

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

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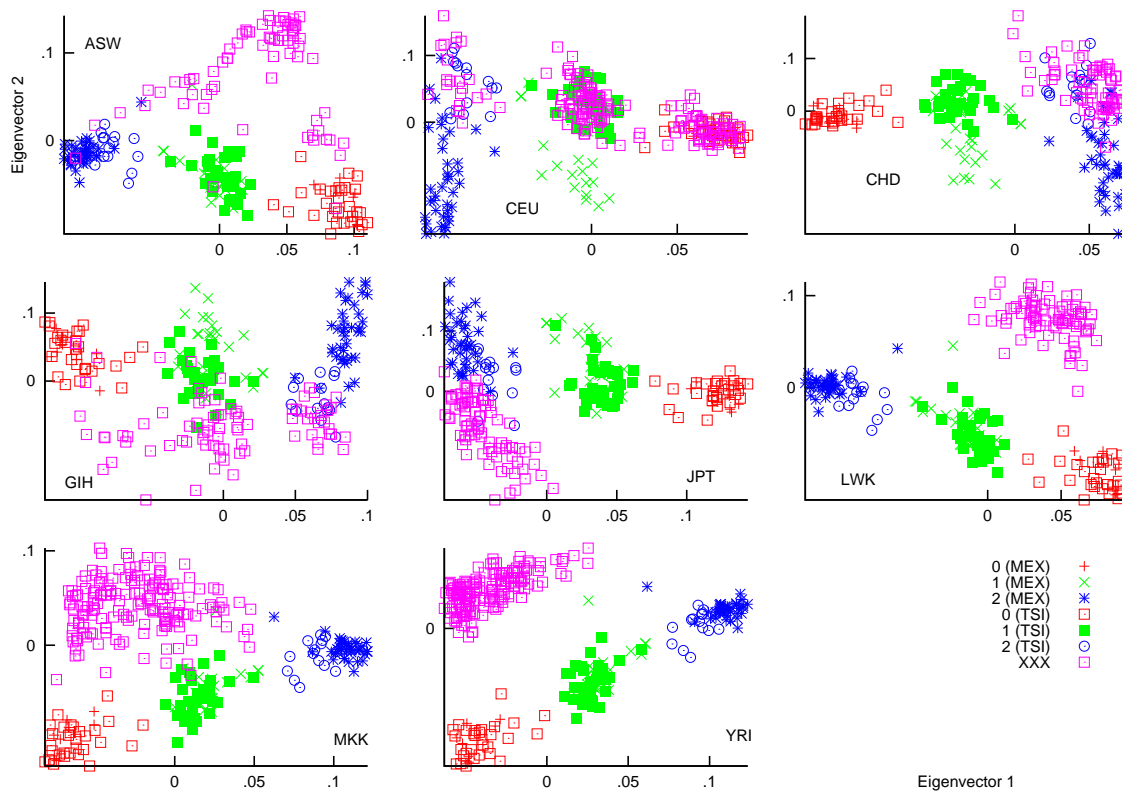


Figure S2. The first two eigenvectors obtained from PCA performed for pooled data of MEX and TSI with data of each of the other HapMap populations, represented by XXX, (except for CHB, for which the results are shown in Figure 4) using markers inside the 8p23.1 inversion region.