

## **Description of Additional Supplementary Files**

**File Name: Supplementary Data 1**

**Description:** Differentially-expressed transcripts in response to HOXA9 depletion.

**File Name: Supplementary Data 2**

**Description:** Functional annotation clustering of regulated genes in HOXA9-depleted cells.

**File Name: Supplementary Data 3**

**Description:** KEGG pathway analysis of regulated genes in HOXA9-depleted cells.