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Author Correction: Snail collaborates with EGR-1 and SP-1 to directly activate transcription of MMP 9 and ZEB1

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This Article contains errors in Figures 6b and 6c where the 'shRNA' axes labels are missing. Additionally, in Figure 6D, 'shRNA' is erroneously included as part of the 'ZEB1/GAPDH' label. The correct Figure 6 appears below as Figure 1.

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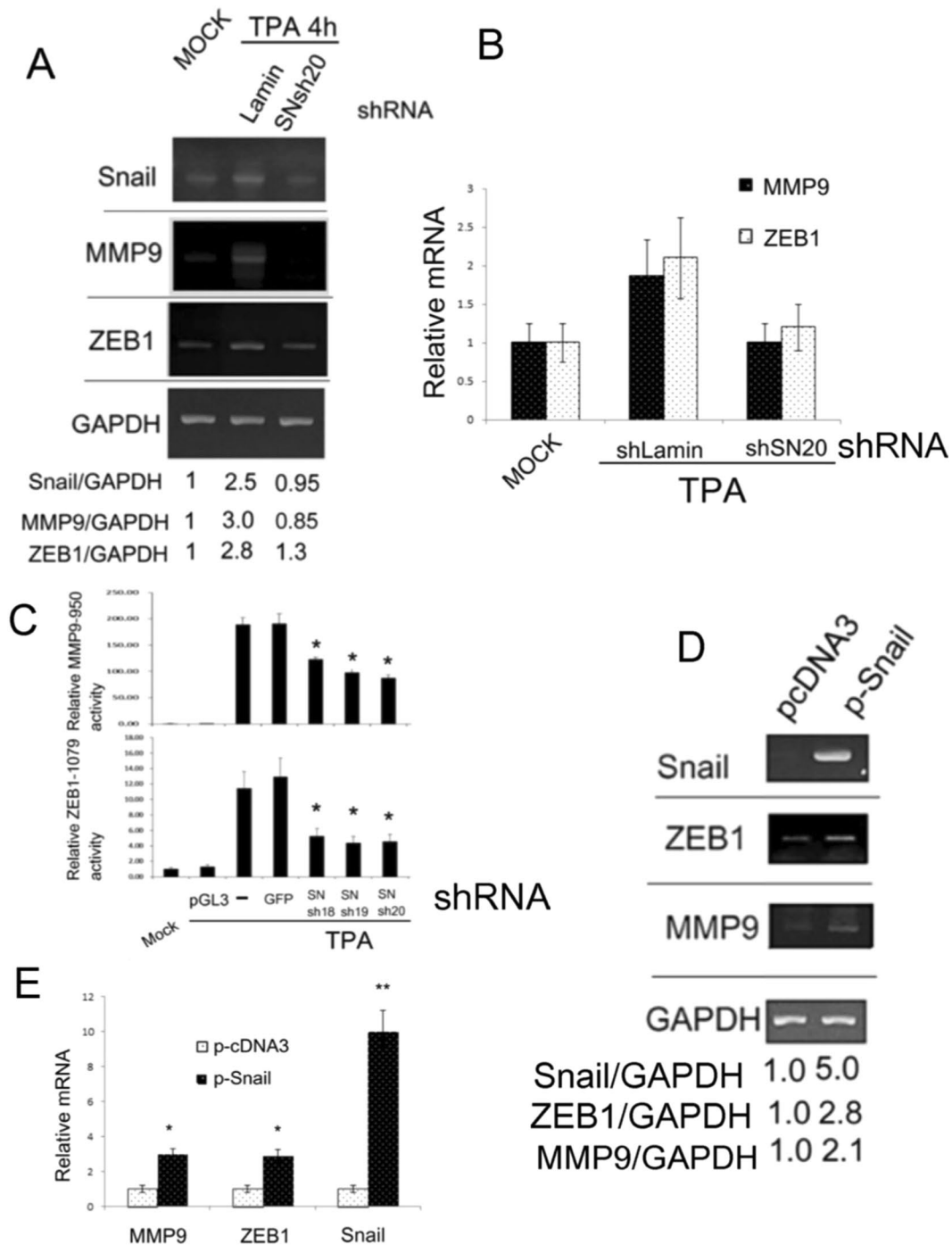


Figure 1. Snail is essential for expression and promoter activation of MMP9 and ZEB1. HepG2 cells were transfected with none (MOCK) or plasmids encoding indicated Snail shRNAs (**A,B,C**) and control shRNA of Lamin (**A**), (**B**) or GFP (**C**) for 24 h followed by untreated (MOCK) or treated with 50 nM TPA for 4 h (**A**); 4 and 6 h for ZEB1 and MMP9, respectively (**B**); or 12 h (**C**). HCC340 were transfected with pcDNA3 vector or Snail expressing plasmid (p-Snail) for 36 h (**D**) and (**E**). RT-PCR (**A**), (**D**) and quantitative RT-PCR (**B**), (**E**) of Snail, MMP9 and ZEB1 and promoter assay of MMP9-950 (**C**, upper panel) and ZEB1-1079 (**C**, lower panel) were performed. In (**A,D**), the numbers below each figure are the ratios of relative mRNA based on RT-PCR of indicated transcriptional factor vs GAPDH, taking the data of MOCK (**A**) and pcDNA3 (**D**) as 1.0. The results are average of 3 reproducible experiments with C.V. of 7.5%. In (**B**) and (**E**), the relative mRNA was calculated based on real time RT-PCR, taking the data of MOCK (**B**) and pcDNA3 (**E**) as 1.0. In (**C**), the relative dual luciferase activity of MMP9-950 or ZEB1-1079 was calculated, taking MOCK as 1.0. The data in (**B**), is average of two representative experiment with C.V. of 12%. In (**C**) and (**E**), (**,*) represent the statistically significant difference ($p < 0.005$, $N = 3$), ($p < 0.05$, $N = 3$) between the indicated samples and the control GFP shRNA (**C**) and pcDNA3 (**E**) group.



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