



Fig. S2 (related to Fig. 1): MiR-137 is less expressed in MCF7 breast cancer cells than in HMEpC normal cells. Levels of miR-137 from RNA-seq data of ENCODE consortium are given as numbers of RNA-seq fragments per kilobase of transcript per million fragments mapped (FPKM) for the breast cancer cell line MCF-7 and human mammary epithelial cell, HMEpC.