantification of DNMT3B protein levels in MDA-MB-231 cells after the transfection of DNMT3B siRNA 1 and 2. **d** Quantification of DNMT3B protein levels in MDA-MB-436 cells after the transfection of DNMT3B siRNA 1 and 2. **e** CCK-8 proliferation assays of MDA-MB-231 cells after the transfection of DNMT3B siRNA. **f** CCK-8 proliferation assays of MDA-MB-436 cells after the transfection of DNMT3B siRNA. **g** Migration and invasion assays of MDA-MB-231 cells after the transfection of DNMT3B siRNA. **h** Migration and invasion assays of MDA-MB-436 cells after the transfection of DNMT3B siRNA. Data are presented as mean ± SD from three independent experiments, and every experiment was repeated three times

Correct Fig. 6.

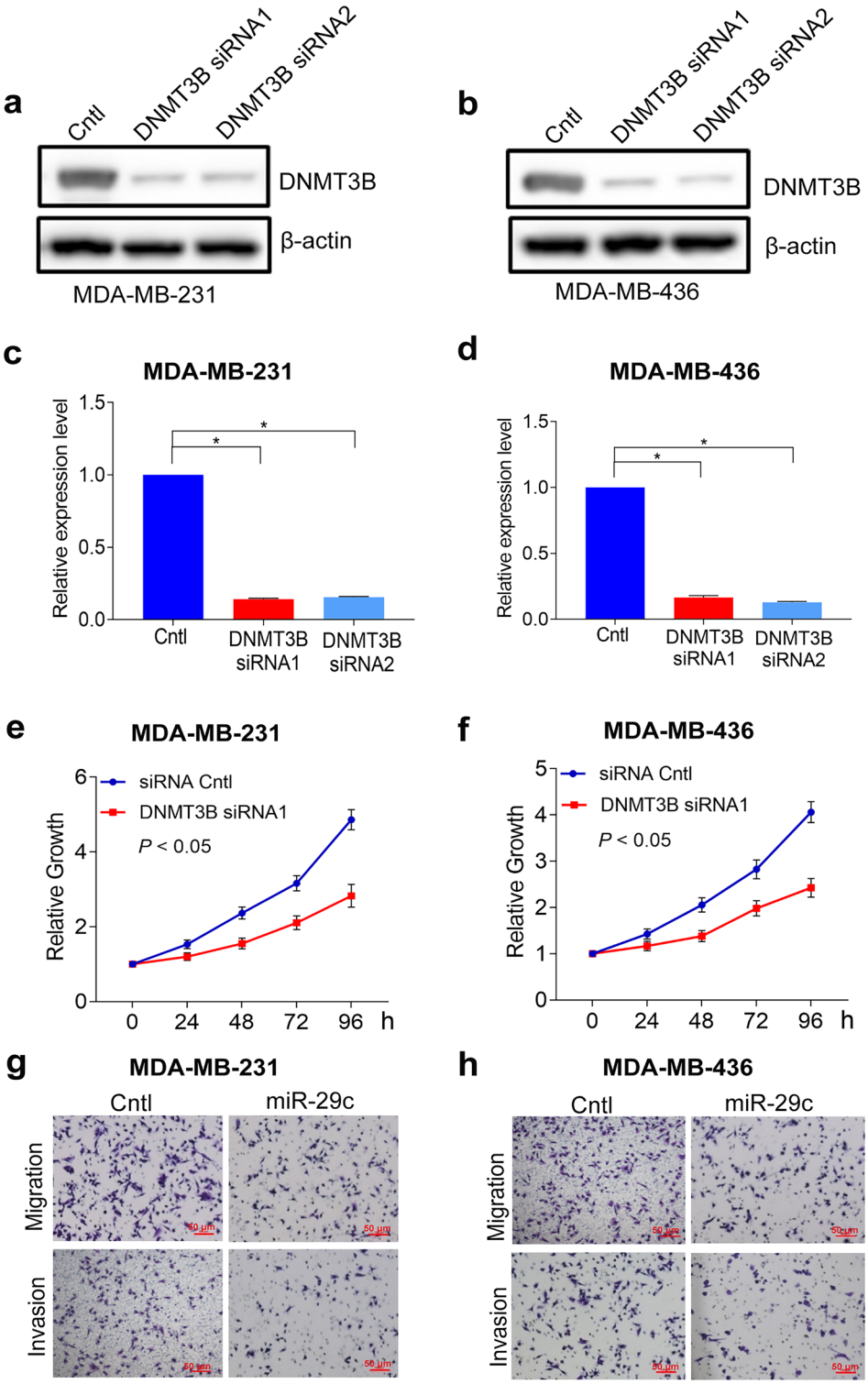


Fig. 6 (See legend on previous page.)

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Reference

1. Li W, Yi J, Zheng X, et al. miR-29c plays a suppressive role in breast cancer by targeting the TIMP3/STAT1/FOXO1 pathway. Clin Epigenet. 2018;10:64.

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