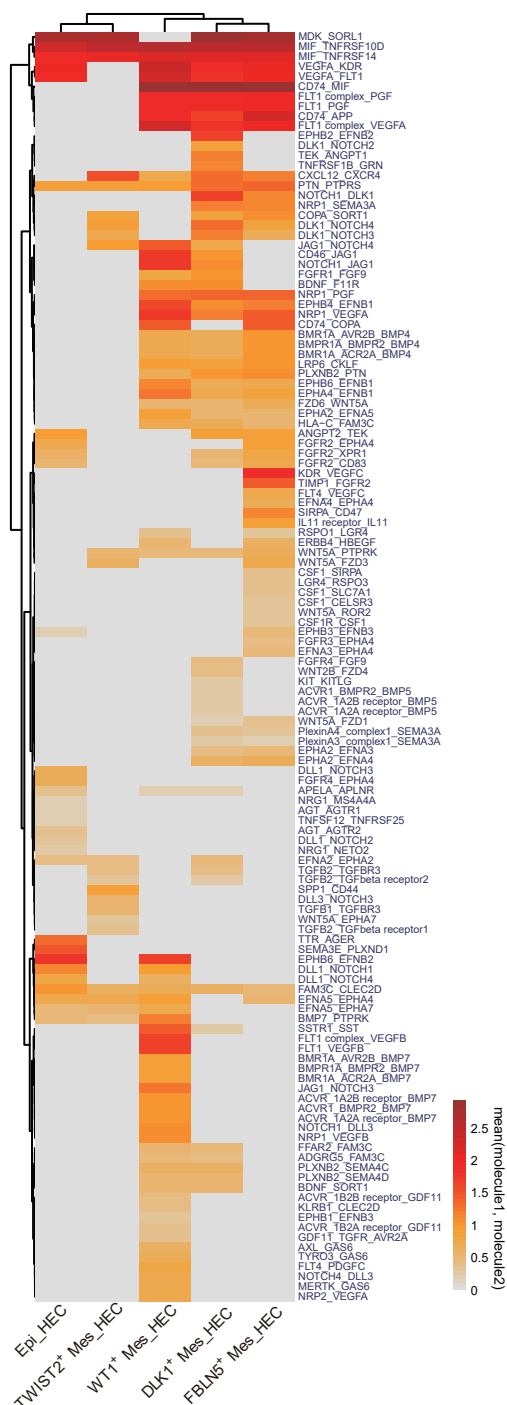




Supplementary Figure 5. The computationally predicted cell-cell interactions that potentially involved in the development of HSC-primed hemogenic ECs

Heatmap showing the scaled average expressions of ligand-receptor pairs in distinct stromal populations when coupled with HEC. Complete list and detailed ligand-receptor genes are shown. The hierarchical clustering result of cell populations (columns) and ligand-receptor pairs (rows) is shown.