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Correction: Resveratrol and Angiogenin-2 combined with PEGDA/TCS hydrogel for the targeted therapy of hypoxic bone defects via activation of the autophagy pathway

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KEYWORDS

resveratrol, ANG2, autophagy, hypoxia condition, vascularization, bone defect

A Correction on

Resveratrol and Angiogenin-2 combined with PEGDA/TCS hydrogel for the targeted therapy of hypoxic bone defects via activation of the autophagy pathway

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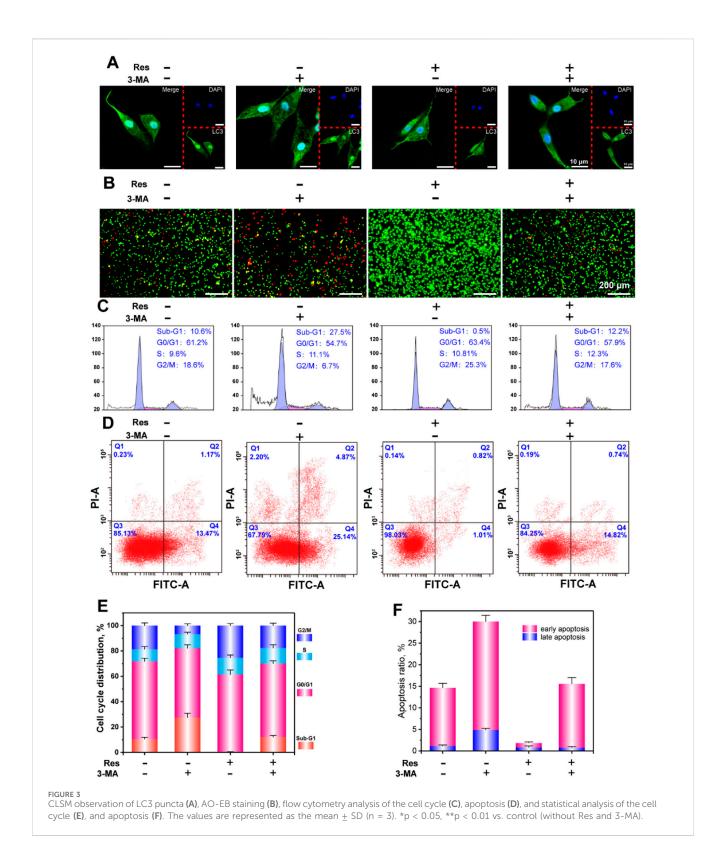
In the published article, there was an error in Figures 3B, 4A,B as published. Images from different groups were stored in the same folder during image shooting, resulting in the misuse of images. The corrected Figures 3B, 4A,B and their captions appear below.

In the published article, there was an error in Figures 7A, 8 as published. The H&E, Masson, OCN and CD31 staining images of the animal tissue sections in Figures 7, 8 overlap with those used in our previous published study. This occurred because the animal experiments for both research projects were conducted concurrently, leading to inadvertent misplacement of data files. The corrected Figures 7A, 8 and their captions appear below.

The original article has been updated.

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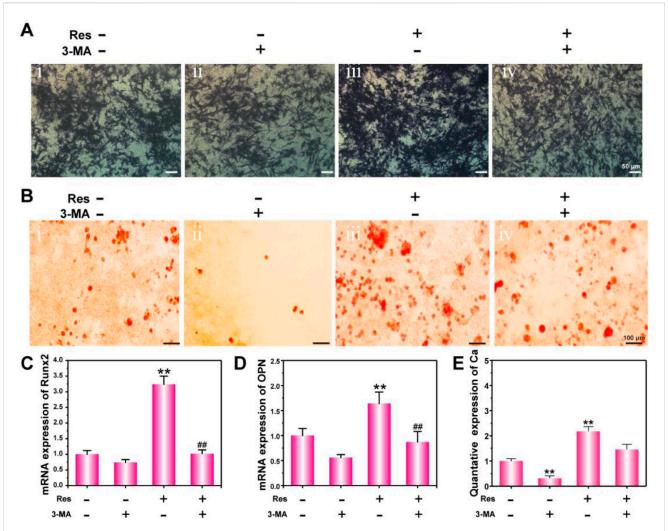


FIGURE 4 Osteogenic differentiation of BMSCs treated with Res and 3-MA for 14 days. ALP activity detection (A) and ARS staining (B). mRNA expression of Runx2 (C) and OPN (D) detected by q-PCR. The quantative data of Ca nodulus analyzed by ARS staining using ImageJ software (E). The values are represented as the mean \pm SD (n = 3). *p < 0.05, **p < 0.01 vs. control (without Res and 3-MA); #p < 0.05, ##p < 0.01 vs. Res group.

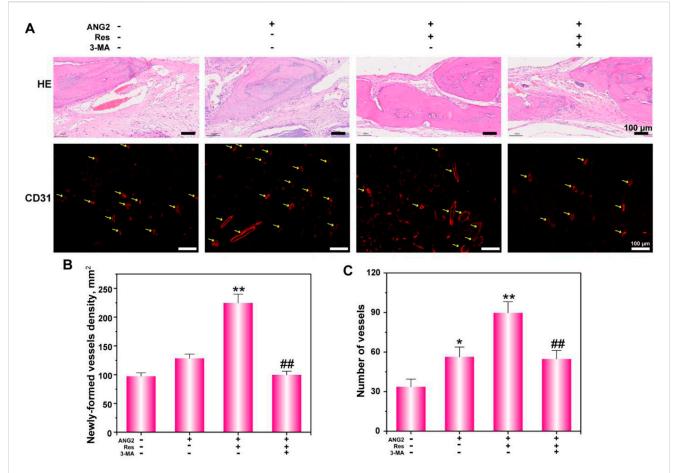


FIGURE 7 Histological analysis (A) of bone defects via H&E staining and CD31 immunofluorescence staining. (B) The newly formed vessel density and (C) number were determined. The values are represented as the mean \pm SD (n = 6).*p < 0.05, **p < 0.01 vs. ANG2 group; #p < 0.05, ##p < 0.01 vs. ANG2/Res group.

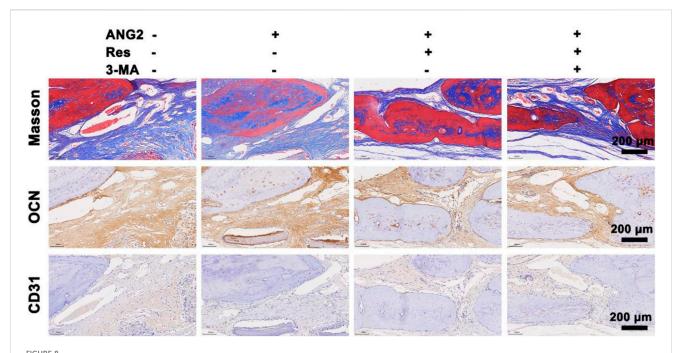


FIGURE 8 Masson staining and immunohistochemical staining for OCN and CD31 in the bone defect area at the 8th week. The values are represented as the mean \pm SD (n = 6).