



Supplementary Figure 1a: Genotype distribution of MEIS 1 SNP among the healthy and case subjects' analysis. MEIS1: The analysis indicates that the T/G heterozygous mutation was mostly present in cases (70% vs.. 54%) while T/T homozygous mutation was mostly distributed among controls (30% vs.. 46%), whereas no case or control showed any wild type G/G allele.