Supplementary Information

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Supplementary Figure S1. Plot of first two principal components of HapMap YRI (Yoruban African) and CEU (European) samples and participants by (A) Case-Control Status and (B) Cohort Study

A.
Supplementary Figure S2. Genome-wide admixture mapping scan for loci associated with diabetic retinopathy in the African American Proliferative Diabetic Retinopathy Study where controls all had type 2 diabetes for at least 10 years and no proliferative diabetic retinopathy. The maximum locus genome statistic score was 2.67 on chromosome 14 which did not meet genome-wide statistical significance. The dashed line indicates the level of genome wide significance (locus genome statistic score of 5).