ESM Figure 1: Q-Q plots of the observed versus expected quantiles of –log10(pvalues) from 841,342 genotyped SNP associations with lnSIF1. A, M1: Univariate; B, M2: adjusted for age, sex, smoking status, skin tone, clinic latitude and any eGFR<60 ml/min/1.73m²; C, M3: adjusted for covariates in M2 + HbA₁c at DCCT eligibility, mean HbA₁c during DCCT, and mean HbA₁c during EDIC. Genomic control lambda for SIF1 were all equal to 1.02 (M1-M3).