Correction

## Correction for: Ubiquitin-specific protease 4 promotes metastasis of hepatocellular carcinoma by increasing TGF- $\beta$ signaling-induced epithelial-mesenchymal transition

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**This article has been corrected:** The authors submitted the wrong Figure 5 (panel F) from other experiment group in this study and requested to replace it. The correct Figure 5 is provided below. The authors declare that this correction does not change the results or conclusions of this paper. The authors sincerely apologize for this error.

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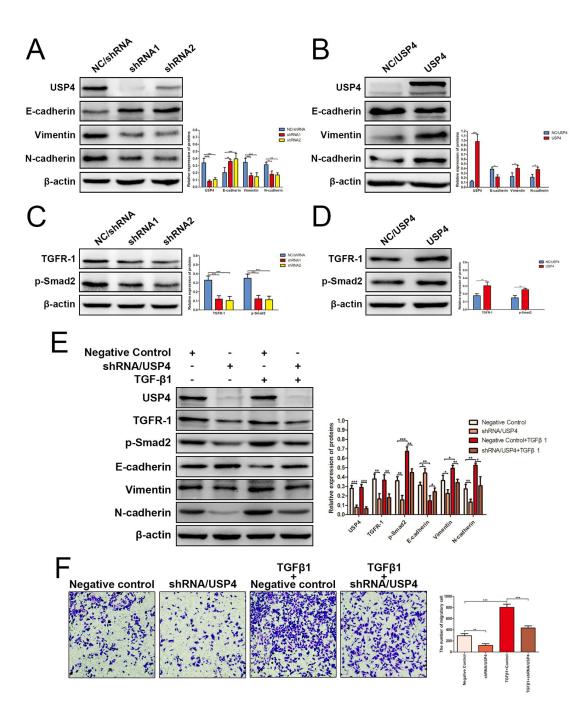


Figure 5. USP4 activated TGF-β signaling pathway to induce epithelial-to-mesenchymal transition (EMT). (A) The expression of EMT markers in SK-Hep1- shRNA/USP4 cells and in negative control cells (\* P < 0.05, \*\* P < 0.01, \*\*\* P < 0.001). (B) The expression of EMT markers in HuH7-USP4 cells and in negative control cells (\* P < 0.05, \*\*\* P < 0.001). (C) The expression of TGFR-1 and p-Smad2 in SK-Hep1- shRNA/USP4 cells and in negative control cells (\*\*\* P < 0.001). (D) The expression of TGFR-1 and p-Smad2 in HuH7-USP4 cells and in negative control cells (\* P < 0.05, \*\* P < 0.01). (E) The expression of TGFR-1, p-Smad2 and EMT markers in SK-Hep1- shRNA/USP4 cells or in negative control cells, with TGF-β1 (10ng/mL) treatment for 24h or without (\* P < 0.05, \*\* P < 0.01, \*\*\* P < 0.01, \*\*\* P < 0.01, \*\*\* P < 0.001). (F) The migration ability of SK-Hep1-shRNA/USP4 cells or negative control cells, with TGF-β1 treatment or without, detected by Transwell migration assay (\*\* P < 0.01, \*\*\* P < 0.001).