A. MAPPER2 prediction

The number of binding sites predicted for different transcription factors (TFs) is shown in the bar chart. The factors include CEBPB, E2F1, PAX6, RELA, and STAT1.

B. Down-regulated genes in TIVE

The bar chart illustrates the number of down-regulated genes for the same TFs (CEBPB, E2F1, PAX6, RELA, STAT1). The genes are categorized into three groups: Not DEGs, Containing TFBS but not miR BS, and Containing both TFBS and miR BS.