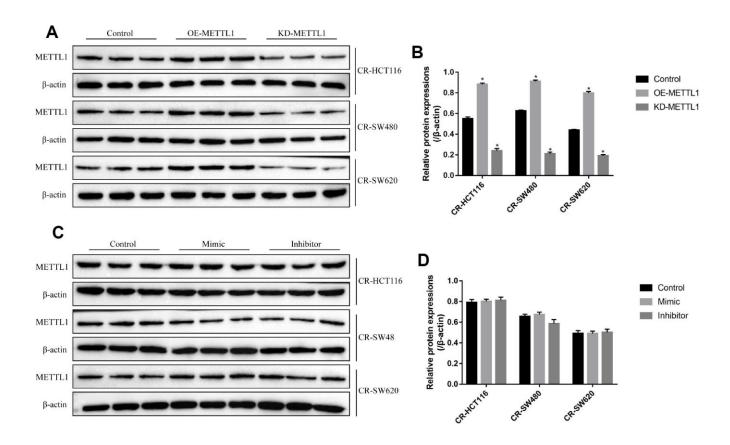
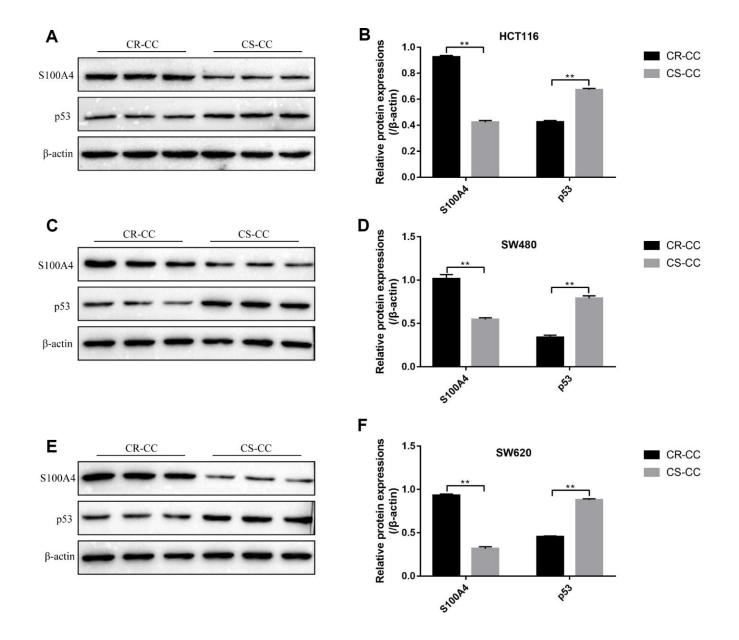
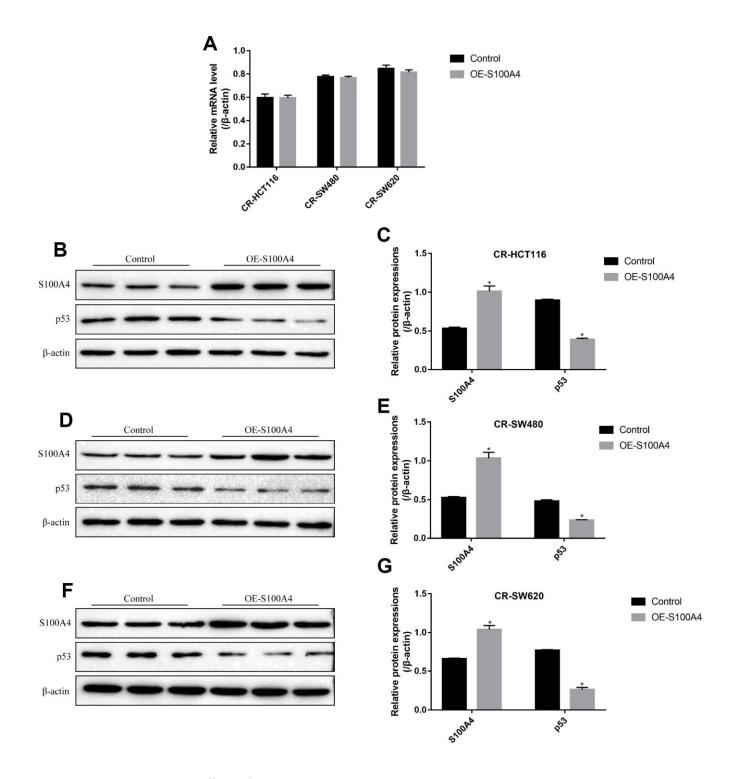
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Western Blot was used to determine (**A**) the expression levels of METTL1 in CR-CC cells and (**B**) the effects of miR-149-3p on METTL1 levels in CR-CC cells. All the experiments repeated at least 3 times. "*" means p < 0.05, "**" means p < 0.01 and "NS" means no statistical significance.



Supplementary Figure 2. Western Blot was used to determine the expression levels of S100A4 and p53 in (**A**, **B**) HCT116 cells, (**C**, **D**) SW480 cells and (**E**, **F**) SW620 cells. All the experiments repeated at least 3 times. "*" means p < 0.05, "**" means p < 0.01 and "NS" means no statistical significance.



Supplementary Figure 3. The effects of **S100A4** on **p53** levels in **CR-CC** cells. (A) Real-Time qPCR was used to detect S100A4 mRNA levels. Western Blot was used to determine the expression levels of S100A4 and p53 in (B, C) CR-HCT116 cells, (D, E) CR-SW480 cells and (F, G) CR-SW620 cells. All the experiments repeated at least 3 times. "*" means p < 0.05, "**" means p < 0.01 and "NS" means no statistical significance.