



**Figure S1. Sequencing analysis of *DMBT1* 3'UTRs.** We sequenced 3'UTR regions in genomic DNAs of all 8 HMECs and identified two SNPs, *rs8441* and *rs7383*, both presented in samples HMEC-1 and -5. We then sequenced the same region using cDNAs isolated from these two HMEC lines. The results revealed a DASE expression pattern for *DMBT1* in HMEC-1 and a bi-allelic expression pattern in HMEC-5.