**Figure S1. Linkage disequilibrium (LD) plot across *PIK3CA-KCNMB3* region in CEU population.** The horizontal white bar depicts the 160-kb DNA segment analyzed in the sample. An LD plot is presented at the bottom of the figure: each diamond represents the magnitude of LD for the single pair of markers. The numbers under the variant name and above the plot indicates the order of the corresponding variants among all the variants in the region. The numbers inside the diamonds indicate the *r*2 value.

