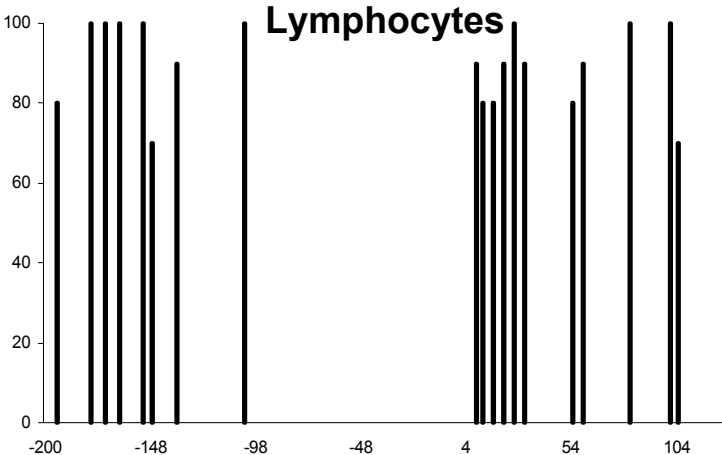
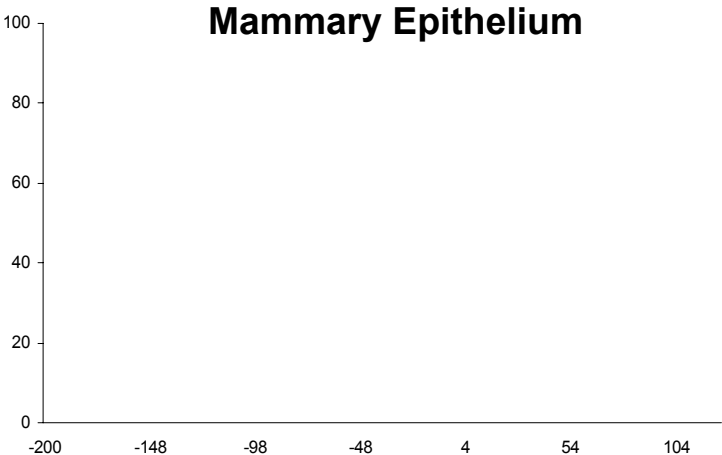
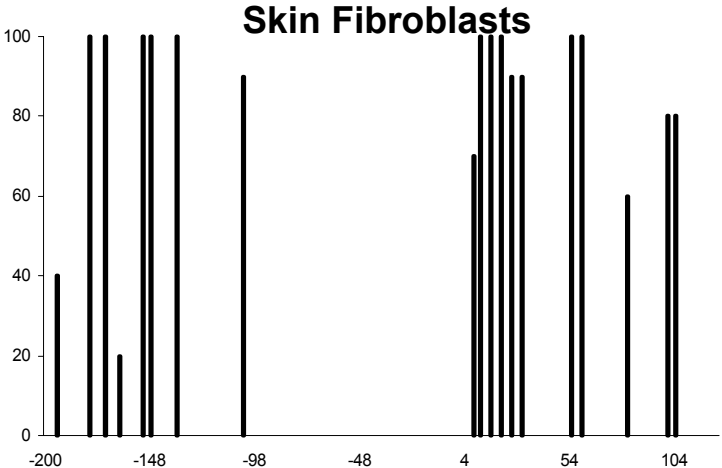
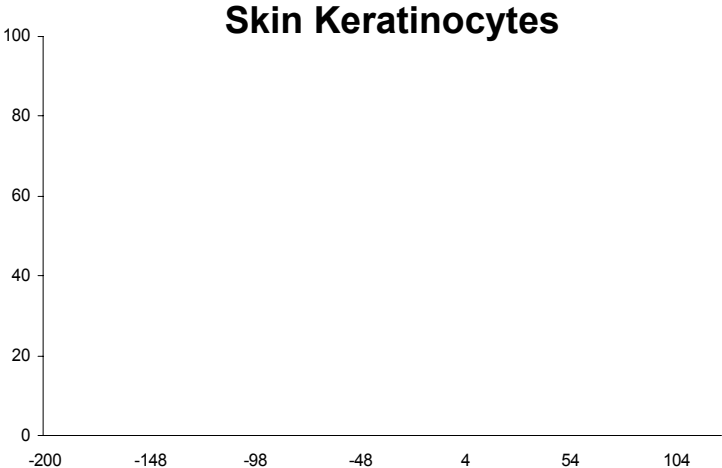


# Web Fig. - A



**Summary of 5-methylcytosine levels of bisulfite genomic sequencing of the opposing strand of the maspin promoter in 4 normal human cell types.** Ten cloned PCR products were sequenced to determine the percent methylation of the 19 CpG sites in the region analyzed. The x-axis is percent cytosine methylation and the y-axis is position relative to transcription start.