

Supplementary Figure 2. A CORE of open chromatin associated with the KLF5 gene promoter in human tracheal epithelial cells.

A region of the human genome encompassing the KLF5 gene is shown on the UCSC genome browser (http://genome.ucsc.edu) with combined DNase Solexa sequence data and peak calls for 2 biological replicates of human tracheal epithelial cells (HTE) and skin fibroblasts (Fibroblasts_park). Above these tracks are HTE-selective COREs (horizontal purple bars, HTE-selective COREs), HTE-selective DHS (vertical purple bars, HTE-selective-sites).