Supplementary S3 Identification of E2A response genes. (A) Volcano plots of differentially expressed E2A relative to the control group. (B) Top 20 up-regulated BP terms during E2A silence in AML cells were ranked by enrichment score. (C) Top 20 up-regulated CC terms during E2A silence in AML cells were ranked by enrichment score. (D) Top 20 up-regulated MF terms during E2A silence in AML cells were ranked by enrichment score. (E) KEGG analysis of up-regulated pathway genes when E2A was knockout.