

Investigation of Inversion Polymorphisms in the Human Genome using Principal Components Analysis

Jianzhong Ma, Christopher I. Amos

Department of Genetics, The University of Texas MD Anderson Cancer Center, Houston, TX 77030, USA

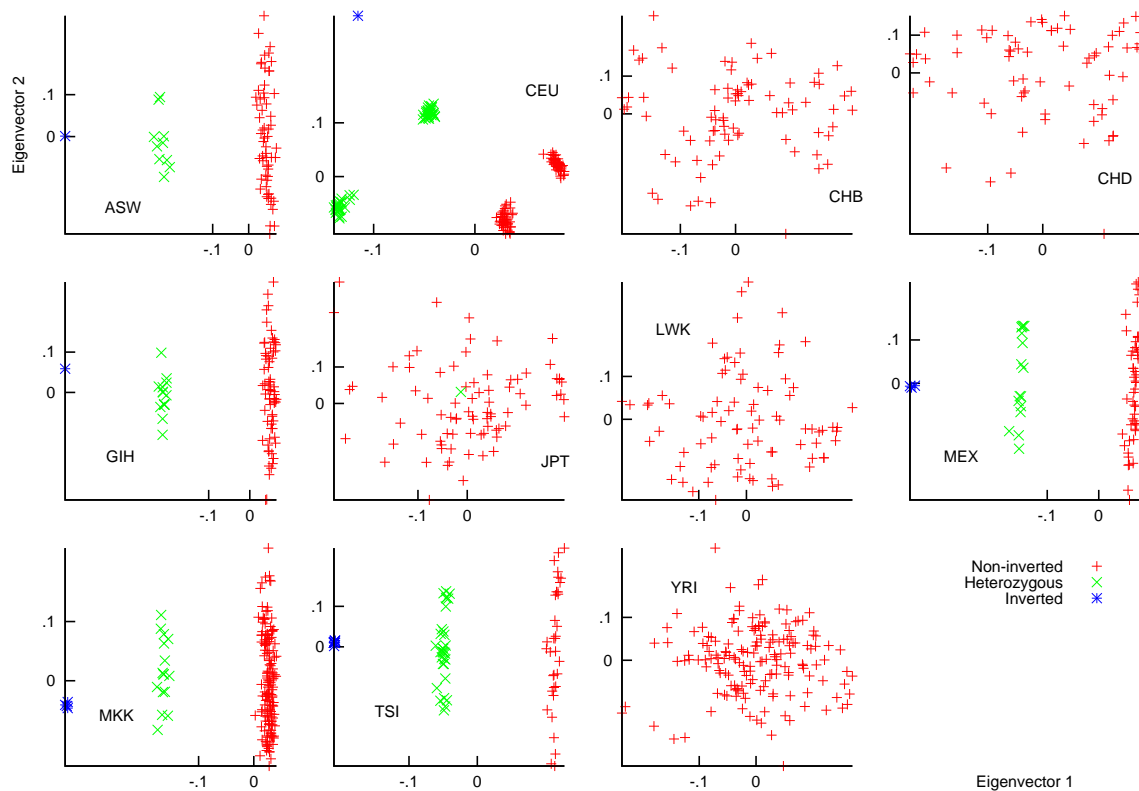


Figure S4. The first two eigenvectors obtained from PCA performed for each of the 11 HapMap populations using markers inside the 17q21.31 inversion region. The inversion genotypes were obtained by inspecting Figures 6 and 7.