

Heatmap visualization showing gene expression data across various cell lines. The y-axis lists 100 genes, and the x-axis lists 10 cell lines: H8_CRX_ROs_D45, EP1_WT_ROs_D45, H9_BRN3B_RO, EP1_BRN3B_RO, EP1_WT_HRO_2, H9_BRN3B_HRO_2, H8_CRX_HRO_2, H9_HRCG_2, and H9_HRCG_1. A dendrogram on the left shows hierarchical clustering of genes, and a dendrogram on the top shows clustering of cell lines. A color scale at the top indicates expression levels from low (blue) to high (red).

