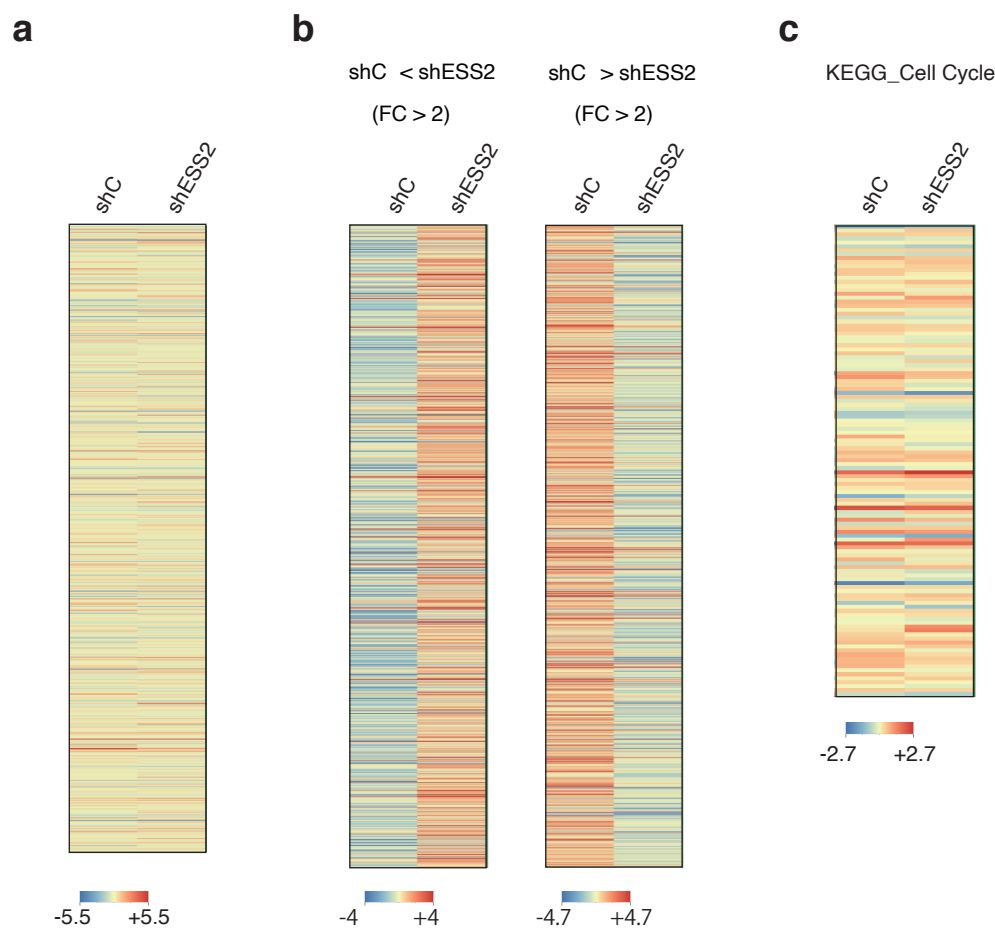


Supplementary Figure 4



Supplementary Figure 4: Microarray analysis of PC3-shC and PC3-shESS2 cells (Data related to Fig. 2). **(a)** Heatmap analysis of whole gene mRNA in PC3-shC and PC3-shESS2 cells. **(b)** Heatmap of gene mRNA groups with more than 2-fold differences in expression in PC3-shC and PC3-shESS2 cells. The left panel shows overexpressed in PC3-shESS2 cells and the right panel shows downregulation in PC3-shESS2 cells. **(c)** Heatmap of expression in cell cycle-related gene sets defined by KEGG.