

ITGA2 Overexpression Promotes Esophageal Squamous Cell Carcinoma Aggression via FAK/AKT Signaling Pathway [Corrigendum]

Huang W, Zhu J, Shi H, Wu Q, Zhang C. *Onco Targets Ther.* 2021;14:3583–3596.

The authors have advised there is an error in Figure 2E on page 3589. The Y axis “Migrative Cells per field” is shown as 0, 20, 40, 60, 80, and 100 and should read 0, 50, 100, 150 and 200. The correct Figure 2 is as follows.

The authors apologize for this error and advise it does not affect the results of the paper.

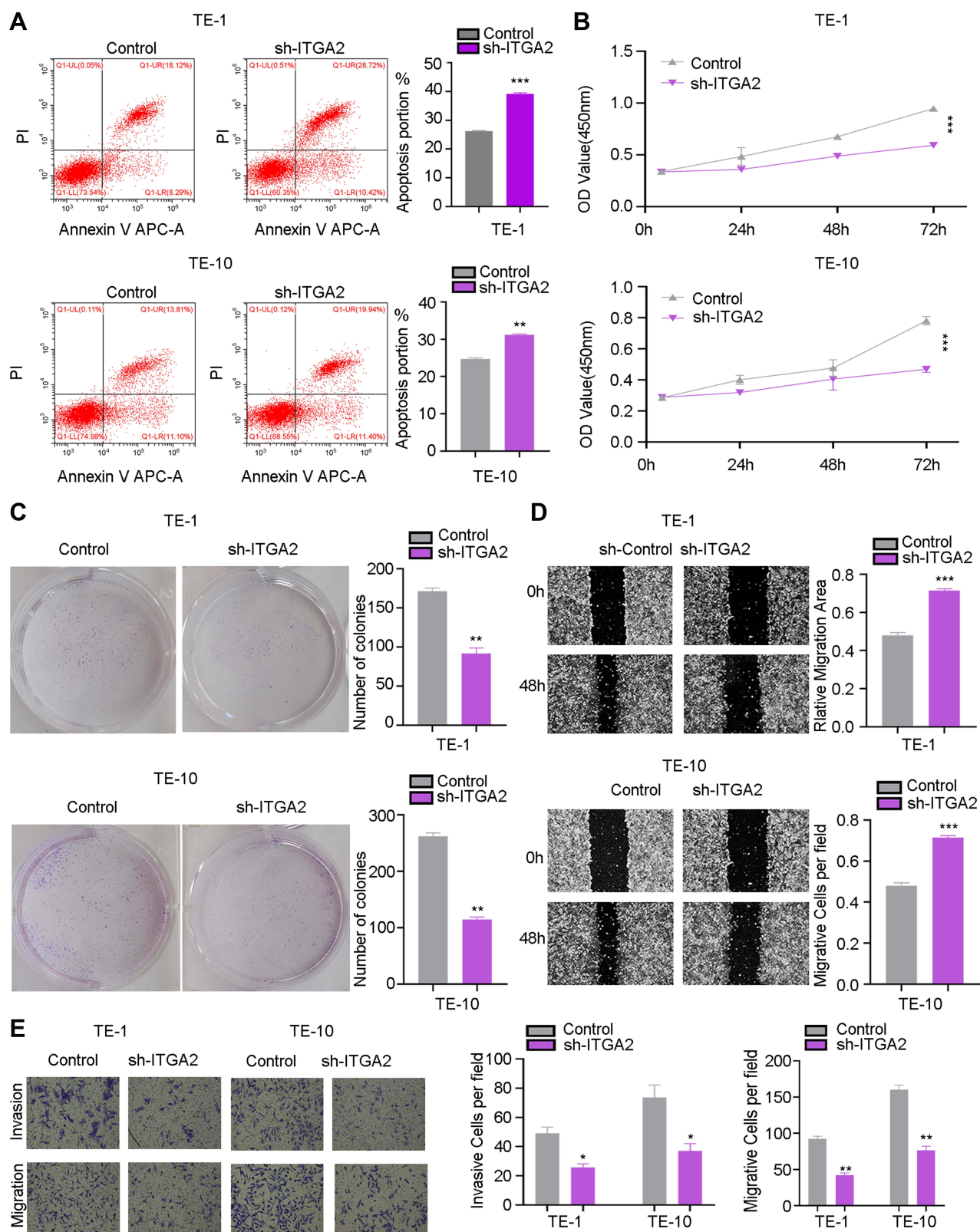


Figure 2 Silencing ITGA2 inhibits ESCC cells aggressiveness. (A–E) TE-1 and TE-10 cells were stably transfected with sh-Control or sh-ITGA2. Flow cytometric analysis of apoptosis (A), CCK8 assay (B), colony formation assay (C), wound healing assay (D), invasion and migration assay (E). For (E), displayed invasion and migration images are based on Transwell experiment. Sh-ITGA2 group and sh-Control group were compared. Each bar displays the mean±SD of 3 independent experiments as analyzed by paired two-tailed students, t-test. * $p<0.05$; ** $p<0.01$; *** $p<0.001$.

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