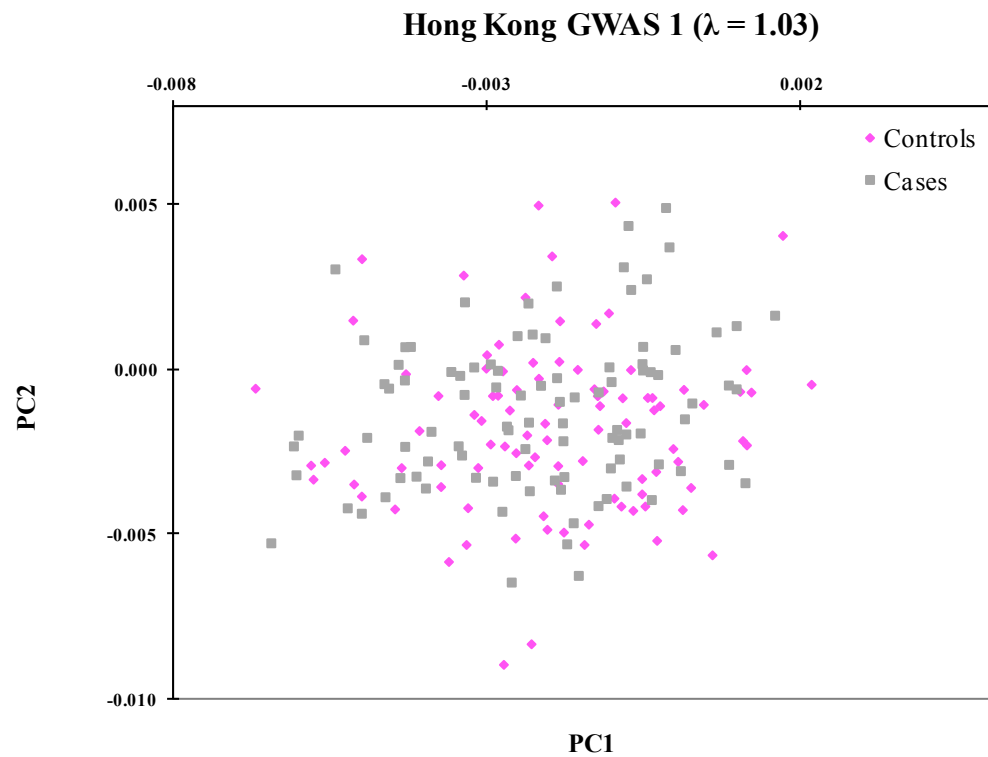
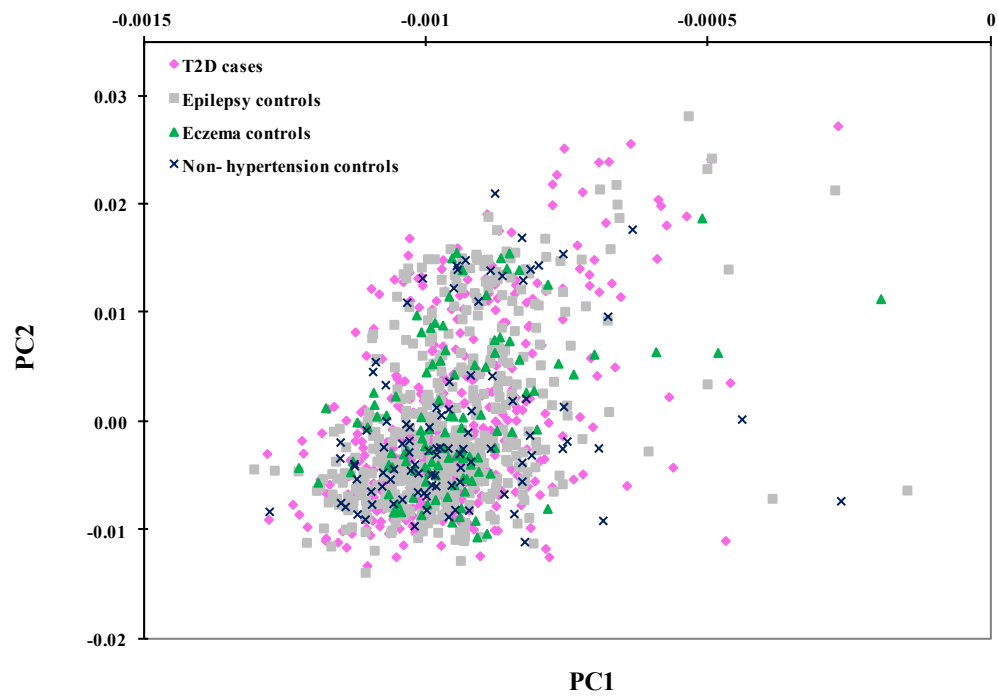


ESM Fig. 2. Multidimensional scaling analysis (MDS) plot shows the first two principal components, based on genotype data of 3 case-controls cohorts (Hong Kong GWAS 1, Hong Kong GWAS 2 and Shanghai GWAS) in the stage 1 genome scan of the present study without HapMap scaling.



Hong Kong GWAS 2 ($\lambda = 1.01$)



Shanghai GWAS ($\lambda = 1.04$)

