

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

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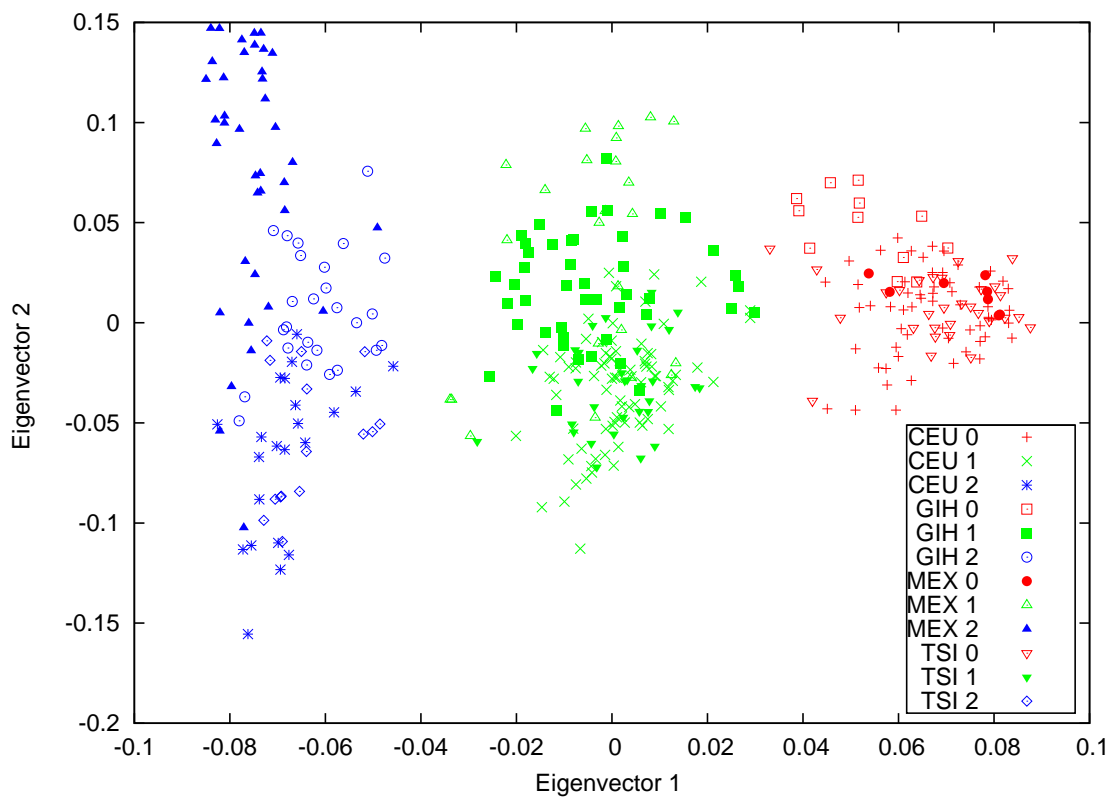


Figure S3. The first two eigenvectors obtained from PCA performed for pooled data of CEU, GIH, MEX, and TSI using markers inside the 8p23.1 inversion region. Genotyping of GIH was mainly based on this figure.