Supplementary FIGURE.  Distinct gene signatures among cHCEC SPs. A.
The gene signatures were compared between CD44-SPs and SPs with CST
using RT² profiler PCR-Array Human Extracellular Matrix and Adhesion
Molecules. The cellular features of #66 passage 5 and #67 passage 5 are as
described in the legend of Figure1. B. The mRNA were extracted from CD44+++ 
cHCEC SPs after transfection of 378a-3p or 378f mimics. The hierarchical 
clustering of the gene signatures was determined using RT² profiler PCR-Array 
of senescence, EMT, Fibrosis, p53, and EMA, and are illustrated as heat maps.